

Supplementary file 1

RBPs	Domain
A2BP1	RRM
ACIN1	RRM
BOLL	RRM
BRUNOL4	RRM
BRUNOL5	RRM
BRUNOL6	RRM
CCBL2	RRM
RCAN3	RRM
CIRBP	RRM
CNOT4	RRM
COVA1	RRM
CPEB1	RRM
CPEB2	RRM
CPEB3	RRM
CPEB4	RRM
CPSF6	RRM
FLJ12529	RRM
CSTF2	RRM
CSTF2T	RRM
CUGBP1	RRM
CUGBP2	RRM
D10S102	RRM
DAZ2	RRM
DAZ3	RRM
DAZ4	RRM
DAZL	RRM
DAZAP1	RRM
DNAJC17	RRM
DKFZp434J0315	RRM
DKFZp686F0223	RRM
DND1	RRM
DRB1	RRM
ELAV1	RRM
ELAVL2	RRM
ELAVL3	RRM
ELAVL4	RRM
EWS	RRM

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RBM44	RRM
FUS	RRM
G3BP1	RRM
G3BP2	RRM
GRSF1	RRM
HNRPH1	RRM
HNRPH2	RRM
HNRPH3	RRM
HNRPLL	RRM
HNRPA0	RRM
HNRPA1	RRM
HNRPCL1	RRM
HNRPC	RRM
HNRPDL	RRM
HNRPD	RRM
HNRPF	RRM
HNRPG	RRM
HNRPGT	RRM
HNRPL	RRM
HNRPM	RRM
SYNCRIP	RRM
HNRPR	RRM
HTF9C	RRM
HTATSF1	RRM
EIF3S4	RRM
EIF3S9	RRM
EIF4B	RRM
EIF4H	RRM
LARP4	RRM
LARP5	RRM
LARP7	RRM
LA	RRM
LKAP	RRM
LOC138046	RRM
MATR3	RRM
MGC10433	RRM
MINT	RRM
MKI67IP	RRM
MSI1	RRM
MSI2H	RRM

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MSSP-1	RRM
MTHFSD	RRM
MYEF2	RRM
NCBP2	RRM
NOL8	RRM
NONO	RRM
NCL	RRM
PABP1	RRM
PABP2	RRM
PABP3	RRM
PABP4	RRM
PABP5	RRM
PABPCP2	RRM
C20orf119	RRM
POLDIP3	RRM
SF3B14	RRM
PPIE	RRM
PPIL4	RRM
PPRC1	RRM
PPARGC1A	RRM
PPARGC1B	RRM
PSPC1	RRM
PTBP2	RRM
Q8TC92	RRM
RALY	RRM
RAVER1	RRM
RAVER2	RRM
RBM12B	RRM
RBM15B	RRM
RBM35A	RRM
RBM35B	RRM
RBM10	RRM
RBM11	RRM
RBM12	RRM
RBM14	RRM
RBM15	RRM
RBM16	RRM
RBM18	RRM
RBM19	RRM
RBM22	RRM

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	RBM23		RRM	
	RBM24		RRM	
	RBM25		RRM	
	RBM26		RRM	
	RBM27		RRM	
	RBM28		RRM	
	RBM33		RRM	
	RBM34		RRM	
	RBM38		RRM	
	RBM39		RRM	
	RBM3		RRM	
	RBM41		RRM	
	RBM43		RRM	
	RBM4		RRM	
	RBM5		RRM	
	RBM6		RRM	
	RBM7		RRM	
	RBM8A		RRM	
	RBM9		RRM	
	RBMS1		RRM	
	RBMS2		RRM	
	RBMS3		RRM	
	RBMX2		RRM	
	RBMX1F		RRM	
	RBMX2FP		RRM	
	RBP56		RRM	
	RPMS		RRM	
	RPMS2		RRM	
	RBMX1A1		RRM	
	RDM1		RRM	
	RNPC3		RRM	
	RNPS1		RRM	
	HNRPA0		RRM	
	HNRPA2B1		RRM	
	HNRPA3		RRM	
	HNRPAB		RRM	
	ROD1		RRM	
	RP11-493K23.3		RRM	
	RP11-78J21.1		RRM	
	SNRP70		RRM	

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SNRPB2	RRM
SAFB1	RRM
SAFB2	RRM
SART3	RRM
SET1A	RRM
SF3B4	RRM
SFPQ	RRM
SFRS11	RRM
SFRS12	RRM
SFRS15	RRM
SFRS1	RRM
SFRPS2B	RRM
SFRPS2	RRM
SFRPS3	RRM
SFRS4	RRM
SFRPS5	RRM
SFRS6	RRM
SFRS7	RRM
SFRPS9	RRM
SIAHBP1	RRM
SLIRP	RRM
SLTM	RRM
SNRPA	RRM
SPF45	RRM
SR140	RRM
SRrp35	RRM
SYNJ1	RRM
SYNJ2	RRM
TARDBP	RRM
THOC4	RRM
TIA1	RRM
TIAR	RRM
TNRC6A	RRM
TNRC6C	RRM
TNRC4	RRM
TRA2A	RRM
TRA2B	RRM
TRSPAP1	RRM
TUT1	RRM
U1SNRNPBP	RRM

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U2AF1L4	RRM
U2AF1	RRM
U2AF2	RRM
ZRSR1	RRM
ZRSR2	RRM
UHMK1	RRM
ZCRB1	RRM
CTA-126B4.3	RRM
ZNF638	RRM

+-----+-----+

AKAP1	KH
ANKHD1	KH
ANKRD17	KH
ASCC1	KH
DDX43	KH
DDX53	KH
ERAL1	KH
FMR1	KH
FUBP1	KH
FUBP2	KH
FUBP3	KH
FXR1	KH
FXR2	KH
HNRPK	KH
KHDRBS3	KH
KHDRBS2	KH
KRR1	KH
NOVA1	KH
NOVA2	KH
PCBP1	KH
PCBP2	KH
PCBP3	KH
PCBP4	KH
PNO1	KH
PNPT1	KH
QKI	KH
RKHD1	KH
RKHD3	KH
RPS3	KH
SAM68	KH

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```
| SF1 | KH |
| TDRKH | KH |
| HDLBP | KH |
+-----+
| CDKN2AIP | dsrm |
| DGCR8 | dsrm |
| DHX30 | dsrm |
| DHX9 | dsrm |
| DICER | dsrm |
| DSRAD | dsrm |
| DUS2L | dsrm |
| E2AK2 | dsrm |
| EXOS9 | dsrm |
| ILF3 | dsrm |
| ADAD2 | dsrm |
| SLC4A1AP | dsrm |
| NKRF | dsrm |
| PRKRA | dsrm |
| RED1 | dsrm |
| RED2 | dsrm |
| MRPL44 | dsrm |
| RNASEN | dsrm |
| SON | dsrm |
| STAU1 | dsrm |
| STAU2 | dsrm |
| STRBP | dsrm |
| Tenr | dsrm |
| TRBP2 | dsrm |
+-----+
| ACF | dsrm RRM |
| MGC27016 | dsrm RRM |
| PTBP1 | dsrm RRM |
+-----+
| IGF2BP2 | KH RRM |
| IGF2BP3 | KH RRM |
| IGF2BP1 | KH RRM |
+-----+
| APOBEC3F | other |
| APOBEC1 | other |
| APOBEC2 | other |
```

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ADAT1	other
CPSF5	other
CSDC2	other
CSTF1	other
CSTF3	other
DDX19B	other
DDX17	other
DKC1	other
DUSP11	other
EPRS	other
EXOSC1	other
EXOSC7	other
FBL	other
HNRPUL1	other
HNRPU	other
IF2B	other
IREB1	other
IREB2	other
JOSD2	other
LRPPRC	other
LSM5	other
MOV10L1	other
RNMT	other
NCBP1	other
NHP2L1	other
NOL3	other
NOL4	other
NOP56	other
NPM	other
NPM1	other
NUFIP1	other
HRB	other
NXF1	other
NXF2	other
NXF5	other
PAIP1	other
PAPOLA	other
PAPOLB	other
PAPOLG	other
PPP1R8	other

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PSMA1	other
PSMA6	other
RAD51AP1	other
RAE1	other
RENT1	other
RIB1	other
RPL12	other
RPL13	other
RL14	other
RPL18	other
RPL19	other
RPL22	other
RPL27A	other
RPL28	other
RPL29	other
RPL3	other
RPL34	other
RPL35A	other
RPL3L	other
RPL4	other
RPL5	other
RPL6	other
RPLP0	other
RPLP1	other
RPLP2	other
MRPL3	other
MRPL12	other
MRPL23	other
RNASEL	other
RNASE1	other
RNH1	other
RO60	other
RPL18A	other
RPL21	other
RPL7	other
RPL9	other
RPP14	other
RPP29	other
RPS13	other
RPS17	other

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RPS27	other
RPS10	other
RPS12	other
RPS19	other
RPS2	other
RPS4Y1	other
RPS5	other
RPS9	other
MRPS15	other
RTCD1	other
SCAND1	other
SLBP	other
SRP14	other
SRP68	other
SRP72	other
SRP9	other
SUPV3L1	other
SURF6	other
GARS	other
HARS	other
MARS	other
SARS	other
SYWC	other
TACC1	other
TACC2	other
TARB1	other
TEP1	other
YBOX1	other
YBOX2	other
ZNF239	other
ZRANB2	other

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List of RBPs with nuclear localization.

```
+-----+
| RBPs |
+-----+
| APOBEC3F |
| ACF |
| ACIN1 |
| BRUNOL4 |
| CIRBP |
| CPSF5 |
| CPSF6 |
| CSDC2 |
| CSTF1 |
| CSTF2 |
| CSTF3 |
| CSTF2T |
| CUGBP1 |
| DAZ2 |
| DAZ3 |
| DAZ4 |
| DAZL |
| DAZAP1 |
| DDX19B |
| DDX17 |
| DDX53 |
| DGCR8 |
| DHX9 |
| DKC1 |
| DRB1 |
| DSRAD |
| DUSP11 |
| EWS |
| EXOSC1 |
| EXOSC7 |
| EXOS9 |
| FBL |
| FMR1 |
| FUBP1 |
| FUBP2 |
| FUBP3 |
| FUS |
| FXR1 |
| FXR2 |
| G3BP1 |
| HNRPH1 |
| HNRPH2 |
| HNRPH3 |
| HNRPUL1 |
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Supplementary file 2

HNRPLL |
HNRPA0 |
HNRPA1 |
HNRPC |
HNRPCL1 |
HNRPD |
HNRPDL |
HNRPF |
HNRPG |
HNRPK |
HNRPL |
HNRPM |
SYNCRIP |
HNRPR |
HNRPU |
HTATSF1 |
ILF3 |
KHDRBS3 |
KRR1 |
LA |
LARP7 |
LRPPRC |
LSM5 |
MATR3 |
RNMT |
MINT |
MKI67IP |
MSI1 |
MSSP-1 |
MYEF2 |
SLC4A1AP |
NCBP1 |
NCBP2 |
NHP2L1 |
NKRF |
NOL3 |
NOL4 |
NONO |
NOP56 |
NOVA1 |
NOVA2 |
NPM |
NPM1 |
NCL |
NUFIP1 |
HRB |
NXF1 |
NXF2 |

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Supplementary file 2

NXF5 |
PABP1 |
PABP2 |
PAPOLA |
PAPOLB |
PAPOLG |
PCBP1 |
PCBP2 |
PCBP3 |
PCBP4 |
POLDIP3 |
SF3B14 |
PPP1R8 |
PPIE |
PPARGC1A |
PPARGC1B |
PSMA1 |
PTBP1 |
PTBP2 |
RAD51AP1 |
RAE1 |
RALY |
RAVER1 |
RAVER2 |
RBM15B |
RBM10 |
RBM12 |
RBM14 |
RBM15 |
RBM19 |
RBM22 |
RBM23 |
RBM26 |
RBM27 |
RBM28 |
RBM34 |
RBM39 |
RBM4 |
RBM5 |
RBM6 |
RBM8A |
RBM9 |
RBMS1 |
RBMS2 |
RBP56 |
RBM1A1 |
RED1 |
RED2 |

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Supplementary file 2

RPL3 |
RPL5 |
RNASEN |
RNPS1 |
HNRPA0 |
HNRPA2B1 |
HNRPA3 |
HNRPAB |
ROD1 |
RPP14 |
RPP29 |
RTCD1 |
SNRP70 |
SNRPB2 |
SAFB1 |
SAFB2 |
SAM68 |
SCAND1 |
SET1A |
SF1 |
SF3B4 |
SFPQ |
SFRS11 |
SFRS12 |
SFRS15 |
SFRS1 |
SFRPS2 |
SFRPS3 |
SFRS4 |
SFRPS5 |
SFRS6 |
SFRS7 |
SFRPS9 |
SLBP |
SNRPA |
SON |
SPF45 |
SRP68 |
SRP9 |
SRrp35 |
STAU2 |
SURF6 |
TACC1 |
TACC2 |
TARDBP |
TARB1 |
TEP1 |
THOC4 |

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Supplementary file 2

TNRC4
TRA2A
TRA2B
TRBP2
TRSPAP1
U2AF1
U2AF1L4
U2AF2
ZRSR2
UHMK1
HDLBP
YBOX1
YBOX2
ZNF239
ZNF638
ZRANB2
ZRSR1

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Supplementary file 3

List of libraries used in the RBP expression analysis . Symbol ' #' identify the tissue name; first column is the library id; second column is the library name; third column is the total of tags (information obtained from SAGEGenie).

===== SAGE: Libraries from normal tissue =====

#Brain_astrocyte

9 SAGE_Brain_astrocyte_normal_CL_NHA5 51481

#Brain_cerebellum

34 SAGE_Brain_normal_cerebellum_B_1 50385

#Brain_cortex

8 SAGE_Brain_normal_cortex_B_BB542 94233

10 SAGE_Brain_normal_cortex_B_pool6 62451

133 SAGE_Brain_normal_peds_cortex_B_H1571 77554

#Breast

47 SAGE_Breast_normal_epithelium_AP_Br_N 50512

420 SAGE_Breast_normal_myoepithelium_AP_IDC7 69006

131 SAGE_Breast_normal_myoepithelium_AP_myoepithelial1 57222

182 SAGE_Breast_normal_organoid_B 58181

#Colon

11 SAGE_Colon_normal_B_NC1 49610

#Eye

1363 SAGE_Eye_lens_B_UIH10 85898

#Heart

136 SAGE_Heart_normal_B_1 83063

#Liver

135 SAGE_Liver_normal_B_1 66308

#Lung

99 SAGE_Lung_normal_B_1 88708

#Muscle

141 SAGE_Muscle_normal_B_old 53889

142 SAGE_Muscle_normal_B_young 53947

#Placenta

160 SAGE_Placenta_first_trimester_normal_B_1 89265

364 SAGE_Placenta_hydatidiform_mole_B_1 101677

161 SAGE_Placenta_normal_B_1 118083

#Prostate

59 SAGE_Prostate_normal_B_2 64058

169 SAGE_Prostate_normal_epithelium_CS_confluent 71897

203 SAGE_Prostate_normal_epithelium_CS_senescent 71717

72 SAGE_Prostate_normal_MD_PR317 59277

#Retina

164 SAGE_Retina_macula_normal_B_HMAC2 102417

162 SAGE_Retina_Peripheral_normal_B_1 59661

163 SAGE_Retina_Peripheral_normal_B_2 105312

165 SAGE_Retina_Pigment_epithelium_normal_B_1 53666

#Spinal_cord

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167 SAGE_Spinal_cord_normal_B_1 54785

#Stomach

1624 SAGE_Stomach_normal_MD_14S 73353

#Thyroid

389 SAGE_Thyroid_normal_B_001 115938

#Vascular_endothelium

180 SAGE_Vascular_endothelium_normal_breast_associated_AP_1 57023

429 SAGE_Vascular_endothelium_normal_liver_associated_AP_NLEC1 77759

29 SAGE_Vascular_normal_CS_control 51642

30 SAGE_Vascular_normal_CS_VEGF + 57316

#White_Blood_Cells

1574 SAGE_White_Blood_Cells_monocytes_normal_AP_A2 98826

1571 SAGE_White_Blood_Cells_monocytes_normal_AP_C1 101722

1570 SAGE_White_Blood_Cells_monocytes_normal_AP_P2 54447

1576 SAGE_White_Blood_Cells_monocyte_cells_normal_AP_A1 53825

1575 SAGE_White_Blood_Cells_monocyte_cells_normal_AP_P1 51653

1573 SAGE_White_Blood_Cells_monocyte_normal_AP_A1 99961

1572 SAGE_White_Blood_Cells_monocyte_normal_AP_P1 101225

#Stem_Cell

1604 SAGE_Embryonic_stem_cell_BG01_normal_p20_CL_SHE19 201668

1332 SAGE_Embryonic_stem_cell_HSF6_normal_p50_CL_SHES9 224488

1386 SAGE_Embryonic_stem_cell_H13_normal_p22_CL_SHE15 221101

1391 SAGE_Embryonic_stem_cell_H1_normal_p31_CL_SHE17 276203

1384 SAGE_Embryonic_stem_cell_HES4_normal_p36_CL_SHE11 209232

1388 SAGE_Embryonic_stem_cell_H1_normal_p54_CL_SHE16 218214

1331 SAGE_Embryonic_stem_cell_HES3_normal_p16_CL_SHE10 205353

1334 SAGE_Embryonic_stem_cell_H14_normal_p22_CL_SHE14 212170

1333 SAGE_Embryonic_stem_cell_H7_normal_p33_CL_SHE13 272465

1003 SAGE_Embryonic_stem_cell_H9_normal_p38_CL_SHES2 401432

844 SAGE_Embryonic_stem_cell_H9_normal_p38_CL_SHES1 151735

===== SAGE: Libraries from tumor tissue =====

#Brain_astrocyte

343 SAGE_Brain_astrocytoma_grade_III_B_343 100158

361 SAGE_Brain_astrocytoma_grade_III_B_407 108312

385 SAGE_Brain_astrocytoma_grade_III_B_439 107824

344 SAGE_Brain_astrocytoma_grade_III_B_584 103008

349 SAGE_Brain_astrocytoma_grade_III_B_828 99939

54 SAGE_Brain_astrocytoma_grade_III_B_H1020 51573

350 SAGE_Brain_astrocytoma_grade_III_B_H1055 109886

563 SAGE_Brain_astrocytoma_grade_III_B_H272 96059

283 SAGE_Brain_astrocytoma_grade_III_B_H970 106982

384 SAGE_Brain_astrocytoma_grade_III_B_R140 118733

351 SAGE_Brain_astrocytoma_grade_III_B_R927 107344

345 SAGE_Brain_astrocytoma_grade_II_B_H127 114489

144 SAGE_Brain_astrocytoma_grade_II_B_H359 105764

145 SAGE_Brain_astrocytoma_grade_II_B_H388 106285

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346 SAGE_Brain_astrocytoma_grade_II_B_H501 128309
347 SAGE_Brain_astrocytoma_grade_II_B_H516 108116
564 SAGE_Brain_astrocytoma_grade_II_B_H518 116022
348 SAGE_Brain_astrocytoma_grade_II_B_H530 102439
146 SAGE_Brain_astrocytoma_grade_II_B_H563 88568
50 SAGE_Brain_astrocytoma_grade_I_B_H1043 75922

#Brain_cortex

137 SAGE_Brain_ependymoma_B_1150 62373
139 SAGE_Brain_ependymoma_B_1394 56314
267 SAGE_Brain_ependymoma_B_H580 68614
147 SAGE_Brain_ependymoma_B_R1023 122690
148 SAGE_Brain_ependymoma_B_R353 73822
149 SAGE_Brain_ependymoma_B_R455 51825
406 SAGE_Brain_ependymoma_B_R510 84073
388 SAGE_Brain_ependymoma_B_R512 75379
150 SAGE_Brain_ependymoma_B_R582 52189
151 SAGE_Brain_ependymoma_B_R628 120431
86 SAGE_Brain_glioblastoma_B_GBM1062 59762
4 SAGE_Brain_glioblastoma_B_H1110 68986
354 SAGE_Brain_glioblastoma_B_H1353 124805
352 SAGE_Brain_glioblastoma_B_H1425C 88990
356 SAGE_Brain_glioblastoma_B_H833 100600
7 SAGE_Brain_glioblastoma_B_pooled 56428
357 SAGE_Brain_glioblastoma_B_R20 101053
358 SAGE_Brain_glioblastoma_B_R336 102322
359 SAGE_Brain_glioblastoma_B_R70 99099
1648 SAGE_Brain_glioblastoma_CL_28d 110624
1650 SAGE_Brain_glioblastoma_CL_4d 115212
3 SAGE_Brain_glioblastoma_CL_H392 55990
68 SAGE_Brain_glioblastoma_CL_H54 + EGFRvIII 56982
67 SAGE_Brain_glioblastoma_CL_H54 +LacZ 66908
1649 SAGE_Brain_glioblastoma_CL_stem_cells 108414
41 SAGE_Brain_glioblastoma_control_CL_H247 60428
42 SAGE_Brain_glioblastoma_hypoxia_CL_H247 71765

#Brain_cerebellum

123 SAGE_Brain_medulloblastoma_B_96-04-P019 52645
126 SAGE_Brain_medulloblastoma_B_97-05-P015 69971
408 SAGE_Brain_medulloblastoma_B_97-05-P312 74295
413 SAGE_Brain_medulloblastoma_B_98-05-P040 89258
386 SAGE_Brain_medulloblastoma_B_98-09-P558 85984
414 SAGE_Brain_medulloblastoma_B_C609 74612
399 SAGE_Brain_medulloblastoma_B_DL5 83671
400 SAGE_Brain_medulloblastoma_B_DL7 68392
412 SAGE_Brain_medulloblastoma_B_H1322 59498
401 SAGE_Brain_medulloblastoma_B_H1413 61853
397 SAGE_Brain_medulloblastoma_B_H275 72318

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383 SAGE_Brain_medulloblastoma_B_H306 60454
407 SAGE_Brain_medulloblastoma_B_H484 57469
415 SAGE_Brain_medulloblastoma_B_H876 67404
263 SAGE_Brain_medulloblastoma_B_H972 85376
303 SAGE_Brain_medulloblastoma_CL_4-HCR 66599
243 SAGE_Brain_medulloblastoma_CL_D283 66316

#Breast

156 SAGE_Breast_carcinoma_associated_myoepithelium_AP_DCIS6 81452
158 SAGE_Breast_carcinoma_associated_myoepithelium_AP_DCIS7 51993
416 SAGE_Breast_carcinoma_associated_myofibroblast_AP_IDC7 65091
1169 SAGE_Breast_carcinoma_associated_myofibroblast_AP_T112603 63539
1164 SAGE_Breast_carcinoma_associated_myofibroblast_AP_T392303 73967
424 SAGE_Breast_carcinoma_associated_stroma_B_DCIS6 57049
419 SAGE_Breast_carcinoma_associated_stroma_B_IDC7 68024
82 SAGE_Breast_carcinoma_B_95-347 67070
431 SAGE_Breast_carcinoma_B_BWHT18 50701
171 SAGE_Breast_carcinoma_B_DCIS-4 60605
173 SAGE_Breast_carcinoma_B_IDC-3 68891
174 SAGE_Breast_carcinoma_B_IDC-4 64095
175 SAGE_Breast_carcinoma_B_IDC-5 60451
77 SAGE_Breast_carcinoma_CL_MCF7control_0h 59877
79 SAGE_Breast_carcinoma_CL_MCF7estradiol_10H 59583
78 SAGE_Breast_carcinoma_CL_MCF7estradiol_3h 59583
187 SAGE_Breast_carcinoma_epithelium_AP_DCIS-3 57402
153 SAGE_Breast_carcinoma_epithelium_AP_DCIS6 72857
157 SAGE_Breast_carcinoma_epithelium_AP_DCIS7 89184
418 SAGE_Breast_carcinoma_epithelium_AP_IDC7 73410
1163 SAGE_Breast_carcinoma_epithelium_AP_T112603 65935
1165 SAGE_Breast_carcinoma_epithelium_AP_T392303 66520
1166 SAGE_Breast_carcinoma_MD_LCIS 67834

#Colon

e13 SAGE_Colon_adenocarcinoma_B_Tu102 55700
2 SAGE_Colon_adenocarcinoma_CL_Caco2 60682
1 SAGE_Colon_adenocarcinoma_CL_HCT116 55641
6 SAGE_Colon_adenocarcinoma_CL_RKO 51294
5 SAGE_Colon_adenocarcinoma_CL_SW837 61148
428 SAGE_Colon_carcinoma_CL_Hct116_p53_knockout_Anoxia 79539
426 SAGE_Colon_carcinoma_CL_Hct116_p53_knockout_Normal_Oxygen 79739
425 SAGE_Colon_carcinoma_CL_Hct116_wildtype_p53_Anoxia 79411
427 SAGE_Colon_carcinoma_CL_Hct116_wildtype_p53_Normal_Oxygen 80826

#Liver

266 SAGE_Liver_cholangiocarcinoma_B_K1 60319
504 SAGE_Liver_cholangiocarcinoma_CL_K3 54842
503 SAGE_Liver_cholangiocarcinoma_CL_K4 52973

#Lung

587 SAGE_Lung_adenocarcinoma_MD_L10 86887

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1366 SAGE_Lung_tumor_associated_chronic_inflammation_B_UIHM0 98962

#Prostate

73 SAGE_Prostate_adenocarcinoma_MD_PR317 64951

60 SAGE_Prostate_carcinoma_B_pool2 66034

57 SAGE_Prostate_carcinoma_CL_LNCaP 60250

58 SAGE_Prostate_carcinoma_CL_LNCaP_no-DHT 62160

#Stomach

97 SAGE_Stomach_adenocarcinoma_B_G234 64925

1443 SAGE_Stomach_adenocarcinoma_MD_HG7 93714

1445 SAGE_Stomach_adenocarcinoma_MD_HS29 58831

186 SAGE_Stomach_carcinoma_B_G189 63075

170 SAGE_Stomach_carcinoma_B_xenograft_X101 69749

244 SAGE_Stomach_carcinoma_B_xenograph_X43 51620

#Thyroid

365 SAGE_Thyroid_follicular_carcinoma_B_TT004 124664

#Vascular_endothelium

159 SAGE_Vascular_endothelium_breast_carcinoma_AP_DCIS6 65314

185 SAGE_Vascular_endothelium_hemangioma_B_146 75680

#White_Blood_Cells

155 SAGE_White_Blood_Cells_breast_carcinoma_associated_AP_DCIS6 66176

List of libraries used in the RBP expression analysis. Symbol '#' identify the tissue name; first column is our local library identification; second column is the library name; third column is the total of tags (information obtained from <http://mps.s.licr.org>).

=====**MPSS : Libraries from normal tissue**=====

#Adrenal

1 HuAG.basl 1365238

#Bladder

2 HuBl.basl 1335841

#Bone_marrow

3 HuBM.basl 1445400

#Kidney

12 HuKi.basl 1461001

#Pituitary_gland

15 HuPG.basl 1379052

#Salivary_gland

21 HuSG.basl 1350021

#Small_intestine

22 HuSI.basl 1336379

#Testis

25 HuTe.basl 1292053

#Thymus

26 HuTm.basl 1341049

#Trachea

27 HuTr.basl 1318303

#Uterus

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29 HuUt.basl 1335188

#Monocytes

49 HuMo.basl 1318365

#Peripheral_blood_lymphocyte

55 HuPB.ctrl 1367070

#Skeletal_muscle

38 skeletal.muscle 1147429

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Supplementary file 4

List of tags used for SAGE and MPSS analyses

```
+-----+-----+-----+
| RBPs | SAGE tags | MPSS tags |
+-----+-----+-----+
| A2BP1 | SAGE: CCGTAAGATC | MPSS: AGTTGGATATTAA |
| APOBEC3F | SAGE: CAGAGCTGTG | MPSS: ACAGACACCAGCA |
| APOBEC1 | SAGE: GATCAACAAA | MPSS: AACAAAATCGGCA |
| ACF | SAGE: AAGTAAAAAA | MPSS: ATAAATAATATGT
TCTGAAATTACTC |
| ACIN1 | SAGE: AAGTAAAGAA | MPSS: GCAGGGACACCAA |
| ADAT1 | SAGE: CAGAGAGCAC | MPSS: AGAAACCCACCGG |
| AKAP1 | SAGE: TCACCTGTAG | MPSS: TGGGAAC TTTTGTG |
| ANKHD1 | SAGE: CCACTGCTGC TATAATTGTT AAGACTATTT
CCACTTGCTG | MPSS: CCTCATGCCACTG ATATGACCTTAGT
CGGCCAAGCATCA CCTCATGCCACTT |
| ANKRD17 | SAGE: GAAAAATTAA AACAGTGTAC AACAGTGTGC |
MPSS: AACAGTTAATTTG CTGAAGTTTTAGA |
| ASCC1 | SAGE: TCGGTACCAC | MPSS: TTGGAAAGCACTA |
| BOLL | SAGE: AGAACAGTAG TTTTATCCAT | MPSS:
TCAAATACATTAA CAGCTTCAACGAA |
| BRUNOL4 | SAGE: AAGAAGAAGA | MPSS: GGCATGAAGAGGC |
| BRUNOL5 | SAGE: AAGTTTCAAC | MPSS: GGCATGAAGAGGC
ATGGGACCAAAT |
| BRUNOL6 | SAGE: TCAGATAGTA TGCTTAACCG | MPSS:
TGATAACATGTCA AATGGTGGTGAAA |
| CCBL2 | SAGE: GCAGATTTAT | MPSS: ACTTTTCACTTTT |
| RCAN3 | SAGE: TCATTGTGGT | MPSS: ATCATTGTGCATT |
| CDKN2AIP | SAGE: AATGTGGGT | MPSS: TAGTACTCCTTGA |
| CIRBP | SAGE: AAGTGGGTGC | MPSS: GAACGGACTGTGA |
| CNOT4 | SAGE: GGGGAAAGCA | MPSS: ACCATCGGGAGCA |
| COVA1 | SAGE: GAGAGTATTT TGAGTGGACC | MPSS:
TGTTTCAGAATTG |
| CPEB2 | SAGE: TTCAAATAAA CTTTTATGGA | MPSS:
CACTTCCGCTGGA ATATTTATTTTAC |
| CPEB4 | SAGE: GTGATTTTTG | MPSS: TTTGCAGTGTAAT |
| CPSF5 | SAGE: TGTTACTGCT TATTTAGAAA | MPSS:
ACTGTTCTTTTTT |
| CPSF6 | SAGE: GATAAAATGA | MPSS: TGAATAAAGCAAA |
| CSDC2 | SAGE: AGGAATAAAG | MPSS: CCTTGCCTTG TAG |
| CSTF2 | SAGE: TTGCCTCCTG | MPSS: ATGAGAAGGCTGC |
| CSTF3 | SAGE: TCAGGAGACG | MPSS: ATTACTGGTGGGG |
| CSTF2T | SAGE: CAGAAAATAA ATTATATTAT | MPSS:
ATTTTGAGATTCT AGATTGCCATGCT |
| CUGBP1 | SAGE: TCGATGTGGC | MPSS: AGGACCTGCTGCA |
| CUGBP2 | SAGE: AAACGTTTCT TCCCCACCCA | MPSS:
AAAGTTTTTTCTG CTAACCCAGAGG |
| DAZAP1 | SAGE: AGCAAGCCCC | MPSS: AGTTTGCCCCTCC |
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| DDX17 | SAGE: CAGATTAGTT | MPSS: CAGTCTGGAATAA |
| DDX53 | SAGE: TTAAAGATAT | MPSS: AAGAAAACCTGGA
AAGAGAACCTGGA |
| DGCR8 | SAGE: CTTCAAGGCC CTTGTCCCTT | MPSS:
CAGAGCTGTTGCC |
DHX30	SAGE: GTGGAGCGGA	MPSS: GGGTGCTGGACC
DHX9	SAGE: CTATGTGTTA	MPSS: TCTTAAAAATACC
DICER	SAGE: CTGCAGAAAT	MPSS: ACAATTACCTTTA
DNAJC17	SAGE: CGCCAGGCGG	MPSS: GCACGGATGCAGC
DKC1	SAGE: CTAGCAGAGC	MPSS: TCATGCTAGCAGA
DKFZp434J0315	SAGE: TATAAGTTTT	MPSS: TTAAAAGTGAAA
DKFZp686F02235	SAGE: TGCCTGTAAA CCCTTAGCAA GTATGTAGTT	
MPSS: CTCATCCCCTGAC TCTCCTTAAGCCA		
DND1	SAGE: GCATTCTGGT	MPSS: ATACCTTCCCCC
DRB1	SAGE: TATTTTTGAA	MPSS: GAAGTTTACCTTG
DSRAD	SAGE: CCCCTCTGAG	MPSS: GCACATCCCCTGCG
DUSP11	SAGE: TTTACCAGAG GTGGTCATTT	MPSS:
CAGGGTCACTCAG AGCAGAGACCATA		
DUS2L	SAGE: GACCAGGGGC	MPSS: TGAGTGACAGGGT
E2AK2	SAGE: TATGTGACAC	MPSS: TTAAGCCAAGTAA
ELAV1	SAGE: AAAAAGTTTT CACAGCGCC	MPSS:
ATTGCGAAAAAAA TAAGCTCCAGGA		
ELAVL2	SAGE: TTCAGTACAT	MPSS: TGAGTTGGATATA
ELAVL3	SAGE: CCAGAAAATA	MPSS: ACAGGCCCTGGCCG
ELAVL4	SAGE: TTAATCTTGC	MPSS: ATTTTAGTATGTT
EPRS	SAGE: AACGAAAAAA	MPSS: TTGAACCTGGTGC
ERAL1	SAGE: AACACTGACT	MPSS: TCTCCCACCCCAT
EWS	SAGE: GACCGAGGTG AGCAGAGGTG	MPSS: GTGGAGGCATGAG
EXOSC7	SAGE: TAAAATTTAA	MPSS: GCTAACACCCTCT
EXOS9	SAGE: AACAAACATC	MPSS: ATTTTGGAAACCAG
FBL	SAGE: CCGTGCTCGT	MPSS: TTTGCTGATGTGG
RBM44	SAGE: CATTTTTTAT	MPSS: TAAAGAAATTGTC
FMR1	SAGE: ATGTTTTTAA	MPSS: TTCCTCTGCTAAA
FUBP1	SAGE: TGAAGCAACT ATTCCAGCTA GTCCTCCAGC	MPSS:
TTTTGGCATACTT CAGCCAGTATAGA		
FUBP2	SAGE: TTGAGCCAGC	MPSS: TAAATCTCTGCCA
FUBP3	SAGE: GGGTTTTTAA	MPSS: ATTTTGTTCAT
FUS	SAGE: GGGGGTAACT	MPSS: AGGAAGGTAGAGA
ACCCAAGGGTTTT		
FXR1	SAGE: TTAGTCTTCA	MPSS: TTCCATTTGCTGA
TTCCATTAAAAAA		
FXR2	SAGE: GCCCTCACAG	MPSS: CAGACAGTAATCC
G3BP2	SAGE: GTGATTGTTT ATGGTGCCTT	MPSS:
ATTTCTTTGATGT		
GRSF1	SAGE: TCAGAACAGT	MPSS: TGATTTAAAAATA

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| HNRPH3 | SAGE: TCATAGTCAA AGTGGAGGTG | MPSS:
AAAATGAAACCTG AGTTGGAAGAATG |
| HNRPUL1 | SAGE: GCACCTCCTA | MPSS: ATACAGTGAGGCT |
| HNRPLL | SAGE: TTGCCTTGCT CCTCTGTTTC ATAGACTGTG | MPSS:
CTGTTCTGTAATA TGCTGACAAATTT |
| HNRPA0 | SAGE: TTGACGTTGA CCTGAAATTT AAGAGCGGGC
TCTAACATTT | MPSS: TTTGTAGCACCAT CCATGCCTGAAAT
CTCCCGGGGCGGC CATTCCGTACCGT |
HNRPCL1	SAGE: AACTCCCGTG	MPSS: TTCCAAGTCTGGA
HNRPD	SAGE: ATGTAGTAGT	MPSS: TAAGATGATTATT
HNRPDL	SAGE: TTAAACCTCA	MPSS: TAGAGGAGGATTT
HNRPF	SAGE: TTTGTTTATT	MPSS: TCCCTTATATTCT
TTTTACACCACAT		
HNRPG	SAGE: CCTATGTAAG TTTAATTTTA AGTTCTTCCA	MPSS:
AAATGTTTATGTA		
HNRPGT	SAGE: GAAAGGGGCT	MPSS: TTCCACAAAACAT
TGATAGAGGGGGA		
HNRPK	SAGE: TCCCTGGCAT	MPSS: ATTACCATTACAG
HNRPL	SAGE: TACTCTTGGC	MPSS: TTGTATTCCTTTT
HNRPM	SAGE: AAGCTGAGTG	MPSS: GCATGGCCACCGG
ACAGCGCCCCCAG		
SYNCRIP	SAGE: GTATCTTAAT	MPSS: ATGGAATCAAATA
HNRPR	SAGE: TAAGATTTCA	MPSS: TACATTCTTCAAA
CCATGAACATTGC		
HNRPU	SAGE: AATTGCATTA GAGTCAGCAT	MPSS:
AGACAAATGTGTC TATATGAAGTTAG		
HTF9C	SAGE: CTGGGGTGCT GCTTGCTGGC	MPSS:
TGGCCAGTCAAGG CTGGCCATCCGGA		
HTATSF1	SAGE: CAGAATAATA	MPSS: TGAAGAGGATAGT
IGF2BP2	SAGE: TACTGGTTTA	MPSS: ACCTGTGTATGTC
IF2B	SAGE: CCGATCACCG	MPSS: ACCGGACACAATC
EIF3S4	SAGE: CAGCCCAACC	MPSS: GTGTCCTGCCGCA
GTGCGCGTGCCAT		
EIF3S9	SAGE: GAAATAAAAG	MPSS: GTTTGAGTCAGTC
EIF4H	SAGE: CATCTAAACT	MPSS: TTTTAAATAAAAG
IGF2BP1	SAGE: TGTCACCCCT	MPSS: CTGGTCCCCAAAA
IGF2BP3	SAGE: TATAATGTGC	MPSS: ATTCCTCATGTAT
ILF3	SAGE: TGAAAGTAAC TGAAAGTAAA	MPSS:
AGCGGTTCTTTTT		
IREB1	SAGE: GATAGGTCGG	MPSS: CACATAAAAAGAA
IREB2	SAGE: GATGTAAATG	MPSS: ATTTGATTGGAAT
CCTCCTGGAACCTG		
JOSD2	SAGE: GACCCTGGGG	MPSS: CTGAACCTGCCCT
KHDRBS3	SAGE: CGTAAAAAGG	MPSS: AGGCAAGCACAGC
KHDRBS2	SAGE: TTAATGGAAA	MPSS: AGAATAGCTCCCA
KRR1	SAGE: AAATCAAGAG TGACATCATT	MPSS:
CCCTAGTTACTCC GATAAAGAATTGG |

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| LA | SAGE: ATGAAAATGG | MPSS: AGACCTGTAGAGA |
| LARP4 | SAGE: CATAAAATGT TTAGTAAATG TTCAGAGTTC | MPSS:
AGATGCTGTGCAG AGTGGCTGGGGTG TTCATAATAATTA |
| LARP5 | SAGE: GATTTAAGAG | MPSS: TCCTGGGCGGAGT |
| LARP7 | SAGE: CTAGATTTAA | MPSS: ACGAACAAGGTAT
ACGAACAAAGGTA |
| LKAP | SAGE: TAAAGAGTGG | MPSS: CACTTAACCTTGT |
| ADAD2 | SAGE: GGTTGTGCGG | MPSS: TACAAGCTGGTGG |
| LRPPRC | SAGE: ACTTATCCCG TTATTTTGAA TTACTIONTAA | MPSS:
CCTATAAAAAAAAA TGTTTCTAAAGCG |
| LSM5 | SAGE: ATTATCCTGG GTTATCCTGG | MPSS:
ACTTAGAAAATTC AGATTTTGCTAAA |
| MOV10L1 | SAGE: TGCTCAGCCT | MPSS: TGCAGTGGCTGAC
CGGCCTGGCACGT |
| MATR3 | SAGE: CACAAAAAAT | MPSS: TCTCCCGTTGAAA |
| RNMT | SAGE: GTTGTATAAT | MPSS: TTAGCTCATAAAA
CCAGGGCACTGTC |
MGC10433	SAGE: TGGAAGGACC	MPSS: CGTGACAAGCGCA
MGC27016	SAGE: TGTTAATTTT	MPSS: ACTATCTGTACTG
MINT	SAGE: GACTCGCAGA	MPSS: TCTTCCCCCAACT
MKI67IP	SAGE: GCGACTTTTT AATTAATAAC	MPSS:
TTTTCATATTTCT		
MSI1	SAGE: GTGTTTCGAA	MPSS: TTTGTGGGGGGGC
MSI2H	SAGE: CCTTATCTTC	MPSS: ACCAAAGCCAGGG
MTHFSD	SAGE: GAAAATTAAC	MPSS: TGGAATATGCCAT
MYEF2	SAGE: GAATACCAAG CCTCCTGTAC GAATGTGATT	MPSS:
CCACGATGTTAGA TCTAAATCTGCTG TATTACATGATTT		
SLC4A1AP	SAGE: ATCCTATTA	MPSS: CTATTAATAAAG
NCBP1	SAGE: TCAAGGTTTT	MPSS: TTCCTACAGCATC
NCBP2	SAGE: AGGACCAGCA	MPSS: TACATTGGGGCCT
NHP2L1	SAGE: TGTTCTCCAT TAGTATTTTC	MPSS:
ATGGAGACCTGGC CAATCCATTCAGC		
NKRF	SAGE: TGATTTCCAG TTCCGGTTGC	MPSS:
ATTTTGTATTAAT TTACTIONTATT		
NOL3	SAGE: GAGCAAGGGG CTGGATAGGA	MPSS:
CCAAACCTAGCCC GAGCCCCGACAAT		
NOL4	SAGE: AAGAGACAAT	MPSS: CAGCTGAAATATT
NOL8	SAGE: AAGCCTTTTT	MPSS: ACAGAACTCCAG
NONO	SAGE: CCCCCTGCTC	MPSS: TACCCTGCCTGTA
TGTTATACCCCAT		
NOP56	SAGE: TTTTCCTTCT	MPSS: TTGAAGAGACCGC
CTTGGGGCTGAAA		
NOVA1	SAGE: TGGTAGGGCA	MPSS: ATCTTAAGCTAAT
NOVA2	SAGE: GTTGCAGCTG	MPSS: TCCAAGAAGGGCG
NPM1	SAGE: CCTCTTCAGG	MPSS: AAAAGGACAAGAA
TCTGTATGGCAGT		
NCL	SAGE: GTTTTTGCTT TACAAAACCA	MPSS: AATGACAGAGCCT

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|
| NUFIP1 | SAGE: TGAAACAGTA | MPSS: CCCACACTGCCTT
CTCTTGGTGTTTT |
HRB	SAGE: TGTACAAAGT	MPSS: ATGTTTTGCTACA
NXF1	SAGE: CCTCAGAAGC	MPSS: CCTCTCCTCGCCT
NXF2	SAGE: ATCACCATTT	MPSS: ACCATTTTATTTT
PABP2	SAGE: AATAAAGTTG	MPSS: TCCGTTCCGCTCC
PABP3	SAGE: GTTCTGATCA	MPSS: ACAGAGATAATCA
PABP4	SAGE: CCAAGAAAGA	MPSS: ACGGGAATGCTGC
PABP5	SAGE: CTCATTGTAA TTAAATTAAC	MPSS:
ATGTTAAATTAAC		
PAIP1	SAGE: TAAATAGTTC CAACTTCAAC	MPSS:
CATACTTGGATGA		
C20orf119	SAGE: GCTGCCAAAA	MPSS: ACGGGCATGCTGC
PAPOLA	SAGE: TTGATCGAAG	MPSS: TACGAGATGACAG
PAPOLB	SAGE: TGATGTACCT	MPSS: AGCAATACTTTAA
PAPOLG	SAGE: AAAAGAGTCT TAATTCTTGT TAAAGAAAAA	MPSS:
ATTATTTACCCCA CTCGGGTAAATCC		
PCBP1	SAGE: AATTTTATTT	MPSS: CATATTAGTTTTA
PCBP2	SAGE: ATCCATCTGT	MPSS: CATCTGTGTAGTT
PCBP3	SAGE: GAGGATGTTG	MPSS: ACCATCACGGGGA
POLDIP3	SAGE: CCTCTATGTT	MPSS: TTGGGAGACCGGG
AGCTTGAAGAGGA		
SF3B14	SAGE: TGATCACCTA	MPSS: CACCAAAATAAAT
PNPT1	SAGE: AGGCAAGAGA	MPSS: AACGAAAGATTAA
PPP1R8	SAGE: CTTTATGTGT	MPSS: AAAAAGAAGGCTT
PPIE	SAGE: GGCAGGCTGT	MPSS: ATCGCCGACTGTG
PPIL4	SAGE: TGTAAGATGT CCAATCAAAA	MPSS:
TTTATGAAAGAGA TTATATTAGATGA		
PPRC1	SAGE: ACTCGATCAG	MPSS: CCCAGTGTTTGGA
TTGACTCCAACCG		
PPARGC1A	SAGE: CCACCCTTTT	MPSS: AGTACTCATTTAC
PPARGC1B	SAGE: GATTTTGACA	MPSS: CGTGCTAGCCTCA
PRKRA	SAGE: TAACAAAGAA TAAGCAAAGA TTTGTTGTTG	MPSS:
AACTTACTGAAAA		
PSMA1	SAGE: GTCTGCGTGC	MPSS: GCTGAGCGATACT
PSPC1	SAGE: AAATATTAAA	MPSS: CGACACAGAGAAC
PTBP1	SAGE: CAGCTGGGGC AGCCACCTGA	MPSS:
CCCTTTCCGTAAA CAGATGGGCTCCG		
PTBP2	SAGE: GCTCAGAATC	MPSS: TGACTTTAATAAAA
Q8TC92	SAGE: TGAACCTGGC	MPSS: CAAGCCTCAACAT
QKI	SAGE: TTGTTTGTA	MPSS: ATTTAAACTGAAC
RAD51AP1	SAGE: TTAGCTGTTA	MPSS: AATTTTAAATAAAA
RAE1	SAGE: GATTTCAACC	MPSS: AGCCCATCTCAGC
RAVR2	SAGE: AAATTTATAT	MPSS: TAATAAAAAGGGTT
RBM12B	SAGE: ATTGACTTTG	MPSS: TAAATGATAGGCC
RBM15B	SAGE: CTCTCCTTGC	MPSS: CAGAACATGACAA

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| RBM35A | SAGE: CCTTCTTGCT | MPSS: TTGGAATAATAAA
GAAAAGGAATTC |
| RBM35B | SAGE: GAATGTCCTT | MPSS: TTCTCTTCCTTAA
CCAATGTTCTACC |
| RBM10 | SAGE: TTGGGTGTCC | MPSS: ACGTCCTGTTTTG |
| RBM11 | SAGE: AAAATGAGCA TTGCCTAGCA | MPSS:
ATTCCCTTTTTAT |
| RBM12 | SAGE: ATATTTTCAA GAGCAGCTGG GTGGCCTTTG
GAGTCTAGAT AATACAGATT | MPSS: ACTAGATACTGCC CCACCTTTGCTCC
TTCATATTGCTGT |
| RBM14 | SAGE: TGGGCCTTCC | MPSS: TCGGTTTCCCCT |
| RBM15 | SAGE: GTTCCTGTTT | MPSS: ATTGTCCGTGGGT
ATTGTCCGTGGTG ATTGTCCGTGCAA |
| RBM16 | SAGE: GTAGGTAAAC | MPSS: CTGAACTTTATGA |
| RBM18 | SAGE: TTAGCATTTG AGTATAAATG | MPSS:
ATAACAAGAATGA |
RBM19	SAGE: GTGGACGCGT	MPSS: CTGGAGCAGCTGG
RBM22	SAGE: AAATTACTGT	MPSS: TTCTCTTTCCTCT
RBM23	SAGE: TGTGAGCCCT	MPSS: TGTCTGTGAGGC
RBM24	SAGE: ACTCTCCTGT	MPSS: TTCTAATCTATTC
RBM25	SAGE: TCTGAGTACT	MPSS: AGAATTTACCAG
RBM26	SAGE: AGTCCTTATG	MPSS: TTGTATTTTGAAA
RBM27	SAGE: CTCATAGAGG GTTTTTATGT GATAATCTCT	MPSS:
AACTTGATGTCTT CACAGCAATGACA		
RBM28	SAGE: ATGCTTCTTG	MPSS: CAAGCCTGCAACT
RBM33	SAGE: GAGAGATGAC	MPSS: TGTGTGAAGAATG
RBM34	SAGE: CGTTCTGTTA	MPSS: GGTTTTTGTGGGG
RBM38	SAGE: AGGTGAGAGG	MPSS: CACAGCATTCAGG
RBM39	SAGE: AACGAAAGAG GCAGGTGGTT	MPSS:
GAAGATTTAGAGG CCTGCATTTCCCTG TACTGTCATTTGT		
RBM3	SAGE: ATTATCCAGG	MPSS: ATCAAACCTGATTA
RBM41	SAGE: TGCTACAGGT	MPSS: TAGGATTTATGTA
RBM4	SAGE: TTCTCTTCTC	MPSS: ACTTTTTTTTCTTT
RBM5	SAGE: TGTAGCCTAT	MPSS: TTAGGGTGAACCA
RBM6	SAGE: GGATACAACC	MPSS: GTAAACTTGTTGA
RBM7	SAGE: TAATGGAGAT	MPSS: CTTTTTAATACCA
RBM8A	SAGE: CAGCCTTGGA AGGAAGCCAC ATAGTGATAG	MPSS:
TTCACGAGGCTGG TCGTGTATGTCCC AAATCTTAAATCA		
RBM9	SAGE: CCCTTGCCCC	MPSS: TCTTACGACAGAA
RBMS2	SAGE: GTGGTCGCAG	MPSS: AAAAAAATTTTC
RBMS3	SAGE: AATCATTCTA	MPSS: GCTAGCAAAAGCA
RBP56	SAGE: ATCCATAGTG	MPSS: CATAGTGAAATTG
RBPMS	SAGE: TTGAATTTGT	MPSS: CTGAAATTCCGCA
RED1	SAGE: ATGGGGGGTG	MPSS: TCCAACATAGAGG
RED2	SAGE: TACTGTGAGG	MPSS: TTCACATCACCTC
RENT1	SAGE: GCGGGGTGAC	MPSS: CGAACACCTCCAG
AGTGCCGGCCCTA |

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| RIB1 | SAGE: TGTTTTTTTT GAGGCCAGTG | MPSS:
GACCACATCCTGG |
| RKHD3 | SAGE: CTCTATAAAT GAGTGCGCCA | MPSS:
CTTTTTGAATGTT CGAGAGGAGCTTG |
| RPL12 | SAGE: ACATCATCGA | MPSS: ATTTGACAACTGA |
| RPL13 | SAGE: CTGGGTCTCC CCCGTCCGGA CTTGGTGCTG | MPSS:
CGAGGAGGCGGAA |
| RPL18 | SAGE: GGAGTGGACA | MPSS: CTA CTCTCTTATT |
| RPL22 | SAGE: TAGTAAGTCA GATGCTGCCA | MPSS:
ATATTTTCTATTA ACCGTGACATCCG |
| RPL28 | SAGE: TGCAGTAATG | MPSS: CATGTGCAGTAAT
CGCAAGAACAAGT |
| RPL34 | SAGE: TTTGATAATG TGTGCTAAAT | MPSS:
ACTAATATAGTAT AAGCGTGCTTTCC |
RPL3L	SAGE: GGCCCCAAA	MPSS: TCTTTGAGCAAAA
RPLP2	SAGE: GGATTTGGCC	MPSS: TTGGACAGCGTGG
MRPL3	SAGE: TTAAAGGCCG	MPSS: TCGGTAAAAATCT
MRPL12	SAGE: TGTCTGTGCC	MPSS: AAGGCGGCCCTGG
MRPL23	SAGE: CTCGAGGAGG	MPSS: TGTTTCCCGAGAA
RNASEL	SAGE: TCAATTA AAA AATCATTGTG CTGCAGGTTG	MPSS:
TG TAGTGATGGAG		
RNASE1	SAGE: TTAACCCCTC	MPSS: CAGGCTTTTCTGG
CAGGTCACCACCT CAGGTCAC TTTGT		
RNH1	SAGE: TGTTTATAAA	MPSS: AACAAAGAGGACT
RO60	SAGE: TCAATGGCCT	MPSS: ACACAAATAACAG
HNRPA2B1	SAGE: GTGTAATAAG	MPSS: ATGGTGT AATAAG
HNRPA3	SAGE: CCAGTGTGCA GAAATTGGTC	MPSS:
CCAAGTCCAGGAA ATTGTTAATTACT CATAGTCAGAAAA		
ROD1	SAGE: GTTTCTACAA	MPSS: TGTGGAAGAAGCA
RPL18A	SAGE: AAGGTGGAGG	MPSS: AAGTTCCCGCTGC
RPL7	SAGE: GAGGGTGTAG	MPSS: AAGCGCCTGAGAA
RPP29	SAGE: TTTTCTGAGT	MPSS: AAATAAAGGAGGT
RPS17	SAGE: GGCCGCGTTC	MPSS: CTGACACTAAGGA
RPS3	SAGE: CCCAGCCAG AGAATCACCC	MPSS:
ATGCTGCCCTGGG CTAGGTGGGTAAG		
RPS4Y1	SAGE: TGAAGGATGC	MPSS: ACCAACAGGGAAA
RPS5	SAGE: CCTTCGAGAT	MPSS: ATACACCTGCTCA
RPS9	SAGE: CCAGTGGCCC GCCATCTCAT	MPSS:
AGCGCTTTACAAA AGCGCTTTAAAAA		
MRPS15	SAGE: CTGAGGGTTCG	MPSS: CGCAGTTATGAAG
RTCD1	SAGE: GCATTAGCCA	MPSS: ATCGCTTGGTAAA
SNRP70	SAGE: GAGTCTGAGG	MPSS: GTGACCGTGACCG
GTGACCGTGCCG		
SNRPB2	SAGE: CTATGAAGAT	MPSS: ACCTATGCCAAGA
SAFB1	SAGE: CAGGGCGGGT	MPSS: ACAAGAGGTGGCA
SAFB2	SAGE: TTGTAGCTCA GACGAATGAT	MPSS:
TTTTTAAAAGCAA AAAATTGAAAAA |

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SAM68	SAGE: TGTAAGTCTG	MPSS: CTCATGAATTAAA
SART3	SAGE: GAGAGGACAG	MPSS: CAAGCTGCGTGAT
SCAND1	SAGE: ATCGGGCCCG	MPSS: GGGCCCGGGGGCC
SF1	SAGE: AAGTGATTCT	MPSS: ACCTGAGCTGCCT
SF3B4	SAGE: GATACACTGG	MPSS: TCCCATGGGTCAC
SFPQ	SAGE: AAAAAAAXXX CGTACTGAGC	MPSS:
TATCCTTTACTTG CCTATGGTTCAGG		
SFRS11	SAGE: GGCATTTTTG	MPSS: TCATTCCAGAGAA
CCAAGTTGAATCA		
SFRS12	SAGE: AGAAGGATCG ACAAGTCTGA GTGGCCGAAA	MPSS:
GAGACAAAAAAA TAAGAGCCCACAT TAACAGCTTTTCC		
SFRS15	SAGE: GCCCCTTTGT	MPSS: TTTTGGAAATAGG
SFRS1	SAGE: GAGCGGGATC	MPSS: AGATTACCAGGAA
SFRPS2B	SAGE: GTTTATCTTT	MPSS: ATATCTACATGTA
SFRPS2	SAGE: GTGTATCTTT	MPSS: ATATCAACATGTA
SFRPS3	SAGE: CATAACCTTC CTGTCAATTTG	MPSS:
CATCATGATGTAA ATACGAGGCATGT		
SFRS4	SAGE: GCCCACAGTA	MPSS: TAGGTCCCCTCA
SFRPS5	SAGE: AGACAAGCTG	MPSS: AGTTGACAGTGGC
SFRS6	SAGE: CCGTAGTGCC AAGACCCTCT	MPSS:
ATACTTGTAATT TGCTGGCAGGGTT		
SFRS7	SAGE: TAGCTAATAT CCTCGGAGAT	MPSS:
TTTATACTAATTG TGGATTACTATGT		
SFRPS9	SAGE: AGGGTGAAAC	MPSS: AACAGTTACTGTG
SIAHBP1	SAGE: GTGGACCCCA	MPSS: TTTGTGGAGTTTT
SLBP	SAGE: TAAAATTGCT	MPSS: CTCCAGCGGAAGA
SLTM	SAGE: TTCCCAACTA ATTGCTTTTC	MPSS:
ACAGCACTATCCT		
SNRPA	SAGE: CCTGCCCCTT	MPSS: TCCTTTGCCAAGA
SON	SAGE: TTAAATAGAG GAACGTCTTA TGTATTACAG	MPSS:
ATTTATTCAATAA TGCTGCCAGCCCG AGCACTAAGTCCT		
SPF45	SAGE: GCTGTTGCAT	MPSS: AGAAATGACAAAT
CCTTTTTTATTTT		
SR140	SAGE: AGCAACTGCA AAACAGATGA TTCTGAGCAA	MPSS:
TGCTGACAGTAGT AAAAAATCCTCC TTCAGATGCACAT		
SRP14	SAGE: TCTGTTTATC	MPSS: AGCACTGTGGTGA
AGCACTGTGGGAG		
SRP68	SAGE: GATTTGAAAT ATATTAAATC	MPSS:
CGACTCTATGACA		
SRP72	SAGE: AAGAAGATAT	MPSS: CAGAGAAAGCCAA
SRrp35	SAGE: GAAGACATTC	MPSS: TCGAAGTTATCGT
STAU1	SAGE: GAATGTAAGT	MPSS: AATAAACAGCTAC
TCCATGTAGTCCA		
STAU2	SAGE: TAGATTGTTT TTGGAATCCA TTCGGATTTT	MPSS:
TCAATTTATATAA AAAAATAAATAAT CAAGGAGATGCTC		
STRBP	SAGE: ACAAGCGCTT	MPSS: ACAAATAAACTGC
SUPV3L1	SAGE: CTGAAACAGC	MPSS: TCCAGAAAGAACT

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SURF6	SAGE: GATGGAAACT	MPSS: CGGGACCTTCAGA
GARS	SAGE: GGCTGATGTG	MPSS: TCTGAGCTGCCCA
HARS	SAGE: TGTGGGGCTC	MPSS: AAGGCTGAGCTGC
MARS	SAGE: CCCACGGTTA	MPSS: TAAAGAAACAGTT
SYNJ1	SAGE: CTCCTTTTAT	MPSS: TAAGGAAGGTGAA
SYNJ2	SAGE: ATACCTATTA ACTGAGCAGC	MPSS:
GTGGCTGTCAAAT ATGGACACAAAAC		
SYWC	SAGE: TGCAGGCCTG GAGGACTCCG	MPSS:
ACTCTGCCCTCTT GCAGAGCACCAGG		
TACC1	SAGE: TTTCATTGCC	MPSS: CAGGCATTTCTTT
TACC2	SAGE: GACAGGTTCT GGCAATGAGA	MPSS:
ATAACCAAGGAGA		
TARDBP	SAGE: TTTATGGTAC GACTGAGCTT	MPSS:
TTGAGTTGTTTTA ATGGTGTACACAGT		
TARB1	SAGE: TGCCTTAAAC	MPSS: TGGGAGTACACCA
TDRKH	SAGE: CTCAGCCCTG	MPSS: TATGTGCCTTCTC
CCTCCCAACCCT TACTTATATGATA		
Tenr	SAGE: TGAAATAGGC	MPSS: CTGGATTGTGAAA
TEP1	SAGE: GACTAGTGCC	MPSS: CATAATCCTTGCC
THOC4	SAGE: TAAAGGTTTT	MPSS: GCTCTGGTCGCAG
TIA1	SAGE: TGAAAATCAA GCAACAGGAA	MPSS:
CTTGTTTCAAACA TCAGCCCAGAAAT		
TIAR	SAGE: GTATAATAAG ATTTTTCTTT	MPSS:
AGAATGTTAAGAC AATCACCTTCTGC		
TNRC6A	SAGE: ACTTTTTGGC	MPSS: TCAATCACTGGAA
TNRC6C	SAGE: GCAACCAAAA	MPSS: TGCGAGAAAGCAC
TNRC4	SAGE: GCTATGCTTC	MPSS: ATTTTCTTTAGGA
TRA2A	SAGE: TGGGGAAAAG	MPSS: TGCATTTATCATG
TRA2B	SAGE: GATGAACTGA TATTTTTTTA	MPSS:
GAAGCCTGTGTAA		
TRBP2	SAGE: TATCTGCGCA	MPSS: CGTCCTCATTTTA
TRSPAP1	SAGE: ATGTGAGGGA	MPSS: TCTTTAAGGGCTG
CCTGCCATGATGT		
TUT1	SAGE: GCCCCTGAAG	MPSS: TCCATCATTTCTT
U2AF1	SAGE: CCATTTTTAC	MPSS: ACTGGCATGGCCG
U2AF1L4	SAGE: GAGACCTTCT	MPSS: TGGGCGATTCTGA
U2AF2	SAGE: CAGGGCCTGA	MPSS: TTTGTGGAGTTCA
HDLBP	SAGE: ACCTCAGGAA	MPSS: AAAAAGAACAGAA
YBOX1	SAGE: CGGAGACCCT GGGTTTTTAT	MPSS:
GGAGAGTGCTCCC CACCAGCTGAGAA		
YBOX2	SAGE: TGCCACCTGA	MPSS: TGTCATGTGCCAC
ZCRB1	SAGE: TTAATACTAT TATTGTCTTT	MPSS:
CTGCTCTTGACAG		
ZNF638	SAGE: GAGAAATATC	MPSS: TAAAAACCATGAT
+-----+-----+-----+-----+-----+

Supplementary file 7

RBPs	Tag freque	Tissue
A2BP1	3.35227	Retina
A2BP1	3.96944	Brain_cerebellum
A2BP1	6.40502	Brain_cortex
A2BP1	7.41468	Muscle
APOBEC3F	3.65063	Spinal_cord
APOBEC3F	37.1575	White_Blood_Cells
APOBEC3F	3.9056	Retina
APOBEC1	11.134	Muscle
ACF	11.1803	Retina
ACF	3.20251	Brain_cortex
ACF	3.48943	Vascular_endothelium
ACF	3.65063	Spinal_cord
ACF	5.45308	Stomach
ACF	6.07128	White_Blood_Cells
ACF	6.76376	Lung
ACF	8.34527	Prostate
ACIN1	15.488	White_Blood_Cells
ACIN1	21.4265	stem_cell
ACIN1	3.01623	Liver
ACIN1	3.20251	Brain_cortex
ACIN1	3.65063	Spinal_cord
ACIN1	3.88493	Brain_astrocyte
ACIN1	3.95946	Breast
ACIN1	5.56351	Prostate
ACIN1	7.01471	Vascular_endothelium
ADAT1	3.72675	Retina
AKAP1	10.4855	Breast
AKAP1	10.5447	stem_cell
AKAP1	11.6548	Brain_astrocyte
AKAP1	12.0391	Heart
AKAP1	15.8777	Brain_cerebellum
AKAP1	16.8699	Prostate
AKAP1	17.5368	Vascular_endothelium
AKAP1	22.268	Muscle
AKAP1	27.2654	Stomach
AKAP1	3.45012	Thyroid
AKAP1	4.03145	Colon
AKAP1	7.43149	White_Blood_Cells
AKAP1	7.59648	Retina
AKAP1	9.60753	Brain_cortex
ANKHD1	10.95189	Spinal_cord
ANKHD1	14.77667	Breast
ANKHD1	18.5367	Muscle
ANKHD1	21.88037	stem_cell
ANKHD1	28.99878	White_Blood_Cells
ANKHD1	3.01623	Liver
ANKHD1	31.72538	Prostate
ANKHD1	3.45012	Thyroid
ANKHD1	4.03145	Colon
ANKHD1	5.30507	Retina
ANKHD1	6.76376	Lung
ANKHD1	8.63353	Vascular_endothelium

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Supplementary file 7

RBPs	Tag freque	Tissue
ANKHD1	9.60753	Brain_cortex
ANKRD17	10.16234	Placenta
ANKRD17	11.8337	stem_cell
ANKRD17	20.6252	Breast
ANKRD17	3.01623	Liver
ANKRD17	3.96944	Brain_cerebellum
ANKRD17	5.45308	Stomach
ANKRD17	5.8584	Retina
ANKRD17	6.98503	Eye
ANKRD17	7.01472	Vascular_endothelium
ANKRD17	7.87355	White_Blood_Cells
ANKRD17	8.98387	Brain_cortex
ASCC1	3.62053	stem_cell
ASCC1	3.65063	Spinal_cord
ASCC1	3.6733	White_Blood_Cells
ASCC1	3.71133	Muscle
ASCC1	3.72675	Retina
ASCC1	3.87282	Vascular_endothelium
ASCC1	4.2448	Brain_cortex
ASCC1	5.57748	Prostate
ASCC1	9.63124	Heart
BRUNOL4	12.81	Brain_cortex
BRUNOL4	3.49516	Breast
BRUNOL4	3.70734	Muscle
BRUNOL4	3.9056	Retina
BRUNOL4	4.03145	Colon
BRUNOL4	7.22343	Heart
BRUNOL4	8.36622	Prostate
BRUNOL5	12.8602	Vascular_endothelium
BRUNOL5	19.2151	Brain_cortex
BRUNOL5	3.71133	Muscle
BRUNOL5	6.70455	Retina
BRUNOL5	7.76986	Brain_astrocyte
BRUNOL6	11.72993	Brain_cortex
BRUNOL6	20.11367	Retina
BRUNOL6	3.65063	Spinal_cord
BRUNOL6	4.03145	Colon
BRUNOL6	7.93888	Brain_cerebellum
CCBL2	3.20251	Brain_cortex
CCBL2	3.49516	Breast
CCBL2	3.96944	Brain_cerebellum
CCBL2	5.17518	Thyroid
CCBL2	5.27235	stem_cell
RCAN3	14.863	White_Blood_Cells
RCAN3	3.37399	Prostate
RCAN3	3.49516	Breast
RCAN3	5.45308	Stomach
RCAN3	5.95037	stem_cell
CDKN2AIP	19.7973	Breast
CDKN2AIP	3.20251	Brain_cortex
CDKN2AIP	3.37399	Prostate
CDKN2AIP	3.45012	Thyroid

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RBPs	Tag freque	Tissue
CDKN2AIP	3.50736	Vascular_endothelium
CDKN2AIP	3.79824	Retina
CDKN2AIP	3.87199	White_Blood_Cells
CDKN2AIP	3.96944	Brain_cerebellum
CDKN2AIP	4.48395	stem_cell
CDKN2AIP	4.50918	Lung
CDKN2AIP	4.81562	Heart
CDKN2AIP	5.08117	Placenta
CDKN2AIP	9.31337	Eye
CIRBP	10.3126	Breast
CIRBP	11.8561	Placenta
CIRBP	12.0754	Thyroid
CIRBP	13.6327	Stomach
CIRBP	20.1136	Retina
CIRBP	23.3096	Brain_astrocyte
CIRBP	28.5093	stem_cell
CIRBP	37.1133	Muscle
CIRBP	37.1575	White_Blood_Cells
CIRBP	39.6944	Brain_cerebellum
CIRBP	43.7104	Prostate
CIRBP	4.50918	Lung
CIRBP	51.2235	Eye
CIRBP	59.6251	Vascular_endothelium
CIRBP	7.22343	Heart
CIRBP	7.30127	Spinal_cord
CIRBP	73.6577	Brain_cortex
CIRBP	8.06289	Colon
CNOT4	3.37399	Prostate
CNOT4	3.50736	Vascular_endothelium
CNOT4	3.6733	White_Blood_Cells
CNOT4	3.95946	Breast
CNOT4	4.03145	Colon
COVA1	3.37399	Prostate
COVA1	3.79824	Retina
COVA1	3.93229	White_Blood_Cells
COVA1	4.93635	stem_cell
CPEB2	11.80785	Prostate
CPEB2	12.14256	White_Blood_Cells
CPEB2	15.43225	Vascular_endothelium
CPEB2	17.63966	stem_cell
CPEB2	3.01623	Liver
CPEB2	3.43755	Breast
CPEB2	3.79824	Retina
CPEB2	3.88493	Brain_astrocyte
CPEB2	3.96944	Brain_cerebellum
CPEB2	4.65669	Eye
CPEB2	5.08117	Placenta
CPEB2	7.30127	Spinal_cord
CPEB2	7.41468	Muscle
CPEB4	10.5221	Vascular_endothelium
CPEB4	3.49516	Breast
CPEB4	3.65063	Spinal_cord

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RBPs	Tag freque	Tissue
CPEB4	3.96944	Brain_cerebellum
CPEB4	4.03145	Colon
CPEB4	4.2448	Brain_cortex
CPEB4	7.74398	White_Blood_Cells
CPSF5	14.16632	White_Blood_Cells
CPSF5	3.01623	Liver
CPSF5	3.35227	Retina
CPSF5	3.45012	Thyroid
CPSF5	3.65063	Spinal_cord
CPSF5	3.71133	Muscle
CPSF5	3.93424	Placenta
CPSF5	3.96944	Brain_cerebellum
CPSF5	43.077	stem_cell
CPSF5	4.81562	Heart
CPSF5	6.39346	Breast
CPSF5	7.74564	Vascular_endothelium
CPSF5	8.68568	Prostate
CPSF6	10.3504	Thyroid
CPSF6	13.7502	Breast
CPSF6	23.8969	stem_cell
CPSF6	4.03145	Colon
CPSF6	4.2448	Brain_cortex
CPSF6	5.8584	Retina
CPSF6	6.24434	Prostate
CPSF6	8.09504	White_Blood_Cells
CSDC2	12.0391	Heart
CSDC2	3.65063	Spinal_cord
CSDC2	3.9056	Retina
CSDC2	3.96944	Brain_cerebellum
CSDC2	5.1441	Vascular_endothelium
CSDC2	6.40502	Brain_cortex
CSTF2	3.6733	White_Blood_Cells
CSTF2	3.88493	Brain_astrocyte
CSTF2	3.9056	Retina
CSTF2	3.96944	Brain_cerebellum
CSTF2	6.9421	stem_cell
CSTF2	6.97885	Vascular_endothelium
CSTF2	8.06289	Colon
CSTF3	3.37399	Prostate
CSTF3	3.38745	Placenta
CSTF3	3.43755	Breast
CSTF3	3.70734	Muscle
CSTF3	3.88493	Brain_astrocyte
CSTF3	3.95159	White_Blood_Cells
CSTF3	4.58266	stem_cell
CSTF3	6.3672	Brain_cortex
CSTF3	6.98503	Eye
CSTF3	7.93887	Brain_cerebellum
CSTF2T	11.53157	stem_cell
CSTF2T	13.6696	Retina
CSTF2T	18.09735	Liver
CSTF2T	3.37399	Prostate

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RBPs	Tag freque	Tissue
CSTF2T	35.13292	Placenta
CSTF2T	3.88493	Brain_astrocyte
CSTF2T	3.96944	Brain_cerebellum
CSTF2T	4.03145	Colon
CSTF2T	4.81562	Heart
CSTF2T	5.17518	Thyroid
CSTF2T	5.45308	Stomach
CSTF2T	7.30127	Spinal_cord
CSTF2T	7.41867	Muscle
CSTF2T	7.74563	Vascular_endothelium
CSTF2T	7.92817	White_Blood_Cells
CSTF2T	8.36021	Brain_cortex
CUGBP1	3.35227	Retina
CUGBP1	3.95159	White_Blood_Cells
CUGBP1	4.03145	Colon
CUGBP2	10.2882	Vascular_endothelium
CUGBP2	15.7471	White_Blood_Cells
CUGBP2	16.0126	Brain_cortex
CUGBP2	3.37399	Prostate
CUGBP2	3.68956	stem_cell
CUGBP2	3.88493	Brain_astrocyte
CUGBP2	3.96944	Brain_cerebellum
CUGBP2	6.76376	Lung
CUGBP2	7.41867	Muscle
DAZAP1	11.127	Prostate
DAZAP1	11.6185	Vascular_endothelium
DAZAP1	12.0943	Colon
DAZAP1	33.0597	White_Blood_Cells
DAZAP1	3.45012	Thyroid
DAZAP1	3.71133	Muscle
DAZAP1	45.7748	stem_cell
DAZAP1	6.40502	Brain_cortex
DAZAP1	6.98503	Eye
DAZAP1	7.76986	Brain_astrocyte
DAZAP1	7.91891	Breast
DDX17	15.193	Retina
DDX17	27.1945	Brain_astrocyte
DDX17	3.37399	Prostate
DDX17	3.45012	Thyroid
DDX17	3.71575	White_Blood_Cells
DDX17	3.96944	Brain_cerebellum
DDX17	4.65669	Eye
DDX17	5.90104	Placenta
DDX17	6.3672	Brain_cortex
DDX17	6.8751	Breast
DDX17	7.01471	Vascular_endothelium
DDX17	7.90852	stem_cell
DDX53	3.37399	Prostate
DDX53	3.65063	Spinal_cord
DDX53	4.00156	White_Blood_Cells
DDX53	4.50918	Lung
DDX53	7.37912	stem_cell

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RBPs	Tag freque	Tissue
DDX53	7.76986	Brain_astrocyte
DGCR8	10.81721	Breast
DGCR8	11.616	White_Blood_Cells
DGCR8	13.40953	stem_cell
DGCR8	23.6042	Placenta
DGCR8	3.01623	Liver
DGCR8	3.37399	Prostate
DGCR8	3.71133	Muscle
DGCR8	3.88493	Brain_astrocyte
DGCR8	3.96944	Brain_cerebellum
DGCR8	5.1577	Brain_cortex
DGCR8	5.8584	Retina
DGCR8	7.71615	Vascular_endothelium
DHX30	10.1188	White_Blood_Cells
DHX30	10.4855	Breast
DHX30	10.5221	Vascular_endothelium
DHX30	11.122	Muscle
DHX30	11.2729	Lung
DHX30	11.6548	Brain_astrocyte
DHX30	13.496	Prostate
DHX30	13.9701	Eye
DHX30	15.6836	Placenta
DHX30	17.2506	Thyroid
DHX30	19.2151	Brain_cortex
DHX30	20.2926	stem_cell
DHX30	27.786	Brain_cerebellum
DHX30	5.45308	Stomach
DHX30	6.03245	Liver
DHX30	7.30127	Spinal_cord
DHX30	7.59648	Retina
DHX30	9.63124	Heart
DHX9	10.9062	Stomach
DHX9	12.0754	Thyroid
DHX9	12.8602	Vascular_endothelium
DHX9	13.496	Prostate
DHX9	25.7131	White_Blood_Cells
DHX9	27.9613	Breast
DHX9	3.01623	Liver
DHX9	3.71133	Muscle
DHX9	3.88493	Brain_astrocyte
DHX9	4.03145	Colon
DHX9	41.8502	stem_cell
DHX9	4.65669	Eye
DHX9	5.1577	Brain_cortex
DHX9	5.8584	Retina
DHX9	7.30127	Spinal_cord
DHX9	9.01835	Lung
DICER	10.3154	Brain_cortex
DICER	13.5498	Placenta
DICER	3.71133	Muscle
DICER	4.81562	Heart
DICER	5.17518	Thyroid

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RBPs	Tag freque	Tissue
DICER	5.92739	White_Blood_Cells
DICER	6.59848	stem_cell
DICER	7.01471	Vascular_endothelium
DICER	7.30127	Spinal_cord
DICER	7.8112	Retina
DICER	7.91891	Breast
DICER	8.17962	Stomach
DICER	9.01835	Lung
DICER	9.31337	Eye
DNAJC17	10.1188	White_Blood_Cells
DNAJC17	15.5397	Brain_astrocyte
DNAJC17	24.426	Vascular_endothelium
DNAJC17	3.70734	Muscle
DNAJC17	3.79824	Retina
DNAJC17	4.48104	Placenta
DNAJC17	4.65669	Eye
DNAJC17	5.48038	stem_cell
DNAJC17	6.24434	Prostate
DNAJC17	6.90024	Thyroid
DNAJC17	8.06289	Colon
DNAJC17	9.60753	Brain_cortex
DKC1	11.127	Prostate
DKC1	20.6253	Breast
DKC1	25.4513	stem_cell
DKC1	3.20251	Brain_cortex
DKC1	3.71133	Muscle
DKC1	3.88493	Brain_astrocyte
DKC1	3.93403	Placenta
DKC1	6.90024	Thyroid
DKC1	7.30127	Spinal_cord
DKC1	7.74398	White_Blood_Cells
DKC1	7.74563	Vascular_endothelium
DKC1	9.49559	Retina
DKFZp434.	3.48751	stem_cell
DKFZp434.	3.95946	Breast
DKFZp434.	4.03145	Colon
DKFZp434.	5.57748	Prostate
DKFZp686f	17.25058	Thyroid
DKFZp686f	18.04128	stem_cell
DKFZp686f	18.63321	Vascular_endothelium
DKFZp686f	19.08582	Stomach
DKFZp686f	21.66911	Breast
DKFZp686f	22.56675	White_Blood_Cells
DKFZp686f	3.01623	Liver
DKFZp686f	3.12217	Prostate
DKFZp686f	3.71133	Muscle
DKFZp686f	3.79824	Retina
DKFZp686f	3.96944	Brain_cerebellum
DKFZp686f	56.4402	Colon
DKFZp686f	5.90103	Placenta
DKFZp686f	6.98503	Eye
DND1	3.49516	Breast

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RBPs	Tag freque	Tissue
DND1	3.88493	Brain_astrocyte
DND1	4.95864	stem_cell
DND1	6.97885	Vascular_endothelium
DND1	7.43149	White_Blood_Cells
DRB1	27.146	Liver
DRB1	3.68956	stem_cell
DRB1	3.71133	Muscle
DRB1	3.88493	Brain_astrocyte
DRB1	4.04752	White_Blood_Cells
DRB1	8.34527	Prostate
DSRAD	11.6548	Brain_astrocyte
DSRAD	19.2151	Brain_cortex
DSRAD	20.2439	Prostate
DSRAD	24.5515	Vascular_endothelium
DSRAD	27.516	stem_cell
DSRAD	3.49516	Breast
DSRAD	3.65063	Spinal_cord
DSRAD	39.3229	White_Blood_Cells
DSRAD	3.96944	Brain_cerebellum
DSRAD	4.03145	Colon
DSRAD	4.65669	Eye
DSRAD	4.81562	Heart
DSRAD	5.45308	Stomach
DSRAD	5.90104	Placenta
DSRAD	6.90024	Thyroid
DSRAD	7.41468	Muscle
DSRAD	9.04868	Liver
DUSP11	3.50736	Vascular_endothelium
DUSP11	3.93229	White_Blood_Cells
DUSP11	4.606092	stem_cell
DUSP11	5.57748	Prostate
DUSP11	5.67955	Retina
DUSP11	6.76376	Lung
DUSP11	6.93271	Breast
DUS2L	15.488	White_Blood_Cells
DUS2L	3.37399	Prostate
DUS2L	3.9056	Retina
DUS2L	4.03145	Colon
DUS2L	5.1441	Vascular_endothelium
DUS2L	7.34039	stem_cell
DUS2L	9.31337	Eye
E2AK2	3.20251	Brain_cortex
E2AK2	3.88493	Brain_astrocyte
E2AK2	4.03145	Colon
E2AK2	6.97885	Vascular_endothelium
E2AK2	8.09504	White_Blood_Cells
ELAV1	11.616	White_Blood_Cells
ELAV1	11.65479	Brain_astrocyte
ELAV1	11.9083	Brain_cerebellum
ELAV1	12.0649	Liver
ELAV1	13.97004	Eye
ELAV1	18.44778	stem_cell

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RBPs	Tag freque	Tissue
ELAV1	19.65166	Prostate
ELAV1	21.04416	Vascular_endothelium
ELAV1	3.66073	Placenta
ELAV1	3.70734	Muscle
ELAV1	3.95946	Breast
ELAV1	4.50918	Lung
ELAV1	4.81562	Heart
ELAV1	7.30127	Spinal_cord
ELAV1	8.4896	Brain_cortex
ELAV1	9.4956	Retina
ELAVL2	15.4731	Brain_cortex
ELAVL2	15.6224	Retina
ELAVL2	3.87282	Vascular_endothelium
ELAVL2	3.96692	stem_cell
ELAVL2	3.96944	Brain_cerebellum
ELAVL2	4.65669	Eye
ELAVL2	7.30127	Spinal_cord
ELAVL3	23.8166	Brain_cerebellum
ELAVL3	29.2051	Spinal_cord
ELAVL3	3.37399	Prostate
ELAVL3	39.056	Retina
ELAVL3	41.2616	Brain_cortex
ELAVL3	5.53434	stem_cell
ELAVL3	6.98503	Eye
ELAVL3	7.71615	Vascular_endothelium
ELAVL4	5.8584	Retina
ELAVL4	7.73655	Brain_cortex
EPRS	3.12217	Prostate
ERAL1	11.122	Muscle
ERAL1	3.38745	Placenta
ERAL1	3.87282	Vascular_endothelium
ERAL1	3.96944	Brain_cerebellum
ERAL1	4.04752	White_Blood_Cells
ERAL1	4.81562	Heart
ERAL1	6.23641	stem_cell
ERAL1	6.24434	Prostate
ERAL1	7.76986	Brain_astrocyte
ERAL1	8.4896	Brain_cortex
EWS	10.0568	Retina
EWS	11.122	Muscle
EWS	15.5397	Brain_astrocyte
EWS	16.1258	Colon
EWS	20.971	Breast
EWS	21.6371	Placenta
EWS	24.80942	Vascular_endothelium
EWS	31.6127	White_Blood_Cells
EWS	32.66403	stem_cell
EWS	3.96944	Brain_cerebellum
EWS	4.50918	Lung
EWS	8.17962	Stomach
EWS	8.98387	Brain_cortex
EWS	9.31337	Eye

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RBPs	Tag freque	Tissue
EWS	9.36651	Prostate
EXOSC7	10.1623	Placenta
EXOSC7	15.7821	Lung
EXOSC7	17.2506	Thyroid
EXOSC7	3.65063	Spinal_cord
EXOSC7	3.70734	Muscle
EXOSC7	47.132	stem_cell
EXOSC7	5.1441	Vascular_endothelium
EXOSC7	5.1577	Brain_cortex
EXOSC7	58.4169	Prostate
EXOSC7	6.07128	White_Blood_Cells
EXOSC7	8.6949	Breast
EXOSC7	9.49559	Retina
EXOSC7	9.63124	Heart
EXOS9	3.87282	Vascular_endothelium
EXOS9	4.03145	Colon
EXOS9	7.97146	stem_cell
FBL	11.2026	Placenta
FBL	11.6548	Brain_astrocyte
FBL	14.4469	Heart
FBL	17.2506	Thyroid
FBL	21.8123	Stomach
FBL	22.2945	White_Blood_Cells
FBL	22.4176	Brain_cortex
FBL	24.9774	Prostate
FBL	34.9251	Eye
FBL	3.65063	Spinal_cord
FBL	3.70734	Muscle
FBL	3.96944	Brain_cerebellum
FBL	4.03145	Colon
FBL	48.1257	Breast
FBL	51.441	Vascular_endothelium
FBL	6.70455	Retina
FBL	89.0649	stem_cell
FBL	9.01835	Lung
FBL	9.04868	Liver
RBM44	3.38745	Placenta
RBM44	3.71575	White_Blood_Cells
RBM44	3.72675	Retina
FMR1	11.0687	stem_cell
FMR1	11.2729	Lung
FMR1	12.0649	Liver
FMR1	14.6025	Spinal_cord
FMR1	18.0519	Brain_cortex
FMR1	18.631	Placenta
FMR1	3.37399	Prostate
FMR1	3.48943	Vascular_endothelium
FMR1	3.49516	Breast
FMR1	3.96944	Brain_cerebellum
FMR1	5.45308	Stomach
FMR1	5.8584	Retina
FMR1	6.90024	Thyroid

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RBPs	Tag freque	Tissue
FMR1	7.43149	White_Blood_Cells
FUBP1	10.4855	Breast
FUBP1	11.64171	Eye
FUBP1	12.0047	White_Blood_Cells
FUBP1	14.59894	stem_cell
FUBP1	17.73273	Prostate
FUBP1	18.63316	Vascular_endothelium
FUBP1	3.96944	Brain_cerebellum
FUBP1	6.40502	Brain_cortex
FUBP1	7.07902	Retina
FUBP1	7.76986	Brain_astrocyte
FUBP2	11.134	Muscle
FUBP2	14.6025	Spinal_cord
FUBP2	15.0811	Liver
FUBP2	15.6109	Prostate
FUBP2	19.6701	Placenta
FUBP2	19.8472	Brain_cerebellum
FUBP2	20.1572	Colon
FUBP2	23.2834	Eye
FUBP2	28.8226	Brain_cortex
FUBP2	30.1705	Retina
FUBP2	34.9644	Brain_astrocyte
FUBP2	37.3567	White_Blood_Cells
FUBP2	4.50918	Lung
FUBP2	61.9651	Vascular_endothelium
FUBP2	65.3134	Breast
FUBP2	7.22343	Heart
FUBP2	73.045	stem_cell
FUBP2	8.6253	Thyroid
FUBP2	89.9759	Stomach
FUBP3	10.9062	Stomach
FUBP3	11.8784	Breast
FUBP3	12.0649	Liver
FUBP3	12.8602	Vascular_endothelium
FUBP3	12.8942	Brain_cortex
FUBP3	14.6025	Spinal_cord
FUBP3	15.7821	Lung
FUBP3	16.2984	Eye
FUBP3	16.8699	Prostate
FUBP3	22.0184	Placenta
FUBP3	22.4258	Thyroid
FUBP3	23.3096	Brain_astrocyte
FUBP3	24.6885	Retina
FUBP3	26.3617	stem_cell
FUBP3	4.03145	Colon
FUBP3	7.42267	Muscle
FUBP3	9.63124	Heart
FUBP3	9.87898	White_Blood_Cells
FUS	11.122	Muscle
FUS	11.2729	Lung
FUS	11.7168	Retina
FUS	119.25	Vascular_endothelium

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RBPs	Tag freque	Tissue
FUS	16.8547	Heart
FUS	20.9551	Eye
FUS	22.4176	Brain_cortex
FUS	23.3096	Brain_astrocyte
FUS	23.6042	Placenta
FUS	24.9774	Prostate
FUS	25.5544	Spinal_cord
FUS	3.01623	Liver
FUS	4.03145	Colon
FUS	47.6332	Brain_cerebellum
FUS	63.3513	Breast
FUS	6.90024	Thyroid
FUS	81.7464	White_Blood_Cells
FUS	91.361	stem_cell
FXR1	10.4683	Vascular_endothelium
FXR1	12.0754	Thyroid
FXR1	15.193	Retina
FXR1	18.5567	Muscle
FXR1	19.7713	stem_cell
FXR1	21.9038	Spinal_cord
FXR1	3.01623	Liver
FXR1	3.87199	White_Blood_Cells
FXR1	3.88493	Brain_astrocyte
FXR1	4.50918	Lung
FXR1	4.65669	Eye
FXR1	4.81562	Heart
FXR1	6.7749	Placenta
FXR1	8.6949	Breast
FXR1	9.36651	Prostate
FXR1	9.60753	Brain_cortex
FXR2	11.134	Muscle
FXR2	12.0943	Colon
FXR2	12.4887	Prostate
FXR2	15.5397	Brain_astrocyte
FXR2	15.8777	Brain_cerebellum
FXR2	3.45012	Thyroid
FXR2	3.79824	Retina
FXR2	4.81562	Heart
FXR2	6.98503	Eye
FXR2	7.01471	Vascular_endothelium
FXR2	7.90319	White_Blood_Cells
FXR2	7.91891	Breast
FXR2	8.90916	stem_cell
FXR2	9.60753	Brain_cortex
G3BP2	11.134	Muscle
G3BP2	11.64171	Eye
G3BP2	14.6025	Spinal_cord
G3BP2	15.08836	Breast
G3BP2	15.53972	Brain_astrocyte
G3BP2	15.8777	Brain_cerebellum
G3BP2	16.8699	Prostate
G3BP2	17.2506	Thyroid

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RBPs	Tag freque	Tissue
G3BP2	18.135	Brain_cortex
G3BP2	18.3665	White_Blood_Cells
G3BP2	20.15725	Colon
G3BP2	20.6354	Vascular_endothelium
G3BP2	3.01623	Liver
G3BP2	5.45308	Stomach
G3BP2	55.85055	stem_cell
G3BP2	6.76377	Lung
G3BP2	8.46862	Placenta
G3BP2	9.63124	Heart
GRSF1	10.9519	Spinal_cord
GRSF1	14.8453	Muscle
GRSF1	16.4976	stem_cell
GRSF1	19.1016	Brain_cortex
GRSF1	3.43755	Breast
GRSF1	3.45012	Thyroid
GRSF1	3.88493	Brain_astrocyte
GRSF1	4.03145	Colon
GRSF1	7.74563	Vascular_endothelium
GRSF1	7.90319	White_Blood_Cells
GRSF1	7.93887	Brain_cerebellum
GRSF1	8.36622	Prostate
GRSF1	9.31337	Eye
HNRPH3	11.12601	Muscle
HNRPH3	12.0943	Colon
HNRPH3	12.40191	Retina
HNRPH3	15.53973	Brain_astrocyte
HNRPH3	15.78211	Lung
HNRPH3	15.8777	Brain_cerebellum
HNRPH3	17.25062	Thyroid
HNRPH3	18.25317	Spinal_cord
HNRPH3	18.34855	Brain_cortex
HNRPH3	23.71212	Placenta
HNRPH3	25.61177	Eye
HNRPH3	29.61041	Prostate
HNRPH3	32.71852	Stomach
HNRPH3	34.4039	White_Blood_Cells
HNRPH3	39.9699	Vascular_endothelium
HNRPH3	47.4511	stem_cell
HNRPH3	59.3025	Breast
HNRPH3	6.03246	Liver
HNRPH3	9.63124	Heart
HNRPUL1	10.122	Prostate
HNRPUL1	13.1809	stem_cell
HNRPUL1	13.6696	Retina
HNRPUL1	13.9577	Vascular_endothelium
HNRPUL1	21.8123	Stomach
HNRPUL1	3.49516	Breast
HNRPUL1	3.88493	Brain_astrocyte
HNRPUL1	4.65669	Eye
HNRPUL1	5.17518	Thyroid
HNRPUL1	5.90104	Placenta

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RBPs	Tag freque	Tissue
HNRPUL1	6.3672	Brain_cortex
HNRPUL1	6.76376	Lung
HNRPUL1	7.22343	Heart
HNRPUL1	7.30127	Spinal_cord
HNRPUL1	7.43149	White_Blood_Cells
HNRPLL	10.37026	Breast
HNRPLL	10.98221	Placenta
HNRPLL	11.27294	Lung
HNRPLL	11.61845	Vascular_endothelium
HNRPLL	14.3369	Retina
HNRPLL	16.35924	Stomach
HNRPLL	18.97564	Thyroid
HNRPLL	19.68645	stem_cell
HNRPLL	3.12217	Prostate
HNRPLL	3.65063	Spinal_cord
HNRPLL	3.71133	Muscle
HNRPLL	3.96944	Brain_cerebellum
HNRPLL	7.2801	Brain_cortex
HNRPLL	7.67486	White_Blood_Cells
HNRPLL	7.76986	Brain_astrocyte
HNRPA0	15.75416	Retina
HNRPA0	18.0367	Lung
HNRPA0	18.25313	Spinal_cord
HNRPA0	18.62678	Eye
HNRPA0	18.97566	Thyroid
HNRPA0	19.42466	Brain_astrocyte
HNRPA0	19.92293	Brain_cortex
HNRPA0	21.81228	Stomach
HNRPA0	23.11433	Prostate
HNRPA0	3.71133	Muscle
HNRPA0	45.13044	Placenta
HNRPA0	54.6362	White_Blood_Cells
HNRPA0	71.21174	stem_cell
HNRPA0	7.93888	Brain_cerebellum
HNRPA0	91.05719	Breast
HNRPA0	93.76325	Vascular_endothelium
HNRPA0	9.63124	Heart
HNRPCL1	3.50736	Vascular_endothelium
HNRPCL1	3.62053	stem_cell
HNRPCL1	3.88493	Brain_astrocyte
HNRPCL1	4.65669	Eye
HNRPCL1	6.24434	Prostate
HNRPCL1	6.99032	Breast
HNRPD	116.054	Placenta
HNRPD	14.8294	Muscle
HNRPD	224.075	stem_cell
HNRPD	23.3096	Brain_astrocyte
HNRPD	23.8166	Brain_cerebellum
HNRPD	25.6118	Eye
HNRPD	25.7885	Brain_cortex
HNRPD	26.4859	Heart
HNRPD	29.326	Thyroid

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RBPs	Tag freque	Tissue
HNRPD	30.1623	Liver
HNRPD	30.9379	Breast
HNRPD	31.2217	Prostate
HNRPD	34.8554	Vascular_endothelium
HNRPD	46.4639	White_Blood_Cells
HNRPD	48.82	Retina
HNRPD	49.6009	Lung
HNRPD	65.7114	Spinal_cord
HNRPD	95.429	Stomach
HNRPDL	119.616	Prostate
HNRPDL	126.257	Lung
HNRPDL	131.529	stem_cell
HNRPDL	144.333	Retina
HNRPDL	146.025	Spinal_cord
HNRPDL	179.695	Breast
HNRPDL	18.6267	Eye
HNRPDL	27.1945	Brain_astrocyte
HNRPDL	27.786	Brain_cerebellum
HNRPDL	29.6907	Muscle
HNRPDL	38.525	Heart
HNRPDL	40.8981	Stomach
HNRPDL	54.013	Vascular_endothelium
HNRPDL	73.5678	White_Blood_Cells
HNRPDL	78.4219	Liver
HNRPDL	79.3528	Thyroid
HNRPDL	8.06289	Colon
HNRPDL	93.1548	Placenta
HNRPDL	97.9962	Brain_cortex
HNRPF	11.6417	Eye
HNRPF	16.1258	Colon
HNRPF	19.8472	Brain_cerebellum
HNRPF	20.5764	Vascular_endothelium
HNRPF	20.7007	Thyroid
HNRPF	22.0086	White_Blood_Cells
HNRPF	24.1298	Liver
HNRPF	24.4661	Breast
HNRPF	24.8005	Lung
HNRPF	26.8862	Placenta
HNRPF	34.9644	Brain_astrocyte
HNRPF	37.466	Prostate
HNRPF	4.81562	Heart
HNRPF	60.8777	stem_cell
HNRPF	6.70455	Retina
HNRPF	7.30127	Spinal_cord
HNRPF	7.42267	Muscle
HNRPF	8.17962	Stomach
HNRPG	10.3154	Brain_cortex
HNRPG	11.2026	Placenta
HNRPG	11.27294	Lung
HNRPG	11.39472	Retina
HNRPG	12.19006	Breast
HNRPG	18.97568	Thyroid

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RBPs	Tag freque	Tissue
HNRPG	19.42463	Brain_astrocyte
HNRPG	20.83395	Prostate
HNRPG	26.37884	Vascular_endothelium
HNRPG	30.42235	White_Blood_Cells
HNRPG	3.71133	Muscle
HNRPG	3.96944	Brain_cerebellum
HNRPG	4.03145	Colon
HNRPG	7.30127	Spinal_cord
HNRPG	84.29059	stem_cell
HNRPG	9.31337	Eye
HNRPGT	3.65063	Spinal_cord
HNRPK	110.135	Breast
HNRPK	11.9083	Brain_cerebellum
HNRPK	12.0943	Colon
HNRPK	128.602	Vascular_endothelium
HNRPK	12.8942	Brain_cortex
HNRPK	13.8005	Thyroid
HNRPK	165.096	stem_cell
HNRPK	24.1298	Liver
HNRPK	25.5544	Spinal_cord
HNRPK	27.1945	Brain_astrocyte
HNRPK	29.3096	Lung
HNRPK	30.3659	Prostate
HNRPK	31.4722	Placenta
HNRPK	40.9328	Heart
HNRPK	42.9616	Retina
HNRPK	5.45308	Stomach
HNRPK	6.98503	Eye
HNRPK	88.1591	White_Blood_Cells
HNRPL	151.58	stem_cell
HNRPL	29.3096	Lung
HNRPL	29.326	Thyroid
HNRPL	29.6587	Muscle
HNRPL	33.1785	Liver
HNRPL	36.5063	Spinal_cord
HNRPL	38.525	Heart
HNRPL	40.3145	Colon
HNRPL	43.8404	Brain_cortex
HNRPL	44.2385	Eye
HNRPL	46.9318	Retina
HNRPL	47.6332	Brain_cerebellum
HNRPL	58.2739	Brain_astrocyte
HNRPL	62.8097	Vascular_endothelium
HNRPL	81.7464	White_Blood_Cells
HNRPL	89.9759	Stomach
HNRPL	93.6651	Prostate
HNRPL	94.4166	Placenta
HNRPL	95.6439	Breast
HNRPM	12.0391	Heart
HNRPM	14.6025	Spinal_cord
HNRPM	14.8453	Muscle
HNRPM	15.193	Retina

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RBPs	Tag freque	Tissue
HNRPM	16.8699	Prostate
HNRPM	20.1572	Colon
HNRPM	20.1647	Placenta
HNRPM	22.5459	Lung
HNRPM	24.4661	Breast
HNRPM	24.5389	Stomach
HNRPM	25.6201	Brain_cortex
HNRPM	25.8759	Thyroid
HNRPM	3.01623	Liver
HNRPM	32.0125	White_Blood_Cells
HNRPM	41.8731	Vascular_endothelium
HNRPM	46.5669	Eye
HNRPM	46.6191	Brain_astrocyte
HNRPM	66.2274	stem_cell
HNRPM	7.93887	Brain_cerebellum
SYNCRIP	12.0754	Thyroid
SYNCRIP	13.5498	Placenta
SYNCRIP	19.8346	stem_cell
SYNCRIP	24.4661	Breast
SYNCRIP	3.01623	Liver
SYNCRIP	3.35227	Retina
SYNCRIP	3.93229	White_Blood_Cells
SYNCRIP	4.50918	Lung
SYNCRIP	7.30127	Spinal_cord
SYNCRIP	8.36622	Prostate
HNRPR	15.4913	Vascular_endothelium
HNRPR	16.8547	Heart
HNRPR	18.6267	Eye
HNRPR	19.0858	Stomach
HNRPR	21.1136	Liver
HNRPR	22.3099	Prostate
HNRPR	25.8759	Thyroid
HNRPR	31.5642	Lung
HNRPR	36.5063	Spinal_cord
HNRPR	3.70734	Muscle
HNRPR	38.6827	Brain_cortex
HNRPR	39.056	Retina
HNRPR	42.5919	White_Blood_Cells
HNRPR	45.7305	Placenta
HNRPR	48.1257	Breast
HNRPR	62.7225	stem_cell
HNRPR	7.76986	Brain_astrocyte
HNRPR	7.93887	Brain_cerebellum
HNRPU	10.9062	Stomach
HNRPU	11.65479	Brain_astrocyte
HNRPU	12.03905	Heart
HNRPU	13.97006	Eye
HNRPU	15.7697	Brain_cortex
HNRPU	17.0921	Retina
HNRPU	18.25014	Placenta
HNRPU	18.25313	Spinal_cord
HNRPU	18.97566	Thyroid

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RBPs	Tag freque	Tissue
HNRPU	22.6107	Prostate
HNRPU	23.37715	White_Blood_Cells
HNRPU	24.80049	Lung
HNRPU	4.03145	Colon
HNRPU	42.1611	Breast
HNRPU	43.725	Vascular_endothelium
HNRPU	6.03245	Liver
HNRPU	7.41468	Muscle
HNRPU	78.8234	stem_cell
HNRPU	7.93887	Brain_cerebellum
HTF9C	12.78692	Breast
HTF9C	13.85233	Brain_cortex
HTF9C	14.83735	Muscle
HTF9C	3.12217	Prostate
HTF9C	3.88493	Brain_astrocyte
HTF9C	3.9056	Retina
HTF9C	4.65668	Eye
HTF9C	5.45308	Stomach
HTF9C	7.04818	Placenta
HTF9C	7.30126	Spinal_cord
HTF9C	7.74564	Vascular_endothelium
HTF9C	7.92219	White_Blood_Cells
HTF9C	8.0629	Colon
HTF9C	9.03506	stem_cell
HTATSF1	10.9062	Stomach
HTATSF1	13.8005	Thyroid
HTATSF1	14.8453	Muscle
HTATSF1	15.4731	Brain_cortex
HTATSF1	19.6701	Placenta
HTATSF1	20.8903	Retina
HTATSF1	23.9821	stem_cell
HTATSF1	24.8005	Lung
HTATSF1	3.65063	Spinal_cord
HTATSF1	4.00156	White_Blood_Cells
HTATSF1	4.65669	Eye
HTATSF1	5.7966	Breast
HTATSF1	6.03245	Liver
HTATSF1	6.74798	Prostate
HTATSF1	7.71615	Vascular_endothelium
IF2B	11.6417	Eye
IF2B	13.8005	Thyroid
IF2B	15.8064	White_Blood_Cells
IF2B	16.8547	Heart
IF2B	19.2151	Brain_cortex
IGF2BP2	10.4683	Vascular_endothelium
IGF2BP2	22.0184	Placenta
IGF2BP2	29.992	Stomach
IGF2BP2	3.01623	Liver
IGF2BP2	3.20251	Brain_cortex
IGF2BP2	3.45012	Thyroid
IGF2BP2	3.49516	Breast
IF2B	23.7567	Breast

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RBPs	Tag freque	Tissue
IGF2BP2	3.9056	Retina
IGF2BP2	40.8537	stem_cell
IGF2BP2	4.65669	Eye
IF2B	25.9793	Muscle
IGF2BP2	7.76986	Brain_astrocyte
IGF2BP2	7.90319	White_Blood_Cells
IF2B	28.0995	Prostate
IGF2BP2	8.36622	Prostate
IF2B	30.7436	stem_cell
IF2B	32.2516	Colon
IF2B	38.8493	Brain_astrocyte
IF2B	4.50918	Lung
IF2B	46.4738	Vascular_endothelium
IF2B	5.45308	Stomach
IF2B	6.03245	Liver
IF2B	6.7749	Placenta
IF2B	7.30127	Spinal_cord
EIF3S4	113.33	White_Blood_Cells
EIF3S4	11.6185	Vascular_endothelium
EIF3S4	18.2532	Spinal_cord
EIF3S4	22.3099	Prostate
EIF3S4	22.4176	Brain_cortex
EIF3S4	23.2834	Eye
EIF3S4	33.402	Muscle
EIF3S4	35.6351	Breast
EIF3S4	3.96944	Brain_cerebellum
EIF3S4	42.5145	stem_cell
EIF3S4	6.76376	Lung
EIF3S4	7.76986	Brain_astrocyte
EIF3S4	7.86805	Placenta
EIF3S4	8.06289	Colon
EIF3S4	9.04868	Liver
EIF3S4	9.49559	Retina
EIF3S9	10.9062	Stomach
EIF3S9	11.616	White_Blood_Cells
EIF3S9	11.9083	Brain_cerebellum
EIF3S9	126.536	stem_cell
EIF3S9	14.6025	Spinal_cord
EIF3S9	14.8453	Muscle
EIF3S9	19.4246	Brain_astrocyte
EIF3S9	21.6703	Heart
EIF3S9	24.1298	Liver
EIF3S9	24.1508	Thyroid
EIF3S9	25.0987	Prostate
EIF3S9	25.7885	Brain_cortex
EIF3S9	27.0551	Lung
EIF3S9	27.9154	Vascular_endothelium
EIF3S9	4.03145	Colon
EIF3S9	45.5788	Retina
EIF3S9	4.65669	Eye
EIF3S9	57.0434	Placenta
EIF3S9	72.1885	Breast

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RBPs	Tag freque	Tissue
EIF4H	12.0649	Liver
EIF4H	14.8453	Muscle
EIF4H	156.727	stem_cell
EIF4H	16.2984	Eye
EIF4H	17.5752	Retina
EIF4H	24.1887	Colon
EIF4H	28.0995	Prostate
EIF4H	32.7761	Thyroid
EIF4H	33.7094	Heart
EIF4H	35.0736	Vascular_endothelium
EIF4H	35.7249	Brain_cerebellum
EIF4H	36.5063	Spinal_cord
EIF4H	38.4301	Brain_cortex
EIF4H	40.8981	Stomach
EIF4H	45.0918	Lung
EIF4H	65.3134	Breast
EIF4H	66.0438	Brain_astrocyte
EIF4H	67.2156	Placenta
EIF4H	88.1591	White_Blood_Cells
IGF2BP1	12.6611	stem_cell
IGF2BP1	3.65063	Spinal_cord
IGF2BP3	3.43755	Breast
IGF2BP3	3.48943	Vascular_endothelium
IGF2BP3	3.71575	White_Blood_Cells
IGF2BP3	3.88493	Brain_astrocyte
IGF2BP3	43.4969	stem_cell
IGF2BP3	54.1992	Placenta
IGF2BP3	8.36622	Prostate
ILF3	10.9062	Stomach
ILF3	11.64172	Eye
ILF3	12.18638	Brain_cortex
ILF3	13.7691	Placenta
ILF3	13.85504	Retina
ILF3	19.36412	Vascular_endothelium
ILF3	20.18804	Prostate
ILF3	24.8005	Lung
ILF3	3.01623	Liver
ILF3	3.45012	Thyroid
ILF3	35.97332	Breast
ILF3	3.71133	Muscle
ILF3	3.88493	Brain_astrocyte
ILF3	3.96944	Brain_cerebellum
ILF3	4.03145	Colon
ILF3	92.8587	stem_cell
ILF3	9.74476	White_Blood_Cells
IREB1	12.0649	Liver
IREB1	13.4431	Placenta
IREB1	14.8294	Muscle
IREB1	17.4471	Vascular_endothelium
IREB1	18.6267	Eye
IREB1	19.7713	stem_cell
IREB1	20.2439	Prostate

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RBPs	Tag freque	Tissue
IREB1	4.03145	Colon
IREB1	4.50918	Lung
IREB1	5.1577	Brain_cortex
IREB1	5.7966	Breast
IREB1	7.22343	Heart
IREB1	7.30127	Spinal_cord
IREB1	7.76986	Brain_astrocyte
IREB1	7.8112	Retina
JOSD2	10.9519	Spinal_cord
JOSD2	11.2729	Lung
JOSD2	13.4431	Placenta
JOSD2	14.0294	Vascular_endothelium
JOSD2	14.8568	Brain_cortex
JOSD2	18.2138	White_Blood_Cells
JOSD2	18.9757	Thyroid
JOSD2	19.5212	Prostate
JOSD2	20.2926	stem_cell
JOSD2	3.71133	Muscle
JOSD2	3.88493	Brain_astrocyte
JOSD2	3.95946	Breast
JOSD2	4.03145	Colon
JOSD2	6.03245	Liver
JOSD2	7.22343	Heart
JOSD2	7.59648	Retina
JOSD2	7.93887	Brain_cerebellum
KHDRBS3	10.122	Prostate
KHDRBS3	10.4855	Breast
KHDRBS3	10.5146	stem_cell
KHDRBS3	12.8942	Brain_cortex
KHDRBS3	3.70734	Muscle
KHDRBS3	3.96944	Brain_cerebellum
KHDRBS3	4.65669	Eye
KHDRBS3	6.03245	Liver
KHDRBS3	6.70455	Retina
KHDRBS3	7.71615	Vascular_endothelium
KHDRBS3	8.17962	Stomach
KHDRBS2	12.0649	Liver
KRR1	10.4683	Vascular_endothelium
KRR1	3.65063	Spinal_cord
KRR1	3.71133	Muscle
KRR1	3.71575	White_Blood_Cells
KRR1	6.90024	Thyroid
KRR1	6.99032	Breast
KRR1	7.52499	Retina
KRR1	7.71709	stem_cell
KRR1	7.93888	Brain_cerebellum
KRR1	9.36651	Prostate
LA	10.1623	Placenta
LA	10.4855	Breast
LA	13.496	Prostate
LA	3.01623	Liver
LA	35.2339	stem_cell

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RBPs	Tag freque	Tissue
LA	3.65063	Spinal_cord
LA	3.71133	Muscle
LA	3.87282	Vascular_endothelium
LA	3.88493	Brain_astrocyte
LA	4.81562	Heart
LA	5.8584	Retina
LA	6.40502	Brain_cortex
LA	8.00312	White_Blood_Cells
LA	8.17962	Stomach
LA	8.6253	Thyroid
LARP4	10.13291	stem_cell
LARP4	12.06491	Liver
LARP4	16.28208	Breast
LARP4	3.37399	Prostate
LARP4	3.45012	Thyroid
LARP4	3.65063	Spinal_cord
LARP4	3.66073	Placenta
LARP4	3.88493	Brain_astrocyte
LARP4	4.65669	Eye
LARP4	5.45308	Stomach
LARP4	6.76377	Lung
LARP4	6.99679	Vascular_endothelium
LARP4	7.15051	Retina
LARP4	7.41468	Muscle
LARP4	9.82804	White_Blood_Cells
LARP5	3.95946	Breast
LARP7	11.9149	stem_cell
LARP7	16.1901	White_Blood_Cells
LARP7	4.50918	Lung
LARP7	5.7966	Breast
LKAP	12.7344	Brain_cortex
LKAP	18.007	White_Blood_Cells
LKAP	3.12217	Prostate
LKAP	3.45012	Thyroid
LKAP	3.49516	Breast
LKAP	3.70734	Muscle
LKAP	3.87282	Vascular_endothelium
LKAP	3.89573	stem_cell
LKAP	3.96944	Brain_cerebellum
LKAP	4.50918	Lung
LKAP	6.98503	Eye
LKAP	7.30127	Spinal_cord
ADAD2	6.40502	Brain_cortex
LRPPRC	11.127	Prostate
LRPPRC	11.16347	Retina
LRPPRC	11.64172	Eye
LRPPRC	13.41787	White_Blood_Cells
LRPPRC	13.80046	Thyroid
LRPPRC	15.7697	Brain_cortex
LRPPRC	19.94362	Placenta
LRPPRC	20.97096	Breast
LRPPRC	39.0256	stem_cell

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RBPs	Tag freque	Tissue
LRPPRC	3.96944	Brain_cerebellum
LRPPRC	5.1441	Vascular_endothelium
LRPPRC	5.45308	Stomach
LRPPRC	7.22343	Heart
LRPPRC	7.30127	Spinal_cord
LRPPRC	7.42267	Muscle
LRPPRC	7.76986	Brain_astrocyte
LRPPRC	9.01836	Lung
LRPPRC	9.04868	Liver
LSM5	10.3126	Breast
LSM5	10.4683	Vascular_endothelium
LSM5	10.9519	Spinal_cord
LSM5	11.6548	Brain_astrocyte
LSM5	12.0391	Heart
LSM5	12.0649	Liver
LSM5	13.2938	Retina
LSM5	13.7691	Placenta
LSM5	14.8294	Muscle
LSM5	15.7821	Lung
LSM5	17.2506	Thyroid
LSM5	22.2541	Prostate
LSM5	32.7185	Stomach
LSM5	37.86953	stem_cell
LSM5	4.70125	Brain_cortex
LSM5	9.31337	Eye
LSM5	9.87898	White_Blood_Cells
MOV10L1	10.4683	Vascular_endothelium
MATR3	3.65063	Spinal_cord
MATR3	5.79284	stem_cell
MATR3	7.59648	Retina
RNMT	10.1623	Placenta
RNMT	15.193	Retina
RNMT	3.37399	Prostate
RNMT	3.95946	Breast
RNMT	3.96944	Brain_cerebellum
RNMT	4.03145	Colon
RNMT	4.65669	Eye
RNMT	4.81562	Heart
RNMT	7.30127	Spinal_cord
RNMT	7.74398	White_Blood_Cells
RNMT	7.76986	Brain_astrocyte
RNMT	7.90852	stem_cell
MGC10433	11.2729	Lung
MGC10433	14.8453	Muscle
MGC10433	15.4913	Vascular_endothelium
MGC10433	18.4478	stem_cell
MGC10433	3.96944	Brain_cerebellum
MGC10433	4.03145	Colon
MGC10433	4.81562	Heart
MGC10433	5.1577	Brain_cortex
MGC10433	6.70455	Retina
MGC10433	7.90319	White_Blood_Cells

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RBPs	Tag freque	Tissue
MGC10433	7.91891	Breast
MGC10433	9.31337	Eye
MGC10433	9.36651	Prostate
MGC10433	9.83507	Placenta
MGC27016	11.2729	Lung
MGC27016	16.1901	White_Blood_Cells
MGC27016	17.8511	stem_cell
MGC27016	21.0441	Vascular_endothelium
MGC27016	3.38745	Placenta
MGC27016	3.79824	Retina
MGC27016	3.96944	Brain_cerebellum
MGC27016	6.40502	Brain_cortex
MGC27016	6.74798	Prostate
MGC27016	6.99032	Breast
MGC27016	7.30127	Spinal_cord
MGC27016	8.6253	Thyroid
MINT	10.2882	Vascular_endothelium
MINT	11.8784	Breast
MINT	3.20251	Brain_cortex
MINT	3.88493	Brain_astrocyte
MINT	3.93403	Placenta
MINT	44.0796	White_Blood_Cells
MINT	6.60635	stem_cell
MKI67IP	11.6548	Brain_astrocyte
MKI67IP	34.6713	stem_cell
MKI67IP	3.65063	Spinal_cord
MKI67IP	3.71133	Muscle
MKI67IP	3.79824	Retina
MKI67IP	3.96944	Brain_cerebellum
MKI67IP	5.1577	Brain_cortex
MKI67IP	5.17518	Thyroid
MKI67IP	5.45308	Stomach
MKI67IP	6.97885	Vascular_endothelium
MKI67IP	6.98503	Eye
MKI67IP	7.45462	Breast
MKI67IP	7.80428	White_Blood_Cells
MKI67IP	9.01835	Lung
MKI67IP	9.04868	Liver
MKI67IP	9.36651	Prostate
MS11	13.9577	Vascular_endothelium
MS11	18.6267	Eye
MS11	19.7713	stem_cell
MS11	20.1136	Retina
MS11	3.20251	Brain_cortex
MS11	3.65063	Spinal_cord
MS11	6.99032	Breast
MS12H	3.96944	Brain_cerebellum
MTHFSD	10.122	Prostate
MTHFSD	11.134	Muscle
MTHFSD	11.8784	Breast
MTHFSD	13.4091	Retina
MTHFSD	16.8547	Heart

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RBPs	Tag freque	Tissue
MTHFSD	18.0367	Lung
MTHFSD	18.2532	Spinal_cord
MTHFSD	18.5787	White_Blood_Cells
MTHFSD	43.827	stem_cell
MTHFSD	5.08117	Placenta
MTHFSD	5.45308	Stomach
MTHFSD	7.73655	Brain_cortex
MTHFSD	7.74563	Vascular_endothelium
MTHFSD	8.6253	Thyroid
MTHFSD	9.04868	Liver
MYEF2	10.98462	Retina
MYEF2	14.73846	stem_cell
MYEF2	17.139	Brain_cortex
MYEF2	3.70734	Muscle
MYEF2	3.96944	Brain_cerebellum
MYEF2	4.65669	Eye
MYEF2	7.30127	Spinal_cord
MYEF2	7.71615	Vascular_endothelium
MYEF2	9.61833	Prostate
SLC4A1AP	10.3504	Thyroid
SLC4A1AP	11.8021	Placenta
SLC4A1AP	12.0649	Liver
SLC4A1AP	15.8777	Brain_cerebellum
SLC4A1AP	20.6308	Brain_cortex
SLC4A1AP	3.49516	Breast
SLC4A1AP	3.71133	Muscle
SLC4A1AP	4.50918	Lung
SLC4A1AP	5.1441	Vascular_endothelium
SLC4A1AP	6.24434	Prostate
SLC4A1AP	7.30127	Spinal_cord
SLC4A1AP	7.37912	stem_cell
SLC4A1AP	7.74398	White_Blood_Cells
SLC4A1AP	9.764	Retina
NCBP1	3.65063	Spinal_cord
NCBP2	10.8548	stem_cell
NCBP2	11.9083	Brain_cerebellum
NCBP2	17.4471	Vascular_endothelium
NCBP2	3.45012	Thyroid
NCBP2	3.65063	Spinal_cord
NCBP2	3.70734	Muscle
NCBP2	3.79824	Retina
NCBP2	3.88493	Brain_astrocyte
NCBP2	4.00156	White_Blood_Cells
NCBP2	4.2448	Brain_cortex
NCBP2	7.91891	Breast
NCBP2	8.34527	Prostate
NCBP2	9.31337	Eye
NHP2L1	141.6359	Prostate
NHP2L1	15.08113	Liver
NHP2L1	18.62674	Eye
NHP2L1	24.5389	Stomach
NHP2L1	25.9634	Muscle

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RBPs	Tag freque	Tissue
NHP2L1	27.10973	Vascular_endothelium
NHP2L1	38.8493	Brain_astrocyte
NHP2L1	39.6944	Brain_cerebellum
NHP2L1	4.03145	Colon
NHP2L1	41.4015	Thyroid
NHP2L1	42.8371	Lung
NHP2L1	43.8076	Spinal_cord
NHP2L1	46.4193	Brain_cortex
NHP2L1	51.6809	White_Blood_Cells
NHP2L1	53.7725	Placenta
NHP2L1	67.4187	Heart
NHP2L1	75.1537	Breast
NHP2L1	92.7738	stem_cell
NHP2L1	95.0863	Retina
NKRF	3.01623	Liver
NKRF	3.12217	Prostate
NKRF	3.43755	Breast
NKRF	3.48943	Vascular_endothelium
NKRF	3.88493	Brain_astrocyte
NKRF	9.18875	stem_cell
NOL3	11.06845	Brain_cortex
NOL3	13.45488	Retina
NOL3	16.11449	Prostate
NOL3	19.26248	Heart
NOL3	25.9634	Muscle
NOL3	3.01623	Liver
NOL3	3.96944	Brain_cerebellum
NOL3	4.65669	Eye
NOL3	5.63944	White_Blood_Cells
NOL3	7.74564	Vascular_endothelium
NOL3	8.14156	Placenta
NOL3	9.01835	Lung
NOL4	3.96944	Brain_cerebellum
NOL4	7.71615	Vascular_endothelium
NOL8	3.01623	Liver
NOL8	3.70734	Muscle
NOL8	3.9056	Retina
NOL8	3.95159	White_Blood_Cells
NOL8	4.50918	Lung
NOL8	5.17518	Thyroid
NOL8	7.01471	Vascular_endothelium
NONO	14.7582	stem_cell
NONO	3.12217	Prostate
NONO	3.35227	Retina
NONO	3.93229	White_Blood_Cells
NONO	5.7966	Breast
NOP56	3.49516	Breast
NOP56	4.48395	stem_cell
NOP56	7.01471	Vascular_endothelium
NOP56	7.45351	Retina
NOP56	7.74398	White_Blood_Cells
NOVA1	11.0687	stem_cell

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RBPs	Tag freque	Tissue
NOVA1	7.01471	Vascular_endothelium
NOVA1	7.30127	Spinal_cord
NOVA2	3.49516	Breast
NOVA2	3.96944	Brain_cerebellum
NOVA2	7.37912	stem_cell
NPM1	10.4683	Vascular_endothelium
NPM1	10.9062	Stomach
NPM1	11.0199	White_Blood_Cells
NPM1	12.4887	Prostate
NPM1	13.9806	Breast
NPM1	15.0811	Liver
NPM1	3.20251	Brain_cortex
NPM1	3.79824	Retina
NPM1	3.96944	Brain_cerebellum
NPM1	4.50918	Lung
NPM1	4.65669	Eye
NPM1	5.17518	Thyroid
NPM1	66.9068	stem_cell
NPM1	6.72156	Placenta
NCL	18.62679	Eye
NCL	226.8787	Breast
NCL	33.402	Muscle
NCL	341.549	stem_cell
NCL	39.6943	Brain_cerebellum
NCL	40.1569	Spinal_cord
NCL	40.9328	Heart
NCL	41.8473	Prostate
NCL	42.2272	Liver
NCL	54.389	Brain_astrocyte
NCL	55.8584	Brain_cortex
NCL	59.4457	Placenta
NCL	59.4625	Retina
NCL	68.516	White_Blood_Cells
NCL	73.4361	Vascular_endothelium
NCL	77.6277	Thyroid
NCL	79.0697	Stomach
NCL	8.06289	Colon
NCL	92.4381	Lung
NUFIP1	3.45012	Thyroid
NUFIP1	3.48943	Vascular_endothelium
NUFIP1	3.49516	Breast
NUFIP1	3.88493	Brain_astrocyte
NUFIP1	5.56351	Prostate
NUFIP1	6.07128	White_Blood_Cells
NUFIP1	7.37912	stem_cell
HRB	3.20251	Brain_cortex
HRB	3.37399	Prostate
HRB	3.48943	Vascular_endothelium
HRB	3.71575	White_Blood_Cells
HRB	3.88493	Brain_astrocyte
HRB	5.27235	stem_cell
HRB	5.45308	Stomach

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RBPs	Tag freque	Tissue
HRB	6.72156	Placenta
HRB	7.30127	Spinal_cord
HRB	7.91891	Breast
NXF1	12.8602	Vascular_endothelium
NXF1	3.65063	Spinal_cord
NXF1	40.0156	White_Blood_Cells
NXF1	6.24434	Prostate
NXF1	7.22343	Heart
NXF1	7.91891	Breast
NXF1	8.14108	stem_cell
NXF2	3.20251	Brain_cortex
NXF2	6.24434	Prostate
PABP2	102.811	stem_cell
PABP2	12.0649	Liver
PABP2	15.5397	Brain_astrocyte
PABP2	15.7821	Lung
PABP2	18.9757	Thyroid
PABP2	24.5515	Vascular_endothelium
PABP2	29.726	White_Blood_Cells
PABP2	30.9462	Brain_cortex
PABP2	31.7555	Brain_cerebellum
PABP2	37.813	Breast
PABP2	39.3403	Placenta
PABP2	43.6247	Stomach
PABP2	46.8326	Prostate
PABP2	58.4101	Spinal_cord
PABP2	58.8727	Retina
PABP2	7.41468	Muscle
PABP2	9.31337	Eye
PABP2	9.63124	Heart
PABP4	14.4915	Breast
PABP4	23.6179	Prostate
PABP4	25.5599	White_Blood_Cells
PABP4	25.9514	Muscle
PABP4	27.9154	Vascular_endothelium
PABP4	3.01623	Liver
PABP4	3.45012	Thyroid
PABP4	34.9644	Brain_astrocyte
PABP4	3.96944	Brain_cerebellum
PABP4	4.81562	Heart
PABP4	50.4092	stem_cell
PABP4	5.8584	Retina
PABP4	6.40502	Brain_cortex
PABP4	6.98503	Eye
PABP4	7.30127	Spinal_cord
PABP4	8.17962	Stomach
PABP5	3.20251	Brain_cortex
PABP5	3.49516	Breast
PABP5	4.797443	stem_cell
PAIP1	10.1188	White_Blood_Cells
PAIP1	10.12197	Prostate
PAIP1	11.122	Muscle

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RBPs	Tag freque	Tissue
PAIP1	12.0391	Heart
PAIP1	18.0043	Vascular_endothelium
PAIP1	22.34347	Retina
PAIP1	3.38745	Placenta
PAIP1	33.94018	stem_cell
PAIP1	3.45012	Thyroid
PAIP1	3.88493	Brain_astrocyte
PAIP1	3.95946	Breast
PAIP1	4.03145	Colon
PAIP1	4.65669	Eye
PAIP1	5.1577	Brain_cortex
PAIP1	7.30127	Spinal_cord
PAIP1	7.93888	Brain_cerebellum
PAIP1	9.01836	Lung
PAIP1	9.04868	Liver
C20orf119	15.7361	Placenta
C20orf119	3.20251	Brain_cortex
C20orf119	3.95946	Breast
C20orf119	6.76376	Lung
C20orf119	7.01471	Vascular_endothelium
C20orf119	7.43149	White_Blood_Cells
C20orf119	7.97146	stem_cell
PAPOLA	10.4855	Breast
PAPOLA	11.6185	Vascular_endothelium
PAPOLA	11.6548	Brain_astrocyte
PAPOLA	12.9536	stem_cell
PAPOLA	3.01623	Liver
PAPOLA	3.45012	Thyroid
PAPOLA	3.71133	Muscle
PAPOLA	4.2448	Brain_cortex
PAPOLA	4.50918	Lung
PAPOLA	4.81562	Heart
PAPOLA	5.45308	Stomach
PAPOLA	8.09504	White_Blood_Cells
PAPOLA	8.36622	Prostate
PAPOLG	10.88753	Vascular_endothelium
PAPOLG	11.14798	Prostate
PAPOLG	3.9056	Retina
PAPOLG	3.96944	Brain_cerebellum
PAPOLG	5.17518	Thyroid
PAPOLG	6.76377	Lung
PAPOLG	7.30126	Spinal_cord
PAPOLG	7.76986	Brain_astrocyte
PAPOLG	9.29176	Breast
PAPOLG	9.56221	Placenta
PAPOLG	9.842092	stem_cell
PCBP1	15.8777	Brain_cerebellum
PCBP1	16.2984	Eye
PCBP1	19.1016	Brain_cortex
PCBP1	20.2439	Prostate
PCBP1	21.1136	Liver
PCBP1	23.4336	Retina

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RBPs	Tag freque	Tissue
PCBP1	29.992	Stomach
PCBP1	36.1172	Heart
PCBP1	37.9513	Thyroid
PCBP1	38.5807	Vascular_endothelium
PCBP1	40.157	Spinal_cord
PCBP1	40.3294	Placenta
PCBP1	40.5826	Lung
PCBP1	40.7808	Muscle
PCBP1	54.389	Brain_astrocyte
PCBP1	61.2497	White_Blood_Cells
PCBP1	8.06289	Colon
PCBP1	85.9387	Breast
PCBP1	89.2556	stem_cell
PCBP2	13.5275	Lung
PCBP2	15.4731	Brain_cortex
PCBP2	15.6224	Retina
PCBP2	16.8547	Heart
PCBP2	17.4471	Vascular_endothelium
PCBP2	18.2532	Spinal_cord
PCBP2	19.4246	Brain_astrocyte
PCBP2	19.6701	Placenta
PCBP2	24.0628	Breast
PCBP2	30.3659	Prostate
PCBP2	31.0511	Thyroid
PCBP2	36.733	White_Blood_Cells
PCBP2	3.96944	Brain_cerebellum
PCBP2	5.45308	Stomach
PCBP2	56.825	stem_cell
PCBP2	7.42267	Muscle
PCBP2	8.06289	Colon
PCBP2	9.04868	Liver
PCBP2	9.31337	Eye
PCBP3	15.6224	Retina
PCBP3	15.8777	Brain_cerebellum
PCBP3	16.2984	Eye
PCBP3	3.71575	White_Blood_Cells
PCBP3	3.88493	Brain_astrocyte
POLDIP3	11.2026	Placenta
POLDIP3	11.6548	Brain_astrocyte
POLDIP3	13.496	Prostate
POLDIP3	16.0249	stem_cell
POLDIP3	18.3665	White_Blood_Cells
POLDIP3	3.01623	Liver
POLDIP3	3.20251	Brain_cortex
POLDIP3	3.45012	Thyroid
POLDIP3	3.49516	Breast
POLDIP3	3.65063	Spinal_cord
POLDIP3	4.50918	Lung
POLDIP3	4.65669	Eye
POLDIP3	5.69736	Retina
POLDIP3	6.97885	Vascular_endothelium
POLDIP3	7.41468	Muscle

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RBPs	Tag freque	Tissue
POLDIP3	7.93887	Brain_cerebellum
SF3B14	13.4431	Placenta
SF3B14	14.0055	White_Blood_Cells
SF3B14	15.5255	Thyroid
SF3B14	15.5397	Brain_astrocyte
SF3B14	23.7567	Breast
SF3B14	27.2654	Stomach
SF3B14	27.9154	Vascular_endothelium
SF3B14	4.03145	Colon
SF3B14	40.5882	Prostate
SF3B14	41.6526	stem_cell
SF3B14	5.1577	Brain_cortex
SF3B14	6.03245	Liver
SF3B14	7.42267	Muscle
SF3B14	7.93887	Brain_cerebellum
SF3B14	9.01835	Lung
SF3B14	9.63124	Heart
PNPT1	3.12217	Prostate
PNPT1	3.87282	Vascular_endothelium
PPP1R8	10.9062	Stomach
PPP1R8	11.1472	White_Blood_Cells
PPP1R8	11.155	Prostate
PPP1R8	11.2729	Lung
PPP1R8	11.6417	Eye
PPP1R8	11.6548	Brain_astrocyte
PPP1R8	12.8942	Brain_cortex
PPP1R8	13.7502	Breast
PPP1R8	14.4469	Heart
PPP1R8	15.6836	Placenta
PPP1R8	23.9821	stem_cell
PPP1R8	3.01623	Liver
PPP1R8	3.45012	Thyroid
PPP1R8	3.65063	Spinal_cord
PPP1R8	3.71133	Muscle
PPP1R8	3.96944	Brain_cerebellum
PPP1R8	4.03145	Colon
PPP1R8	5.1441	Vascular_endothelium
PPP1R8	9.49559	Retina
PPIE	11.6185	Vascular_endothelium
PPIE	11.8628	stem_cell
PPIE	18.5787	White_Blood_Cells
PPIE	3.01623	Liver
PPIE	3.38745	Placenta
PPIE	3.88493	Brain_astrocyte
PPIE	3.9056	Retina
PPIE	4.50918	Lung
PPIE	5.56351	Prostate
PPIE	5.7966	Breast
PPIE	6.98503	Eye
PPIE	7.30127	Spinal_cord
PPIE	7.41468	Muscle
PPIL4	17.4471	Vascular_endothelium

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RBPs	Tag freque	Tissue
PPIL4	3.01623	Liver
PPIL4	3.20251	Brain_cortex
PPIL4	3.43755	Breast
PPIL4	4.48104	Placenta
PPIL4	4.876646	stem_cell
PPIL4	7.30127	Spinal_cord
PPIL4	8.36622	Prostate
PPRC1	3.12217	Prostate
PPRC1	3.20251	Brain_cortex
PPRC1	3.71133	Muscle
PPRC1	3.93229	White_Blood_Cells
PPRC1	3.96944	Brain_cerebellum
PPRC1	5.53434	stem_cell
PPARGC1E	3.87199	White_Blood_Cells
PPARGC1E	7.93887	Brain_cerebellum
PRKRA	11.27294	Lung
PRKRA	11.35647	Breast
PRKRA	12.0391	Heart
PRKRA	12.86025	Vascular_endothelium
PRKRA	13.80048	Thyroid
PRKRA	14.60253	Spinal_cord
PRKRA	14.83735	Muscle
PRKRA	16.36634	Prostate
PRKRA	19.77131	stem_cell
PRKRA	22.78939	Retina
PRKRA	4.00156	White_Blood_Cells
PRKRA	4.65669	Eye
PRKRA	5.35446	Placenta
PRKRA	7.76986	Brain_astrocyte
PRKRA	7.93888	Brain_cerebellum
PRKRA	9.4025	Brain_cortex
PSMA1	10.0568	Retina
PSMA1	12.81	Brain_cortex
PSMA1	13.6327	Stomach
PSMA1	14.4469	Heart
PSMA1	15.8777	Brain_cerebellum
PSMA1	18.5367	Muscle
PSMA1	19.7973	Breast
PSMA1	20.1572	Colon
PSMA1	25.0987	Prostate
PSMA1	31.0794	Brain_astrocyte
PSMA1	40.0156	White_Blood_Cells
PSMA1	4.48104	Placenta
PSMA1	46.4738	Vascular_endothelium
PSMA1	54.9919	stem_cell
PSMA1	6.03245	Liver
PSMA1	6.90024	Thyroid
PSMA1	7.30127	Spinal_cord
PSMA1	9.31337	Eye
PSPC1	11.616	White_Blood_Cells
PSPC1	3.01623	Liver
PSPC1	3.20251	Brain_cortex

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RBPs	Tag freque	Tissue
PSPC1	3.37399	Prostate
PSPC1	3.45012	Thyroid
PSPC1	3.50736	Vascular_endothelium
PSPC1	3.65063	Spinal_cord
PSPC1	3.70734	Muscle
PSPC1	3.79824	Retina
PSPC1	3.96944	Brain_cerebellum
PSPC1	4.7132	stem_cell
PSPC1	6.8751	Breast
PSPC1	8.46862	Placenta
PTBP1	11.6417	Eye
PTBP1	121.5538	stem_cell
PTBP1	12.81	Brain_cortex
PTBP1	13.91574	Prostate
PTBP1	23.75676	Breast
PTBP1	24.1887	Colon
PTBP1	3.65063	Spinal_cord
PTBP1	3.71133	Muscle
PTBP1	3.88493	Brain_astrocyte
PTBP1	3.96944	Brain_cerebellum
PTBP1	5.69736	Retina
PTBP1	71.7839	Vascular_endothelium
PTBP1	7.59476	Placenta
PTBP1	82.83626	White_Blood_Cells
PTBP2	3.12217	Prostate
PTBP2	3.35227	Retina
PTBP2	3.38745	Placenta
PTBP2	3.49516	Breast
PTBP2	3.65063	Spinal_cord
PTBP2	6.9421	stem_cell
Q8TC92	4.65669	Eye
QKI	11.122	Muscle
QKI	11.6548	Brain_astrocyte
QKI	13.4091	Retina
QKI	15.2435	Placenta
QKI	19.8472	Brain_cerebellum
QKI	22.0086	White_Blood_Cells
QKI	226.339	Spinal_cord
QKI	26.0847	Breast
QKI	26.4859	Heart
QKI	29.3096	Lung
QKI	4.65669	Eye
QKI	5.17518	Thyroid
QKI	6.03245	Liver
QKI	71.9464	stem_cell
QKI	77.3654	Brain_cortex
QKI	7.74563	Vascular_endothelium
RAD51AP1	15.2062	stem_cell
RAD51AP1	4.48104	Placenta
RAD51AP1	5.69736	Retina
RAD51AP1	6.97885	Vascular_endothelium
RAE1	10.2765	stem_cell

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RBPs	Tag freque	Tissue
RAE1	14.1663	White_Blood_Cells
RAE1	3.79824	Retina
RAE1	3.88493	Brain_astrocyte
RAE1	6.74798	Prostate
RAE1	6.99032	Breast
RAE1	7.74563	Vascular_endothelium
RBM12B	3.48943	Vascular_endothelium
RBM12B	5.73526	stem_cell
RBM15B	3.01623	Liver
RBM15B	4.45458	stem_cell
RBM15B	7.43149	White_Blood_Cells
RBM35A	3.93403	Placenta
RBM35A	4.03145	Colon
RBM35A	9.41337	stem_cell
RBM35B	13.7691	Placenta
RBM35B	15.0811	Liver
RBM35B	3.49516	Breast
RBM35B	4.50918	Lung
RBM35B	5.45308	Stomach
RBM35B	5.57748	Prostate
RBM35B	8.06289	Colon
RBM35B	8.60289	stem_cell
RBM10	11.9083	Brain_cerebellum
RBM10	13.496	Prostate
RBM10	3.45012	Thyroid
RBM10	3.65063	Spinal_cord
RBM10	3.71133	Muscle
RBM10	3.88493	Brain_astrocyte
RBM10	43.7248	Vascular_endothelium
RBM10	5.90104	Placenta
RBM10	62.9166	White_Blood_Cells
RBM10	6.70455	Retina
RBM10	6.76376	Lung
RBM10	75.636	stem_cell
RBM10	7.73655	Brain_cortex
RBM10	7.91891	Breast
RBM11	3.65063	Spinal_cord
RBM11	3.96944	Brain_cerebellum
RBM11	5.1577	Brain_cortex
RBM11	9.764	Retina
RBM12	10.02616	Brain_cortex
RBM12	11.64172	Eye
RBM12	14.83734	Muscle
RBM12	24.0308	Prostate
RBM12	25.30871	Breast
RBM12	3.01623	Liver
RBM12	31.07939	Brain_astrocyte
RBM12	39.05608	White_Blood_Cells
RBM12	3.96944	Brain_cerebellum
RBM12	5.45308	Stomach
RBM12	5.67955	Retina
RBM12	60.6854	stem_cell

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RBPs	Tag freque	Tissue
RBM12	7.86827	Placenta
RBM12	8.6253	Thyroid
RBM12	94.26034	Vascular_endothelium
RBM14	11.1472	White_Blood_Cells
RBM14	12.4554	stem_cell
RBM14	14.0294	Vascular_endothelium
RBM14	16.2984	Eye
RBM14	3.71133	Muscle
RBM14	3.95946	Breast
RBM14	4.2448	Brain_cortex
RBM14	6.24434	Prostate
RBM15	3.87282	Vascular_endothelium
RBM15	4.65669	Eye
RBM16	3.01623	Liver
RBM16	3.49516	Breast
RBM16	3.88493	Brain_astrocyte
RBM16	4.03145	Colon
RBM16	5.89843	White_Blood_Cells
RBM16	9.54251	stem_cell
RBM18	11.134	Muscle
RBM18	11.2026	Placenta
RBM18	3.20251	Brain_cortex
RBM18	3.85192	Retina
RBM18	3.96944	Brain_cerebellum
RBM18	5.87277	White_Blood_Cells
RBM18	7.74563	Vascular_endothelium
RBM18	7.98604	stem_cell
RBM18	8.95147	Prostate
RBM18	9.63124	Heart
RBM19	3.88493	Brain_astrocyte
RBM22	10.122	Prostate
RBM22	10.9062	Stomach
RBM22	11.9083	Brain_cerebellum
RBM22	3.20251	Brain_cortex
RBM22	3.38745	Placenta
RBM22	3.48943	Vascular_endothelium
RBM22	3.65063	Spinal_cord
RBM22	6.8751	Breast
RBM22	7.43149	White_Blood_Cells
RBM22	7.76986	Brain_astrocyte
RBM22	9.46611	stem_cell
RBM23	10.9519	Spinal_cord
RBM23	11.2729	Lung
RBM23	11.6417	Eye
RBM23	11.6548	Brain_astrocyte
RBM23	12.0943	Colon
RBM23	18.4478	stem_cell
RBM23	18.631	Placenta
RBM23	19.6614	White_Blood_Cells
RBM23	21.8123	Stomach
RBM23	22.4176	Brain_cortex
RBM23	24.1508	Thyroid

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RBPs	Tag freque	Tissue
RBM23	3.01623	Liver
RBM23	30.1705	Retina
RBM23	6.74798	Prostate
RBM23	7.42267	Muscle
RBM23	7.74563	Vascular_endothelium
RBM23	7.91891	Breast
RBM23	9.63124	Heart
RBM24	21.6703	Heart
RBM24	3.12217	Prostate
RBM24	3.35227	Retina
RBM24	3.87282	Vascular_endothelium
RBM24	3.88493	Brain_astrocyte
RBM24	3.93403	Placenta
RBM24	63.0927	Muscle
RBM25	10.122	Prostate
RBM25	11.3947	Retina
RBM25	13.6327	Stomach
RBM25	3.01623	Liver
RBM25	3.70734	Muscle
RBM25	3.88493	Brain_astrocyte
RBM25	4.81562	Heart
RBM25	4.86966	stem_cell
RBM25	5.1441	Vascular_endothelium
RBM25	5.90104	Placenta
RBM25	6.90024	Thyroid
RBM25	7.30127	Spinal_cord
RBM25	7.74398	White_Blood_Cells
RBM25	7.93887	Brain_cerebellum
RBM25	9.01835	Lung
RBM26	11.459	stem_cell
RBM26	13.6327	Stomach
RBM26	3.01623	Liver
RBM26	30.9759	White_Blood_Cells
RBM26	3.88493	Brain_astrocyte
RBM26	4.2448	Brain_cortex
RBM26	5.90104	Placenta
RBM26	6.90024	Thyroid
RBM26	6.98503	Eye
RBM26	6.99032	Breast
RBM26	7.01471	Vascular_endothelium
RBM26	7.30127	Spinal_cord
RBM26	7.59648	Retina
RBM27	3.87282	Vascular_endothelium
RBM27	3.88493	Brain_astrocyte
RBM27	3.96944	Brain_cerebellum
RBM27	5.550247	stem_cell
RBM28	3.35227	Retina
RBM28	3.49516	Breast
RBM28	3.65063	Spinal_cord
RBM28	3.71133	Muscle
RBM28	4.03145	Colon
RBM28	4.48104	Placenta

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RBPs	Tag freque	Tissue
RBM28	5.57748	Prostate
RBM28	7.74563	Vascular_endothelium
RBM3	10.9519	Spinal_cord
RBM3	11.122	Muscle
RBM3	12.0943	Colon
RBM3	12.81	Brain_cortex
RBM3	15.8777	Brain_cerebellum
RBM3	16.2984	Eye
RBM3	24.9774	Prostate
RBM3	27.526	White_Blood_Cells
RBM3	29.218	stem_cell
RBM33	11.1803	Retina
RBM33	11.8548	White_Blood_Cells
RBM3	31.3015	Heart
RBM3	31.4564	Breast
RBM33	3.70734	Muscle
RBM33	3.87282	Vascular_endothelium
RBM33	3.95946	Breast
RBM33	3.96944	Brain_cerebellum
RBM33	4.50918	Lung
RBM33	6.40502	Brain_cortex
RBM3	38.1716	Stomach
RBM34	11.7593	stem_cell
RBM34	3.37399	Prostate
RBM34	3.45012	Thyroid
RBM34	3.88493	Brain_astrocyte
RBM34	3.93403	Placenta
RBM34	3.95159	White_Blood_Cells
RBM34	3.95946	Breast
RBM34	6.3672	Brain_cortex
RBM3	46.6191	Brain_astrocyte
RBM34	6.97885	Vascular_endothelium
RBM3	47.3464	Lung
RBM3	53.7725	Placenta
RBM3	6.03245	Liver
RBM38	10.122	Prostate
RBM38	11.6548	Brain_astrocyte
RBM38	11.9083	Brain_cerebellum
RBM38	22.6141	stem_cell
RBM38	24.1887	Colon
RBM38	3.01623	Liver
RBM38	37.0734	Muscle
RBM38	38.7199	White_Blood_Cells
RBM38	4.2448	Brain_cortex
RBM38	4.50918	Lung
RBM3	8.6253	Thyroid
RBM38	6.7749	Placenta
RBM38	6.8751	Breast
RBM38	7.01471	Vascular_endothelium
RBM38	7.22343	Heart
RBM38	9.31337	Eye
RBM39	10.9519	Spinal_cord

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RBPs	Tag freque	Tissue
RBM39	11.122	Muscle
RBM39	12.0943	Colon
RBM39	13.2938	Retina
RBM39	13.80052	Thyroid
RBM39	13.9701	Eye
RBM39	15.6109	Prostate
RBM39	18.0367	Lung
RBM39	20.77698	stem_cell
RBM39	3.88493	Brain_astrocyte
RBM39	39.3403	Placenta
RBM3	94.2145	Vascular_endothelium
RBM39	43.62468	Stomach
RBM39	46.2969	Vascular_endothelium
RBM39	6.3672	Brain_cortex
RBM39	69.413	Breast
RBM39	7.22343	Heart
RBM39	7.93887	Brain_cerebellum
RBM39	82.03198	White_Blood_Cells
RBM39	9.04868	Liver
RBM4	12.0754	Thyroid
RBM41	3.48943	Vascular_endothelium
RBM4	13.9577	Vascular_endothelium
RBM4	13.9806	Breast
RBM4	15.8777	Brain_cerebellum
RBM4	16.1901	White_Blood_Cells
RBM4	20.2926	stem_cell
RBM4	20.9551	Eye
RBM4	23.6179	Prostate
RBM4	3.65063	Spinal_cord
RBM4	3.79824	Retina
RBM4	6.03245	Liver
RBM4	6.40502	Brain_cortex
RBM4	7.42267	Muscle
RBM4	7.76986	Brain_astrocyte
RBM4	8.06289	Colon
RBM4	8.96208	Placenta
RBM5	10.122	Prostate
RBM5	10.9062	Stomach
RBM5	11.122	Muscle
RBM5	13.3823	stem_cell
RBM5	13.6696	Retina
RBM5	15.7361	Placenta
RBM5	15.8777	Brain_cerebellum
RBM5	19.36	White_Blood_Cells
RBM5	25.5544	Spinal_cord
RBM5	27.1097	Vascular_endothelium
RBM5	3.20251	Brain_cortex
RBM5	3.88493	Brain_astrocyte
RBM5	4.65669	Eye
RBM5	5.7966	Breast
RBM5	6.76376	Lung
RBM5	8.06289	Colon

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RBPs	Tag freque	Tissue
RBM6	11.9083	Brain_cerebellum
RBM6	13.5275	Lung
RBM6	13.6327	Stomach
RBM6	15.5255	Thyroid
RBM6	21.4265	stem_cell
RBM6	21.6371	Placenta
RBM6	23.2096	Brain_cortex
RBM6	23.7567	Breast
RBM6	29.2051	Spinal_cord
RBM6	32.285	Retina
RBM6	43.7248	Vascular_endothelium
RBM6	44.5227	White_Blood_Cells
RBM6	4.65669	Eye
RBM6	6.74798	Prostate
RBM6	9.04868	Liver
RBM7	15.4447	stem_cell
RBM7	15.8378	Breast
RBM7	3.87282	Vascular_endothelium
RBM7	5.56351	Prostate
RBM7	6.00234	White_Blood_Cells
RBM7	6.03245	Liver
RBM7	6.40502	Brain_cortex
RBM7	8.17962	Stomach
RBM7	9.01835	Lung
RBM8A	10.3504	Thyroid
RBM8A	10.90616	Stomach
RBM8A	10.9519	Spinal_cord
RBM8A	11.2729	Lung
RBM8A	17.97662	Placenta
RBM8A	20.82011	Retina
RBM8A	27.8874	Prostate
RBM8A	28.2925	Vascular_endothelium
RBM8A	3.01623	Liver
RBM8A	31.0794	Brain_astrocyte
RBM8A	3.71133	Muscle
RBM8A	4.03145	Colon
RBM8A	41.0024	White_Blood_Cells
RBM8A	4.2448	Brain_cortex
RBM8A	42.56781	Breast
RBM8A	56.08299	stem_cell
RBM8A	6.98503	Eye
RBM8A	7.22343	Heart
RBM8A	7.93888	Brain_cerebellum
RBMS3	3.50736	Vascular_endothelium
RBMS3	5.7966	Breast
RBP56	11.6185	Vascular_endothelium
RBP56	11.6548	Brain_astrocyte
RBP56	27.1039	White_Blood_Cells
RBP56	3.01623	Liver
RBP56	3.35227	Retina
RBP56	3.37399	Prostate
RBP56	34.3734	stem_cell

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RBPs	Tag freque	Tissue
RBP56	3.71133	Muscle
RBP56	3.96944	Brain_cerebellum
RBP56	4.03145	Colon
RBP56	4.50918	Lung
RBP56	4.65669	Eye
RBP56	6.40502	Brain_cortex
RBP56	7.30127	Spinal_cord
RBP56	7.86805	Placenta
RBPMS	10.122	Prostate
RBPMS	10.3504	Thyroid
RBPMS	15.7821	Lung
RBPMS	17.5368	Vascular_endothelium
RBPMS	39.5946	Breast
RBPMS	4.48104	Placenta
RBPMS	4.81562	Heart
RBPMS	59.5037	stem_cell
RBPMS	7.76986	Brain_astrocyte
RBPMS	9.04868	Liver
RBPMS	9.764	Retina
RED1	3.20251	Brain_cortex
RED1	3.65063	Spinal_cord
RED1	3.71133	Muscle
RED1	3.72675	Retina
RED2	3.49516	Breast
RENT1	3.70734	Muscle
RENT1	3.95946	Breast
RENT1	4.00156	White_Blood_Cells
RENT1	7.01471	Vascular_endothelium
RIB1	10.9519	Spinal_cord
RIB1	13.40119	Retina
RIB1	14.14157	Brain_cortex
RIB1	15.5397	Brain_astrocyte
RIB1	19.17753	Placenta
RIB1	23.55038	White_Blood_Cells
RIB1	24.15088	Thyroid
RIB1	24.18865	Colon
RIB1	24.53882	Stomach
RIB1	26.40664	Prostate
RIB1	27.055	Lung
RIB1	29.6587	Muscle
RIB1	33.17853	Liver
RIB1	33.7968	Vascular_endothelium
RIB1	35.3752	stem_cell
RIB1	3.96944	Brain_cerebellum
RIB1	40.93282	Heart
RIB1	47.51351	Breast
RIB1	6.98503	Eye
RPL12	120.433	Brain_astrocyte
RPL12	129.897	Muscle
RPL12	149.676	Spinal_cord
RPL12	151.692	Heart
RPL12	160.656	Eye

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<u>RBPs</u>	<u>Tag freque</u>	<u>Tissue</u>
RPL12	194.685	White_Blood_Cells
RPL12	236.732	Lung
RPL12	309.825	Vascular_endothelium
RPL12	318.154	Placenta
RPL12	371.255	Breast
RPL12	40.3145	Colon
RPL12	452.606	Stomach
RPL12	469.216	Thyroid
RPL12	54.1558	Brain_cortex
RPL12	542.443	Prostate
RPL12	556.158	stem_cell
RPL12	63.511	Brain_cerebellum
RPL12	90.4868	Liver
RPL12	93.8637	Retina
RPL13	102.2177	Spinal_cord
RPL13	106.9537	Thyroid
RPL13	1086.816	Vascular_endothelium
RPL13	124.975	Colon
RPL13	126.6817	Liver
RPL13	158.9157	Heart
RPL13	174.6547	Brain_cerebellum
RPL13	179.1387	White_Blood_Cells
RPL13	225.4593	Lung
RPL13	239.8312	Brain_cortex
RPL13	248.6358	Brain_astrocyte
RPL13	250.9387	Placenta
RPL13	307.737	Muscle
RPL13	341.305	stem_cell
RPL13	381.7162	Stomach
RPL13	637.9665	Eye
RPL13	771.9887	Prostate
RPL13	87.4196	Retina
RPL13	878.9986	Breast
RPL18	100.175	stem_cell
RPL18	126.703	Breast
RPL18	18.0367	Lung
RPL18	224.796	Prostate
RPL18	24.0781	Heart
RPL18	24.1887	Colon
RPL18	25.8759	Thyroid
RPL18	27.1945	Brain_astrocyte
RPL18	31.3673	Placenta
RPL18	31.7555	Brain_cerebellum
RPL18	32.8557	Spinal_cord
RPL18	6.03245	Liver
RPL18	67.9168	Brain_cortex
RPL18	7.59648	Retina
RPL18	88.0343	White_Blood_Cells
RPL18	93.1337	Eye
RPL18	94.2145	Vascular_endothelium
RPL18	96.4946	Muscle
RPL22	129.0307	Breast

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RBPs	Tag freque	Tissue
RPL22	146.5563	Vascular_endothelium
RPL22	15.08113	Liver
RPL22	186.9302	White_Blood_Cells
RPL22	192.4234	Prostate
RPL22	20.1572	Colon
RPL22	21.89755	Retina
RPL22	33.11432	Placenta
RPL22	35.72487	Brain_cerebellum
RPL22	357.6104	stem_cell
RPL22	36.07345	Lung
RPL22	36.5063	Spinal_cord
RPL22	38.52501	Heart
RPL22	65.437	Stomach
RPL22	74.2266	Muscle
RPL22	78.5288	Brain_cortex
RPL22	79.16369	Eye
RPL22	86.253	Thyroid
RPL22	97.12326	Brain_astrocyte
RPL28	3.12217	Prostate
RPL28	3.66613	stem_cell
RPL28	3.72675	Retina
RPL28	3.87282	Vascular_endothelium
RPL28	3.96944	Brain_cerebellum
RPL28	6.99032	Breast
RPL28	8.00312	White_Blood_Cells
RPL34	100.098	Muscle
RPL34	102.5517	Liver
RPL34	130.9914	Brain_cerebellum
RPL34	16.1258	Colon
RPL34	164.279	Spinal_cord
RPL34	173.362	Heart
RPL34	174.761	Placenta
RPL34	181.6103	Eye
RPL34	204.8186	White_Blood_Cells
RPL34	261.5322	Lung
RPL34	263.352	Vascular_endothelium
RPL34	294.466	Stomach
RPL34	301.378	stem_cell
RPL34	379.5131	Thyroid
RPL34	395.4236	Breast
RPL34	46.6191	Brain_astrocyte
RPL34	479.6423	Prostate
RPL34	58.8727	Retina
RPL34	95.4174	Brain_cortex
RPL3L	7.22343	Heart
RPL3L	77.938	Muscle
RPLP2	108.849	Colon
RPLP2	1198.91	stem_cell
RPLP2	127.022	Brain_cerebellum
RPLP2	1374.19	Prostate
RPLP2	1883.78	Breast
RPLP2	223.201	Liver

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RBPs	Tag freque	Tissue
RPLP2	225.995	Retina
RPLP2	273.789	Brain_cortex
RPLP2	306.653	Spinal_cord
RPLP2	318.831	Muscle
RPLP2	321.311	Eye
RPLP2	426.09	Thyroid
RPLP2	445.445	Heart
RPLP2	485.616	Brain_astrocyte
RPLP2	542.194	Vascular_endothelium
RPLP2	616.198	Stomach
RPLP2	635.794	Lung
RPLP2	710.245	Placenta
RPLP2	902.926	White_Blood_Cells
MRPL3	10.1623	Placenta
MRPL3	11.6548	Brain_astrocyte
MRPL3	11.9083	Brain_cerebellum
MRPL3	15.0811	Liver
MRPL3	15.488	White_Blood_Cells
MRPL3	15.4913	Vascular_endothelium
MRPL3	16.8547	Heart
MRPL3	20.6253	Breast
MRPL3	21.4808	Retina
MRPL3	21.8123	Stomach
MRPL3	22.4258	Thyroid
MRPL3	27.8176	Prostate
MRPL3	36.1039	Brain_cortex
MRPL3	49.8091	stem_cell
MRPL3	6.98503	Eye
MRPL3	7.42267	Muscle
MRPL3	9.01835	Lung
MRPL12	12.0943	Colon
MRPL12	15.5397	Brain_astrocyte
MRPL12	15.7291	White_Blood_Cells
MRPL12	16.0126	Brain_cortex
MRPL12	16.6905	Prostate
MRPL12	17.4471	Vascular_endothelium
MRPL12	24.0781	Heart
MRPL12	29.6587	Muscle
MRPL12	3.01623	Liver
MRPL12	3.95946	Breast
MRPL12	4.48104	Placenta
MRPL12	4.50918	Lung
MRPL12	50.6487	stem_cell
MRPL12	7.93887	Brain_cerebellum
MRPL23	10.3504	Thyroid
MRPL23	10.612	Brain_cortex
MRPL23	11.8021	Placenta
MRPL23	12.0391	Heart
MRPL23	12.0649	Liver
MRPL23	16.1258	Colon
MRPL23	23.2369	Vascular_endothelium
MRPL23	25.7131	White_Blood_Cells

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RBPs	Tag freque	Tissue
MRPL23	3.65063	Spinal_cord
MRPL23	3.72675	Retina
MRPL23	40.4879	Prostate
MRPL23	40.7808	Muscle
MRPL23	4.50918	Lung
MRPL23	52.2423	stem_cell
MRPL23	7.76986	Brain_astrocyte
MRPL23	7.91891	Breast
MRPL23	7.93887	Brain_cerebellum
MRPL23	8.17962	Stomach
MRPL23	9.31337	Eye
RNASEL	3.20251	Brain_cortex
RNASEL	3.45012	Thyroid
RNASEL	7.63235	Retina
RNASE1	10.0039	White_Blood_Cells
RNASE1	115.529	Retina
RNASE1	137.069	Colon
RNASE1	182.994	Heart
RNASE1	18.5367	Muscle
RNASE1	18.6267	Eye
RNASE1	21.1136	Liver
RNASE1	239.201	Vascular_endothelium
RNASE1	272.654	Stomach
RNASE1	281.824	Lung
RNASE1	37.1139	Prostate
RNASE1	43.1265	Thyroid
RNASE1	48.8152	Brain_cortex
RNASE1	72.4575	Breast
RNASE1	7.93887	Brain_cerebellum
RNASE1	82.9925	Placenta
RNASE1	83.9646	Spinal_cord
RNH1	10.5146	stem_cell
RNH1	3.01623	Liver
RNH1	3.70734	Muscle
RNH1	4.00156	White_Blood_Cells
RNH1	4.48104	Placenta
RNH1	4.50918	Lung
RNH1	6.24434	Prostate
RNH1	7.22343	Heart
HNRPA2B1	114.515	Stomach
HNRPA2B1	119.029	Thyroid
HNRPA2B1	199.378	Breast
HNRPA2B1	251.755	stem_cell
HNRPA2B1	31.0794	Brain_astrocyte
HNRPA2B1	32.2516	Colon
HNRPA2B1	32.5968	Eye
HNRPA2B1	33.7399	Prostate
HNRPA2B1	38.525	Heart
HNRPA2B1	42.4989	White_Blood_Cells
HNRPA2B1	43.6638	Brain_cerebellum
HNRPA2B1	44.4881	Muscle
HNRPA2B1	53.1753	Retina

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RBPs	Tag freque	Tissue
HNRPA2B1	54.7595	Spinal_cord
HNRPA2B1	59.1571	Vascular_endothelium
HNRPA2B1	63.3408	Liver
HNRPA2B1	71.1364	Placenta
HNRPA2B1	85.102	Brain_cortex
HNRPA2B1	96.9473	Lung
HNRPA3	10.07284	White_Blood_Cells
HNRPA3	10.83456	Breast
HNRPA3	10.9519	Spinal_cord
HNRPA3	11.27294	Lung
HNRPA3	12.07542	Thyroid
HNRPA3	12.0943	Colon
HNRPA3	13.40907	Retina
HNRPA3	13.6327	Stomach
HNRPA3	15.4323	Vascular_endothelium
HNRPA3	15.87774	Brain_cerebellum
HNRPA3	16.86998	Prostate
HNRPA3	18.6267	Eye
HNRPA3	28.15881	stem_cell
HNRPA3	32.02512	Brain_cortex
HNRPA3	3.71133	Muscle
HNRPA3	4.48104	Placenta
HNRPA3	6.03246	Liver
HNRPA3	7.76986	Brain_astrocyte
ROD1	3.93229	White_Blood_Cells
RPL18A	104.818	Colon
RPL18A	133.711	Brain_cortex
RPL18A	134.961	Brain_cerebellum
RPL18A	161.323	Heart
RPL18A	170.937	Brain_astrocyte
RPL18A	187.131	Lung
RPL18A	237.808	White_Blood_Cells
RPL18A	251.859	Thyroid
RPL18A	337.368	Muscle
RPL18A	344.473	Breast
RPL18A	378.648	Placenta
RPL18A	407.46	Eye
RPL18A	441.501	Vascular_endothelium
RPL18A	458.614	stem_cell
RPL18A	54.7595	Spinal_cord
RPL18A	59.9839	Stomach
RPL18A	63.6932	Retina
RPL18A	684.92	Prostate
RPL18A	93.503	Liver
RPL7	3.12217	Prostate
RPL7	3.68956	stem_cell
RPL7	3.87282	Vascular_endothelium
RPL7	3.95946	Breast
RPP29	10.9519	Spinal_cord
RPP29	12.0943	Colon
RPP29	12.81	Brain_cortex
RPP29	3.01623	Liver

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RBPs	Tag freque	Tissue
RPP29	3.71133	Muscle
RPP29	3.79824	Retina
RPP29	3.96944	Brain_cerebellum
RPP29	6.07128	White_Blood_Cells
RPP29	6.7749	Placenta
RPP29	6.90024	Thyroid
RPP29	6.99032	Breast
RPP29	7.22343	Heart
RPP29	7.74563	Vascular_endothelium
RPP29	9.36651	Prostate
RPS17	11.1803	Retina
RPS17	19.4246	Brain_astrocyte
RPS17	21.1136	Liver
RPS17	216.536	Eye
RPS17	245.486	Breast
RPS17	256.386	White_Blood_Cells
RPS17	340.033	stem_cell
RPS17	38.1716	Stomach
RPS17	38.2032	Brain_cortex
RPS17	40.157	Spinal_cord
RPS17	431.172	Prostate
RPS17	43.3406	Heart
RPS17	47.3464	Lung
RPS17	48.2473	Muscle
RPS17	620.802	Vascular_endothelium
RPS17	70.7275	Thyroid
RPS17	7.93887	Brain_cerebellum
RPS17	8.06289	Colon
RPS17	85.1398	Placenta
RPS3	103.806	Muscle
RPS3	106.954	Thyroid
RPS3	125.421	Stomach
RPS3	142.039	Lung
RPS3	219.571	Placenta
RPS3	230.506	Eye
RPS3	364.143	White_Blood_Cells
RPS3	389.446	Prostate
RPS3	39.69434	Brain_cerebellum
RPS3	42.2272	Liver
RPS3	455.337	Breast
RPS3	48.3773	Colon
RPS3	535.1141	stem_cell
RPS3	54.389	Brain_astrocyte
RPS3	551.7198	Vascular_endothelium
RPS3	58.4101	Spinal_cord
RPS3	61.5496	Brain_cortex
RPS3	67.4187	Heart
RPS3	68.3683	Retina
RPS4Y1	106.154	Prostate
RPS4Y1	111.472	White_Blood_Cells
RPS4Y1	12.81	Brain_cortex
RPS4Y1	13.6696	Retina

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RBPs	Tag freque	Tissue
RPS4Y1	20.1572	Colon
RPS4Y1	29.6587	Muscle
RPS4Y1	31.5642	Lung
RPS4Y1	33.7094	Heart
RPS4Y1	3.49516	Breast
RPS4Y1	3.65063	Spinal_cord
RPS4Y1	41.3073	Placenta
RPS4Y1	44.2385	Eye
RPS4Y1	50.3466	Vascular_endothelium
RPS4Y1	5.45308	Stomach
RPS4Y1	89.8201	stem_cell
RPS4Y1	9.04868	Liver
RPS5	102.946	Breast
RPS5	103.917	Muscle
RPS5	104.775	Eye
RPS5	12.0649	Liver
RPS5	144.914	White_Blood_Cells
RPS5	15.6224	Retina
RPS5	157.024	Vascular_endothelium
RPS5	21.9038	Spinal_cord
RPS5	24.1887	Colon
RPS5	27.0551	Lung
RPS5	311.621	stem_cell
RPS5	359.05	Prostate
RPS5	39.6764	Thyroid
RPS5	42.7342	Brain_astrocyte
RPS5	44.5704	Brain_cortex
RPS5	46.3512	Stomach
RPS5	50.564	Heart
RPS5	55.5721	Brain_cerebellum
RPS5	80.6587	Placenta
RPS9	142.029	Eye
RPS9	144.586	Muscle
RPS9	154.419	Breast
RPS9	160.8968	Vascular_endothelium
RPS9	18.99124	Retina
RPS9	246.651	Prostate
RPS9	27.786	Brain_cerebellum
RPS9	32.8557	Spinal_cord
RPS9	36.19475	Liver
RPS9	402.0561	White_Blood_Cells
RPS9	40.5525	Brain_cortex
RPS9	40.9328	Heart
RPS9	46.6191	Brain_astrocyte
RPS9	47.34639	Lung
RPS9	56.013	Placenta
RPS9	56.44025	Colon
RPS9	60.37706	Thyroid
RPS9	674.0737	stem_cell
MRPS15	12.0649	Liver
MRPS15	14.8453	Muscle
MRPS15	19.4723	Prostate

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RBPs	Tag freque	Tissue
MRPS15	19.6614	White_Blood_Cells
MRPS15	25.6914	stem_cell
MRPS15	3.45012	Thyroid
MRPS15	3.65063	Spinal_cord
MRPS15	3.87282	Vascular_endothelium
MRPS15	4.2448	Brain_cortex
MRPS15	4.50918	Lung
MRPS15	4.81562	Heart
MRPS15	7.76986	Brain_astrocyte
RTCD1	13.9577	Vascular_endothelium
RTCD1	19.4246	Brain_astrocyte
RTCD1	22.2945	White_Blood_Cells
RTCD1	3.20251	Brain_cortex
RTCD1	3.71133	Muscle
RTCD1	8.06289	Colon
RTCD1	9.2239	stem_cell
RTCD1	9.36651	Prostate
SNRP70	12.0943	Colon
SNRP70	15.7821	Lung
SNRP70	15.8777	Brain_cerebellum
SNRP70	22.3605	Retina
SNRP70	23.3096	Brain_astrocyte
SNRP70	24.9774	Prostate
SNRP70	25.8366	stem_cell
SNRP70	3.01623	Liver
SNRP70	33.4392	Placenta
SNRP70	3.71133	Muscle
SNRP70	38.4468	Breast
SNRP70	48.8689	Vascular_endothelium
SNRP70	50.3359	White_Blood_Cells
SNRP70	5.45308	Stomach
SNRP70	6.90024	Thyroid
SNRP70	7.30127	Spinal_cord
SNRP70	8.4896	Brain_cortex
SNRP70	9.31337	Eye
SNRPB2	12.4887	Prostate
SNRPB2	13.9577	Vascular_endothelium
SNRPB2	22.1374	stem_cell
SNRPB2	3.01623	Liver
SNRPB2	3.45012	Thyroid
SNRPB2	3.65063	Spinal_cord
SNRPB2	5.45308	Stomach
SNRPB2	6.70455	Retina
SNRPB2	6.76376	Lung
SNRPB2	6.8751	Breast
SNRPB2	6.98503	Eye
SNRPB2	7.93887	Brain_cerebellum
SNRPB2	8.09504	White_Blood_Cells
SAFB1	10.612	Brain_cortex
SAFB1	11.122	Muscle
SAFB1	17.3898	Breast
SAFB1	17.4471	Vascular_endothelium

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RBPs	Tag freque	Tissue
SAFB1	18.5787	White_Blood_Cells
SAFB1	20.4526	stem_cell
SAFB1	20.9551	Eye
SAFB1	3.01623	Liver
SAFB1	3.65063	Spinal_cord
SAFB1	3.96944	Brain_cerebellum
SAFB1	5.45308	Stomach
SAFB1	5.69736	Retina
SAFB1	6.24434	Prostate
SAFB1	7.76986	Brain_astrocyte
SAFB1	7.86805	Placenta
SAFB1	9.01835	Lung
SAFB2	10.122	Prostate
SAFB2	10.54469	stem_cell
SAFB2	10.9519	Spinal_cord
SAFB2	12.81	Brain_cortex
SAFB2	13.5275	Lung
SAFB2	13.7502	Breast
SAFB2	17.51883	Vascular_endothelium
SAFB2	27.26414	Retina
SAFB2	4.65668	Eye
SAFB2	5.45308	Stomach
SAFB2	5.90104	Placenta
SAFB2	6.03245	Liver
SAFB2	7.22343	Heart
SAFB2	7.34659	White_Blood_Cells
SAFB2	8.06289	Colon
SAFB2	8.6253	Thyroid
SAM68	10.9519	Spinal_cord
SAM68	11.6548	Brain_astrocyte
SAM68	12.0943	Colon
SAM68	13.9701	Eye
SAM68	14.4469	Heart
SAM68	15.0811	Liver
SAM68	18.007	White_Blood_Cells
SAM68	19.3641	Vascular_endothelium
SAM68	24.0628	Breast
SAM68	24.5389	Stomach
SAM68	25.8759	Thyroid
SAM68	30.487	Placenta
SAM68	30.9462	Brain_cortex
SAM68	33.8188	Lung
SAM68	3.70734	Muscle
SAM68	37.1139	Prostate
SAM68	45.5788	Retina
SAM68	47.6332	Brain_cerebellum
SAM68	64.0995	stem_cell
SART3	10.9062	Stomach
SART3	3.20251	Brain_cortex
SART3	3.49516	Breast
SART3	6.24434	Prostate
SART3	7.74563	Vascular_endothelium

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RBPs	Tag freque	Tissue
SCAND1	10.3126	Breast
SCAND1	11.9083	Brain_cerebellum
SCAND1	12.0649	Liver
SCAND1	19.0858	Stomach
SCAND1	21.9038	Spinal_cord
SCAND1	22.0184	Placenta
SCAND1	24.1508	Thyroid
SCAND1	24.8005	Lung
SCAND1	26.8182	Retina
SCAND1	29.6907	Muscle
SCAND1	31.6127	White_Blood_Cells
SCAND1	36.0808	Brain_cortex
SCAND1	36.9064	stem_cell
SCAND1	37.2535	Eye
SCAND1	37.466	Prostate
SCAND1	40.3145	Colon
SCAND1	74.5894	Vascular_endothelium
SCAND1	7.76986	Brain_astrocyte
SF1	10.9519	Spinal_cord
SF1	12.0649	Liver
SF1	12.0943	Colon
SF1	18.0367	Lung
SF1	18.5567	Muscle
SF1	18.9757	Thyroid
SF1	19.6701	Placenta
SF1	23.1484	Vascular_endothelium
SF1	23.4336	Retina
SF1	24.5389	Stomach
SF1	26.3617	stem_cell
SF1	27.1945	Brain_astrocyte
SF1	27.9613	Breast
SF1	31.7555	Brain_cerebellum
SF1	34.3439	Prostate
SF1	44.8352	Brain_cortex
SF1	4.65669	Eye
SF1	8.00312	White_Blood_Cells
SF1	9.63124	Heart
SF3B4	17.4758	Breast
SF3B4	20.2376	White_Blood_Cells
SF3B4	20.9551	Eye
SF3B4	3.35227	Retina
SF3B4	3.45012	Thyroid
SF3B4	3.71133	Muscle
SF3B4	4.2448	Brain_cortex
SF3B4	5.08117	Placenta
SF3B4	7.30127	Spinal_cord
SF3B4	7.74563	Vascular_endothelium
SF3B4	7.76986	Brain_astrocyte
SF3B4	8.06289	Colon
SF3B4	9.36651	Prostate
SF3B4	9.73933	stem_cell
SFPQ	12.0943	Colon

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RBPs	Tag freque	Tissue
SFPQ	12.81	Brain_cortex
SFPQ	13.9701	Eye
SFPQ	15.7291	White_Blood_Cells
SFPQ	23.3096	Brain_astrocyte
SFPQ	25.7205	Vascular_endothelium
SFPQ	31.4564	Breast
SFPQ	3.65063	Spinal_cord
SFPQ	3.79824	Retina
SFPQ	41.1027	stem_cell
SFPQ	6.72156	Placenta
SFPQ	6.76376	Lung
SFPQ	7.93887	Brain_cerebellum
SFPQ	9.36651	Prostate
SFRS11	19.8472	Brain_cerebellum
SFRS11	3.01623	Liver
SFRS11	3.37399	Prostate
SFRS11	3.45012	Thyroid
SFRS11	5.1441	Vascular_endothelium
SFRS11	5.7966	Breast
SFRS11	7.33225	stem_cell
SFRS11	9.87898	White_Blood_Cells
SFRS12	10.4683	Vascular_endothelium
SFRS12	13.9806	Breast
SFRS12	15.17547	White_Blood_Cells
SFRS12	15.53973	Brain_astrocyte
SFRS12	3.12217	Prostate
SFRS12	3.20251	Brain_cortex
SFRS12	3.96944	Brain_cerebellum
SFRS12	4.65669	Eye
SFRS12	7.30126	Spinal_cord
SFRS12	7.86805	Placenta
SFRS12	8.17962	Stomach
SFRS15	12.0943	Colon
SFRS15	3.35227	Retina
SFRS15	3.49516	Breast
SFRS15	3.50736	Vascular_endothelium
SFRS15	3.71133	Muscle
SFRS15	3.87199	White_Blood_Cells
SFRS15	3.88493	Brain_astrocyte
SFRS15	6.23641	stem_cell
SFRS15	9.31337	Eye
SFRS1	13.2938	Retina
SFRS1	13.496	Prostate
SFRS1	13.763	White_Blood_Cells
SFRS1	14.0294	Vascular_endothelium
SFRS1	15.5397	Brain_astrocyte
SFRS1	15.8777	Brain_cerebellum
SFRS1	20.1572	Colon
SFRS1	22.918	stem_cell
SFRS1	31.6756	Breast
SFRS1	4.81562	Heart
SFRS1	6.03245	Liver

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RBPs	Tag freque	Tissue
SFRS1	6.40502	Brain_cortex
SFRS1	6.72156	Placenta
SFRS1	6.98503	Eye
SFRS1	7.30127	Spinal_cord
SFRS1	7.41468	Muscle
SFRS1	8.6253	Thyroid
SFRPS2	11.122	Muscle
SFRPS2	14.4469	Heart
SFRPS2	18.0974	Liver
SFRPS2	23.1484	Vascular_endothelium
SFRPS2	23.3096	Brain_astrocyte
SFRPS2	23.7096	White_Blood_Cells
SFRPS2	25.5544	Spinal_cord
SFRPS2	25.7885	Brain_cortex
SFRPS2	30.3659	Prostate
SFRPS2	32.7185	Stomach
SFRPS2	32.7761	Thyroid
SFRPS2	35.1504	Retina
SFRPS2	36.0734	Lung
SFRPS2	3.96944	Brain_cerebellum
SFRPS2	4.03145	Colon
SFRPS2	45.7305	Placenta
SFRPS2	89.8473	Breast
SFRPS2	9.31337	Eye
SFRPS2	94.9023	stem_cell
SFRPS2B	10.3126	Breast
SFRPS2B	12.4887	Prostate
SFRPS2B	3.72675	Retina
SFRPS2B	3.77056	stem_cell
SFRPS2B	3.87199	White_Blood_Cells
SFRPS2B	3.93403	Placenta
SFRPS2B	3.96944	Brain_cerebellum
SFRPS2B	4.50918	Lung
SFRPS2B	5.1577	Brain_cortex
SFRPS2B	7.22343	Heart
SFRPS2B	7.74563	Vascular_endothelium
SFRPS2B	8.6253	Thyroid
SFRPS3	11.122	Muscle
SFRPS3	11.9083	Brain_cerebellum
SFRPS3	197.7721	stem_cell
SFRPS3	27.146	Liver
SFRPS3	31.07946	Brain_astrocyte
SFRPS3	36.11712	Heart
SFRPS3	37.54008	White_Blood_Cells
SFRPS3	37.9824	Retina
SFRPS3	39.5818	Eye
SFRPS3	4.03145	Colon
SFRPS3	44.85154	Thyroid
SFRPS3	45.48012	Prostate
SFRPS3	46.35124	Stomach
SFRPS3	46.4193	Brain_cortex
SFRPS3	47.4583	Spinal_cord

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RBPs	Tag freque	Tissue
SFRPS3	63.8337	Vascular_endothelium
SFRPS3	67.6377	Lung
SFRPS3	73.38002	Placenta
SFRPS3	99.891	Breast
SFRS4	10.122	Prostate
SFRS4	10.4855	Breast
SFRS4	11.8021	Placenta
SFRS4	23.232	White_Blood_Cells
SFRS4	25.6914	stem_cell
SFRS4	3.01623	Liver
SFRS4	3.88493	Brain_astrocyte
SFRS4	4.03145	Colon
SFRS4	7.30127	Spinal_cord
SFRS4	7.73655	Brain_cortex
SFRS4	7.74563	Vascular_endothelium
SFRS4	8.17962	Stomach
SFRPS5	13.9701	Eye
SFRPS5	15.2435	Placenta
SFRPS5	15.4731	Brain_cortex
SFRPS5	15.5397	Brain_astrocyte
SFRPS5	17.2506	Thyroid
SFRPS5	24.1887	Colon
SFRPS5	24.4661	Breast
SFRPS5	28.0995	Prostate
SFRPS5	28.2925	Vascular_endothelium
SFRPS5	3.71133	Muscle
SFRPS5	4.50918	Lung
SFRPS5	4.81562	Heart
SFRPS5	52.5924	stem_cell
SFRPS5	7.30127	Spinal_cord
SFRPS5	7.59648	Retina
SFRPS5	7.93887	Brain_cerebellum
SFRPS5	85.4621	White_Blood_Cells
SFRS6	12.0943	Colon
SFRS6	13.52749	Lung
SFRS6	15.4731	Brain_cortex
SFRS6	15.68364	Placenta
SFRS6	18.6267	Eye
SFRS6	18.97568	Thyroid
SFRS6	19.1018	Vascular_endothelium
SFRS6	19.1804	Breast
SFRS6	19.43959	White_Blood_Cells
SFRS6	21.00558	Retina
SFRS6	21.55455	stem_cell
SFRS6	25.5544	Spinal_cord
SFRS6	27.19453	Brain_astrocyte
SFRS6	3.70734	Muscle
SFRS6	40.89808	Stomach
SFRS6	63.9543	Prostate
SFRS6	7.22343	Heart
SFRS6	9.04868	Liver
SFRS7	10.9519	Spinal_cord

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RBPs	Tag freque	Tissue
SFRS7	11.65479	Brain_astrocyte
SFRS7	14.90923	Breast
SFRS7	15.86269	Prostate
SFRS7	28.29882	Vascular_endothelium
SFRS7	3.01623	Liver
SFRS7	3.71133	Muscle
SFRS7	4.03145	Colon
SFRS7	4.48104	Placenta
SFRS7	45.7322	White_Blood_Cells
SFRS7	4.65669	Eye
SFRS7	4.81562	Heart
SFRS7	5.1577	Brain_cortex
SFRS7	87.1501	stem_cell
SFRPS9	12.81	Brain_cortex
SFRPS9	13.9806	Breast
SFRPS9	15.488	White_Blood_Cells
SFRPS9	23.3096	Brain_astrocyte
SFRPS9	29.8506	stem_cell
SFRPS9	3.65063	Spinal_cord
SFRPS9	3.70734	Muscle
SFRPS9	3.72675	Retina
SFRPS9	4.03145	Colon
SFRPS9	46.4738	Vascular_endothelium
SFRPS9	4.65669	Eye
SFRPS9	6.03245	Liver
SFRPS9	62.4434	Prostate
SFRPS9	6.72156	Placenta
SFRPS9	6.90024	Thyroid
SFRPS9	7.93887	Brain_cerebellum
SFRPS9	9.01835	Lung
SIAHBP1	10.122	Prostate
SIAHBP1	10.9519	Spinal_cord
SIAHBP1	11.6548	Brain_astrocyte
SIAHBP1	14.8294	Muscle
SIAHBP1	19.7973	Breast
SIAHBP1	19.8472	Brain_cerebellum
SIAHBP1	21.224	Brain_cortex
SIAHBP1	23.5187	stem_cell
SIAHBP1	28.0589	Vascular_endothelium
SIAHBP1	34.9251	Eye
SIAHBP1	4.03145	Colon
SIAHBP1	4.50918	Lung
SIAHBP1	4.81562	Heart
SIAHBP1	5.08117	Placenta
SIAHBP1	5.89843	White_Blood_Cells
SLBP	11.2729	Lung
SLBP	11.6185	Vascular_endothelium
SLBP	12.0649	Liver
SLBP	13.9088	Prostate
SLBP	15.2435	Placenta
SLBP	15.4731	Brain_cortex
SLBP	16.4411	stem_cell

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RBPs	Tag freque	Tissue
SLBP	22.4258	Thyroid
SLBP	25.7131	White_Blood_Cells
SLBP	3.71133	Muscle
SLBP	3.88493	Brain_astrocyte
SLBP	3.96944	Brain_cerebellum
SLBP	5.45308	Stomach
SLBP	6.98503	Eye
SLBP	7.30127	Spinal_cord
SLBP	8.6949	Breast
SLBP	9.764	Retina
SLTM	10.81721	Breast
SLTM	15.30822	Retina
SLTM	16.34963	Vascular_endothelium
SLTM	16.86999	Prostate
SLTM	20.90583	stem_cell
SLTM	22.58026	White_Blood_Cells
SLTM	3.71133	Muscle
SLTM	3.88493	Brain_astrocyte
SLTM	3.93403	Placenta
SLTM	3.96944	Brain_cerebellum
SLTM	4.03145	Colon
SLTM	5.17518	Thyroid
SLTM	5.78136	Brain_cortex
SLTM	6.03245	Liver
SLTM	7.30127	Spinal_cord
SNRPA	11.6185	Vascular_endothelium
SNRPA	12.0943	Colon
SNRPA	19.36	White_Blood_Cells
SNRPA	31.6341	stem_cell
SNRPA	3.65063	Spinal_cord
SNRPA	3.93403	Placenta
SNRPA	5.8584	Retina
SNRPA	6.8751	Breast
SNRPA	7.42267	Muscle
SNRPA	7.76986	Brain_astrocyte
SNRPA	7.93887	Brain_cerebellum
SNRPA	8.36622	Prostate
SNRPA	9.31337	Eye
SNRPA	9.60753	Brain_cortex
SON	12.03905	Heart
SON	12.0649	Liver
SON	16.12578	Colon
SON	18.67561	Brain_cortex
SON	19.42466	Brain_astrocyte
SON	19.84717	Brain_cerebellum
SON	19.99207	Prostate
SON	22.26797	Muscle
SON	23.46587	Retina
SON	24.8005	Lung
SON	27.96132	Breast
SON	29.20503	Spinal_cord
SON	29.32606	Thyroid

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RBPs	Tag freque	Tissue
SON	29.99194	Stomach
SON	31.35254	White_Blood_Cells
SON	35.0578	stem_cell
SON	36.36883	Vascular_endothelium
SON	37.80872	Placenta
SON	6.98503	Eye
SPF45	10.122	Prostate
SPF45	10.2882	Vascular_endothelium
SPF45	11.0199	White_Blood_Cells
SPF45	11.6417	Eye
SPF45	13.5498	Placenta
SPF45	13.6327	Stomach
SPF45	18.0367	Lung
SPF45	20.6253	Breast
SPF45	26.2962	stem_cell
SPF45	3.20251	Brain_cortex
SPF45	3.71133	Muscle
SPF45	3.88493	Brain_astrocyte
SPF45	3.96944	Brain_cerebellum
SPF45	4.81562	Heart
SPF45	5.17518	Thyroid
SPF45	6.03245	Liver
SPF45	6.70455	Retina
SPF45	7.30127	Spinal_cord
SR140	10.9519	Spinal_cord
SR140	11.23507	Vascular_endothelium
SR140	11.65479	Brain_astrocyte
SR140	17.07657	stem_cell
SR140	18.71774	White_Blood_Cells
SR140	3.01623	Liver
SR140	3.70734	Muscle
SR140	3.93403	Placenta
SR140	5.17518	Thyroid
SR140	6.16273	Prostate
SR140	6.40502	Brain_cortex
SR140	7.91891	Breast
SR140	8.65735	Retina
SRP14	103.498	Retina
SRP14	112.037	stem_cell
SRP14	126.364	Brain_cortex
SRP14	16.2984	Eye
SRP14	170.113	Prostate
SRP14	25.9793	Muscle
SRP14	31.7555	Brain_cerebellum
SRP14	34.8479	White_Blood_Cells
SRP14	38.8493	Brain_astrocyte
SRP14	42.601	Vascular_endothelium
SRP14	55.8929	Placenta
SRP14	62.7105	Stomach
SRP14	66.357	Liver
SRP14	69.5592	Breast
SRP14	8.06289	Colon

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RBPs	Tag freque	Tissue
SRP14	93.1532	Thyroid
SRP14	96.3124	Heart
SRP14	98.5671	Spinal_cord
SRP14	99.2019	Lung
SRP68	12.31149	Prostate
SRP68	13.92305	Breast
SRP68	16.44334	stem_cell
SRP68	18.09733	Liver
SRP68	3.65063	Spinal_cord
SRP68	3.71133	Muscle
SRP68	6.90024	Thyroid
SRP68	6.97885	Vascular_endothelium
SRP68	7.59648	Retina
SRP68	7.76986	Brain_astrocyte
SRP68	8.06289	Colon
SRP68	9.37035	White_Blood_Cells
SRP68	9.60753	Brain_cortex
SRP72	11.0199	White_Blood_Cells
SRP72	11.6548	Brain_astrocyte
SRP72	27.1097	Vascular_endothelium
SRP72	3.49516	Breast
SRP72	6.24434	Prostate
SRP72	6.33195	stem_cell
SRP72	6.40502	Brain_cortex
SRP72	7.42267	Muscle
SRrp35	3.37399	Prostate
SRrp35	3.48943	Vascular_endothelium
STAU1	11.2729	Lung
STAU1	12.8602	Vascular_endothelium
STAU1	13.8005	Thyroid
STAU1	15.0811	Liver
STAU1	15.4731	Brain_cortex
STAU1	16.9372	Placenta
STAU1	17.0921	Retina
STAU1	18.8267	stem_cell
STAU1	19.0858	Stomach
STAU1	20.971	Breast
STAU1	23.6179	Prostate
STAU1	29.2051	Spinal_cord
STAU1	4.65669	Eye
STAU1	7.93887	Brain_cerebellum
STAU1	8.06289	Colon
STAU1	8.09504	White_Blood_Cells
STAU2	11.60944	Retina
STAU2	12.29343	stem_cell
STAU2	14.8453	Muscle
STAU2	15.5397	Brain_astrocyte
STAU2	26.6246	Brain_cortex
STAU2	3.01623	Liver
STAU2	3.65063	Spinal_cord
STAU2	3.87282	Vascular_endothelium
STAU2	3.96944	Brain_cerebellum

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RBPs	Tag freque	Tissue
STAU2	4.50918	Lung
STAU2	4.65669	Eye
STAU2	5.17518	Thyroid
STAU2	7.86805	Placenta
STAU2	7.91892	Breast
STAU2	8.68568	Prostate
STAU2	9.63124	Heart
STRBP	15.4148	stem_cell
STRBP	3.50736	Vascular_endothelium
STRBP	7.43149	White_Blood_Cells
SUPV3L1	3.01623	Liver
SUPV3L1	4.03145	Colon
SUPV3L1	5.1577	Brain_cortex
SUPV3L1	5.27235	stem_cell
SUPV3L1	6.74798	Prostate
SUPV3L1	6.98503	Eye
SUPV3L1	6.99032	Breast
SUPV3L1	7.74398	White_Blood_Cells
SURF6	3.01623	Liver
SURF6	3.37399	Prostate
SURF6	3.45012	Thyroid
SURF6	3.49516	Breast
SURF6	3.88493	Brain_astrocyte
SURF6	5.49919	stem_cell
SURF6	7.74398	White_Blood_Cells
GARS	104.893	Brain_astrocyte
GARS	11.134	Muscle
GARS	11.2729	Lung
GARS	11.9083	Brain_cerebellum
GARS	15.7361	Placenta
GARS	19.36	White_Blood_Cells
GARS	19.528	Retina
GARS	32.0251	Brain_cortex
GARS	3.45012	Thyroid
GARS	34.9251	Eye
GARS	37.466	Prostate
GARS	4.03145	Colon
GARS	45.4371	Breast
GARS	49.9147	stem_cell
GARS	55.8308	Vascular_endothelium
GARS	68.1635	Stomach
GARS	7.22343	Heart
GARS	7.30127	Spinal_cord
HARS	10.122	Prostate
HARS	11.134	Muscle
HARS	17.5368	Vascular_endothelium
HARS	18.5787	White_Blood_Cells
HARS	19.4246	Brain_astrocyte
HARS	19.8472	Brain_cerebellum
HARS	22.4176	Brain_cortex
HARS	3.01623	Liver
HARS	3.38745	Placenta

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RBPs	Tag freque	Tissue
HARS	3.49516	Breast
HARS	8.07443	stem_cell
MARS	11.0199	White_Blood_Cells
MARS	15.5397	Brain_astrocyte
MARS	15.8378	Breast
MARS	17.4471	Vascular_endothelium
MARS	21.0802	stem_cell
MARS	3.35227	Retina
MARS	3.65063	Spinal_cord
MARS	3.96944	Brain_cerebellum
MARS	4.03145	Colon
MARS	6.40502	Brain_cortex
MARS	7.41468	Muscle
MARS	9.31337	Eye
MARS	9.36651	Prostate
SYNJ2	11.8561	Placenta
SYNJ2	21.9038	Spinal_cord
SYNJ2	3.6733	White_Blood_Cells
SYNJ2	3.87282	Vascular_endothelium
SYNJ2	3.9056	Retina
SYNJ2	5.7966	Breast
SYNJ2	6.40502	Brain_cortex
SYNJ2	6.90024	Thyroid
SYNJ2	7.76986	Brain_astrocyte
SYWC	108.7294	Vascular_endothelium
SYWC	11.134	Muscle
SYWC	11.6417	Eye
SYWC	11.87837	Breast
SYWC	12.0943	Colon
SYWC	122.4993	White_Blood_Cells
SYWC	14.44686	Heart
SYWC	14.60253	Spinal_cord
SYWC	22.0664	Retina
SYWC	24.5389	Stomach
SYWC	29.33818	stem_cell
SYWC	32.91611	Brain_cortex
SYWC	37.95134	Thyroid
SYWC	38.9557	Placenta
SYWC	50.50403	Brain_astrocyte
SYWC	76.656	Lung
SYWC	8.36622	Prostate
TACC1	10.0039	White_Blood_Cells
TACC1	16.8547	Heart
TACC1	17.1877	Breast
TACC1	21.8123	Stomach
TACC1	22.1374	stem_cell
TACC1	22.3605	Retina
TACC1	24.1887	Colon
TACC1	3.01623	Liver
TACC1	31.5662	Vascular_endothelium
TACC1	46.5766	Thyroid
TACC1	57.0434	Placenta

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RBPs	Tag freque	Tissue
TACC1	58.6193	Lung
TACC1	62.0608	Spinal_cord
TACC1	64.4712	Brain_cortex
TACC1	67.4798	Prostate
TACC1	6.98503	Eye
TACC1	7.42267	Muscle
TACC1	7.93887	Brain_cerebellum
TACC2	15.2435	Placenta
TACC2	3.72675	Retina
TACC2	3.95946	Breast
TACC2	8.06289	Colon
TARDBP	12.8602	Vascular_endothelium
TARDBP	13.4431	Placenta
TARDBP	15.52552	Thyroid
TARDBP	15.8777	Brain_cerebellum
TARDBP	16.86999	Prostate
TARDBP	18.6267	Eye
TARDBP	20.97096	Breast
TARDBP	24.42651	Brain_cortex
TARDBP	24.53884	Stomach
TARDBP	31.0794	Brain_astrocyte
TARDBP	34.3792	White_Blood_Cells
TARDBP	3.71133	Muscle
TARDBP	51.70272	stem_cell
TARDBP	5.69736	Retina
TARDBP	7.22343	Heart
TARDBP	9.01835	Lung
TARB1	11.8021	Placenta
TARB1	11.9083	Brain_cerebellum
TARB1	15.4731	Brain_cortex
TARB1	3.37399	Prostate
TARB1	3.65063	Spinal_cord
TARB1	3.95946	Breast
TARB1	5.1441	Vascular_endothelium
TARB1	6.59044	stem_cell
TARB1	7.74398	White_Blood_Cells
TARB1	9.764	Retina
TDRKH	15.6836	Placenta
TDRKH	3.20251	Brain_cortex
TDRKH	3.72675	Retina
TEP1	10.122	Prostate
TEP1	10.9062	Stomach
TEP1	3.65063	Spinal_cord
TEP1	3.71133	Muscle
TEP1	3.79824	Retina
TEP1	3.88493	Brain_astrocyte
TEP1	6.99032	Breast
TEP1	8.00312	White_Blood_Cells
THOC4	10.909	stem_cell
THOC4	11.5932	Breast
THOC4	13.9088	Prostate
THOC4	3.48943	Vascular_endothelium

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RBPs	Tag freque	Tissue
THOC4	3.70734	Muscle
THOC4	3.87199	White_Blood_Cells
THOC4	4.03145	Colon
THOC4	4.2448	Brain_cortex
THOC4	4.65669	Eye
THOC4	5.08117	Placenta
THOC4	5.8584	Retina
THOC4	7.76986	Brain_astrocyte
TIA1	10.1623	Placenta
TIA1	10.5221	Vascular_endothelium
TIA1	11.127	Prostate
TIA1	13.45185	stem_cell
TIA1	14.6025	Spinal_cord
TIA1	17.39635	Retina
TIA1	3.43755	Breast
TIA1	3.70734	Muscle
TIA1	3.88493	Brain_astrocyte
TIA1	3.96944	Brain_cerebellum
TIA1	6.90024	Thyroid
TIA1	7.74398	White_Blood_Cells
TIAR	14.6025	Spinal_cord
TIAR	15.4323	Vascular_endothelium
TIAR	17.0548	Brain_cortex
TIAR	17.5752	Retina
TIAR	17.70966	Breast
TIAR	20.7007	Thyroid
TIAR	22.2541	Prostate
TIAR	22.29445	White_Blood_Cells
TIAR	22.54589	Lung
TIAR	31.27516	stem_cell
TIAR	4.81562	Heart
TIAR	51.08501	Placenta
TIAR	5.45308	Stomach
TIAR	6.03245	Liver
TIAR	7.42267	Muscle
TIAR	7.76986	Brain_astrocyte
TIAR	7.93888	Brain_cerebellum
TNRC6A	11.1803	Retina
TNRC6A	11.6185	Vascular_endothelium
TNRC6A	21.9038	Spinal_cord
TNRC6A	3.37399	Prostate
TNRC6A	3.45012	Thyroid
TNRC6A	3.49516	Breast
TNRC6A	3.71133	Muscle
TNRC6A	3.88493	Brain_astrocyte
TNRC6A	6.03245	Liver
TNRC6A	6.40502	Brain_cortex
TNRC6A	6.98503	Eye
TNRC6A	7.93887	Brain_cerebellum
TNRC6A	8.46862	Placenta
TNRC6A	9.22661	stem_cell
TNRC6C	3.49516	Breast

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RBPs	Tag freque	Tissue
TNRC4	3.79824	Retina
TNRC4	8.4896	Brain_cortex
TRA2A	10.5221	Vascular_endothelium
TRA2A	10.9062	Stomach
TRA2A	11.1472	White_Blood_Cells
TRA2A	12.0754	Thyroid
TRA2A	12.4887	Prostate
TRA2A	13.5275	Lung
TRA2A	15.2435	Placenta
TRA2A	20.6308	Brain_cortex
TRA2A	21.0894	stem_cell
TRA2A	25.5544	Spinal_cord
TRA2A	31.4564	Breast
TRA2A	36.875	Retina
TRA2A	3.70734	Muscle
TRA2A	3.96944	Brain_cerebellum
TRA2A	4.65669	Eye
TRA2A	6.03245	Liver
TRA2A	8.06289	Colon
TRA2B	10.9519	Spinal_cord
TRA2B	12.06491	Liver
TRA2B	14.09633	Placenta
TRA2B	18.03666	Lung
TRA2B	18.05931	Brain_cortex
TRA2B	19.84714	Brain_cerebellum
TRA2B	21.81228	Stomach
TRA2B	23.0042	Retina
TRA2B	28.78973	Vascular_endothelium
TRA2B	29.5986	White_Blood_Cells
TRA2B	3.71133	Muscle
TRA2B	4.03145	Colon
TRA2B	41.40144	Thyroid
TRA2B	4.65669	Eye
TRA2B	52.6797	stem_cell
TRA2B	54.0861	Breast
TRA2B	7.76986	Brain_astrocyte
TRA2B	9.87015	Prostate
TRBP2	13.748	stem_cell
TRBP2	3.96944	Brain_cerebellum
TRBP2	6.74798	Prostate
TRBP2	6.98503	Eye
TRBP2	7.34659	White_Blood_Cells
TRBP2	7.74563	Vascular_endothelium
TRSPAP1	11.127	Prostate
TRSPAP1	12.0391	Heart
TRSPAP1	13.7691	Placenta
TRSPAP1	3.01623	Liver
TRSPAP1	3.49516	Breast
TRSPAP1	3.72675	Retina
TRSPAP1	3.96944	Brain_cerebellum
TRSPAP1	4.2448	Brain_cortex
TRSPAP1	6.98503	Eye

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RBPs	Tag freque	Tissue
TRSPAP1	7.34659	White_Blood_Cells
TRSPAP1	7.74563	Vascular_endothelium
TRSPAP1	8.17962	Stomach
TRSPAP1	9.55877	stem_cell
TUT1	12.0649	Liver
TUT1	13.1809	stem_cell
TUT1	13.2938	Retina
TUT1	13.5498	Placenta
TUT1	3.65063	Spinal_cord
TUT1	3.71133	Muscle
TUT1	3.88493	Brain_astrocyte
TUT1	3.96944	Brain_cerebellum
TUT1	4.65669	Eye
TUT1	4.81562	Heart
TUT1	6.40502	Brain_cortex
TUT1	6.97885	Vascular_endothelium
TUT1	7.91891	Breast
TUT1	8.17962	Stomach
U2AF1	12.0391	Heart
U2AF1	12.0649	Liver
U2AF1	12.81	Brain_cortex
U2AF1	13.8005	Thyroid
U2AF1	13.9701	Eye
U2AF1	14.8453	Muscle
U2AF1	15.5397	Brain_astrocyte
U2AF1	15.7821	Lung
U2AF1	16.1258	Colon
U2AF1	18.2532	Spinal_cord
U2AF1	23.232	White_Blood_Cells
U2AF1	24.426	Vascular_endothelium
U2AF1	28.0995	Prostate
U2AF1	30.1705	Retina
U2AF1	32.1808	Placenta
U2AF1	44.6881	Breast
U2AF1	5.45308	Stomach
U2AF1	63.0879	stem_cell
U2AF1	7.93887	Brain_cerebellum
U2AF1L4	11.9083	Brain_cerebellum
U2AF1L4	12.8942	Brain_cortex
U2AF1L4	14.863	White_Blood_Cells
U2AF1L4	18.9912	Retina
U2AF1L4	3.01623	Liver
U2AF1L4	3.45012	Thyroid
U2AF1L4	3.87282	Vascular_endothelium
U2AF1L4	3.93403	Placenta
U2AF1L4	6.74798	Prostate
U2AF1L4	7.42267	Muscle
U2AF1L4	7.76986	Brain_astrocyte
U2AF1L4	7.90852	stem_cell
U2AF2	15.4323	Vascular_endothelium
U2AF2	15.8378	Breast
U2AF2	18.5047	stem_cell

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RBPs	Tag freque	Tissue
U2AF2	30.9759	White_Blood_Cells
U2AF2	3.88493	Brain_astrocyte
U2AF2	3.93403	Placenta
U2AF2	6.24434	Prostate
HDLBP	104.893	Brain_astrocyte
HDLBP	14.6932	White_Blood_Cells
HDLBP	16.1258	Colon
HDLBP	20.7007	Thyroid
HDLBP	25.5544	Spinal_cord
HDLBP	31.3015	Heart
HDLBP	32.5968	Eye
HDLBP	32.7185	Stomach
HDLBP	33.3661	Muscle
HDLBP	33.8548	stem_cell
HDLBP	35.2276	Brain_cortex
HDLBP	36.875	Retina
HDLBP	3.96944	Brain_cerebellum
HDLBP	51.2759	Liver
HDLBP	51.8555	Lung
HDLBP	60.7318	Prostate
HDLBP	65.8379	Vascular_endothelium
HDLBP	68.8455	Placenta
HDLBP	71.2702	Breast
YBOX1	107.6286	Muscle
YBOX1	14.5602	Brain_cortex
YBOX1	153.0986	Breast
YBOX1	24.12983	Liver
YBOX1	25.5544	Spinal_cord
YBOX1	286.9061	stem_cell
YBOX1	31.0511	Thyroid
YBOX1	38.0888	Placenta
YBOX1	38.1716	Stomach
YBOX1	56.3647	Lung
YBOX1	57.0272	Retina
YBOX1	64.5031	Colon
YBOX1	75.74385	White_Blood_Cells
YBOX1	76.42415	Vascular_endothelium
YBOX1	7.93887	Brain_cerebellum
YBOX1	85.4684	Brain_astrocyte
YBOX1	93.6651	Prostate
YBOX1	96.31242	Heart
YBOX1	97.7904	Eye
YBOX2	18.5047	stem_cell
YBOX2	4.03145	Colon
YBOX2	7.45351	Retina
YBOX2	7.91891	Breast
YBOX2	8.17962	Stomach
ZCRB1	3.01623	Liver
ZCRB1	3.87199	White_Blood_Cells
ZCRB1	3.87282	Vascular_endothelium
ZCRB1	3.93424	Placenta
ZCRB1	7.76986	Brain_astrocyte

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<u>RBPs</u>	<u>Tag freque</u>	<u>Tissue</u>
ZNF638	10.9519	Spinal_cord
ZNF638	11.7168	Retina
ZNF638	11.8548	White_Blood_Cells
ZNF638	11.8561	Placenta
ZNF638	12.8602	Vascular_endothelium
ZNF638	15.4731	Brain_cortex
ZNF638	15.7821	Lung
ZNF638	27.6798	stem_cell
ZNF638	3.37399	Prostate
ZNF638	3.88493	Brain_astrocyte
ZNF638	3.96944	Brain_cerebellum
ZNF638	4.65669	Eye
ZNF638	6.90024	Thyroid
ZNF638	7.41468	Muscle
ZNF638	7.91891	Breast
ZNF638	8.17962	Stomach
ZNF638	9.04868	Liver

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RBPs	Tag frequency	Tissue
APOBEC3F	108.261	peripheral_blood_lymphocytes
APOBEC3F	10.9871	adrenal_gland
APOBEC3F	11.6022	pituitary_gland
APOBEC3F	17.226	uterus
APOBEC3F	18.9629	Monocytes
APOBEC3F	32.6177	trachea
APOBEC3F	3.70365	salivary_gland
APOBEC3F	44.8974	small_intestine
APOBEC3F	47.2117	testis
APOBEC3F	52.1979	thymus
APOBEC3F	6.9185	bone_marrow
APOBEC1	4.48974	small_intestine
ADAT1	13.1574	testis
AKAP1	10.9724	peripheral_blood_lymphocytes
AKAP1	115.985	small_intestine
AKAP1	129.506	bladder
AKAP1	15.6594	thymus
AKAP1	16.1144	adrenal_gland
AKAP1	24.2148	bone_marrow
AKAP1	24.7668	testis
AKAP1	25.1848	salivary_gland
AKAP1	25.7895	Monocytes
AKAP1	44.9375	uterus
AKAP1	50.0344	pituitary_gland
AKAP1	55.3742	trachea
AKAP1	84.8733	kidney
AKAP1	92.3804	skeletal_muscle
ANKHD1	105.1416	thymus
ANKHD1	105.1447	pituitary_gland
ANKHD1	16.82427	peripheral_blood_lymphocytes
ANKHD1	18.5182	salivary_gland
ANKHD1	32.20517	uterus
ANKHD1	3.45925	bone_marrow
ANKHD1	4.10677	kidney
ANKHD1	54.20299	adrenal_gland
ANKHD1	71.08756	small_intestine
ANKHD1	74.33448	Monocytes
ANKHD1	83.5879	testis
ANKHD1	85.4083	skeletal_muscle
ANKHD1	88.7505	trachea
ANKHD1	89.831	bladder
ANKRD17	16.68731	Monocytes
ANKRD17	17.1116	kidney
ANKRD17	22.96263	salivary_gland
ANKRD17	25.6022	peripheral_blood_lymphocytes
ANKRD17	30.9585	testis
ANKRD17	36.6237	adrenal_gland
ANKRD17	38.3466	skeletal_muscle

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RBPs	Tag frequency	Tissue
ANKRD17	47.16134	bladder
ANKRD17	50.1802	uterus
ANKRD17	6.22665	bone_marrow
ANKRD17	70.84	thymus
ANKRD17	77.3722	trachea
ANKRD17	89.7949	small_intestine
ANKRD17	97.1682	pituitary_gland
ASCC1	4.64377	testis
ASCC1	6.58342	peripheral_blood_lymphocytes
BOLL	235.2848	testis
BOLL	6.1006	skeletal_muscle
BRUNOL6	14.2302	uterus
BRUNOL6	14.7053	testis
BRUNOL6	28.336	thymus
BRUNOL6	5.30986	trachea
CCBL2	147.928	pituitary_gland
CCBL2	17.7775	salivary_gland
CCBL2	177.805	bone_marrow
CCBL2	21.7878	skeletal_muscle
CCBL2	263.976	trachea
CCBL2	287.6	uterus
CCBL2	324.373	thymus
CCBL2	32.5064	testis
CCBL2	38.769	peripheral_blood_lymphocytes
CCBL2	51.2731	adrenal_gland
CCBL2	83.4367	Monocytes
CCBL2	84.5569	small_intestine
CCBL2	97.8781	kidney
CCBL2	99.5627	bladder
RCAN3	11.9775	bladder
RCAN3	15.5553	salivary_gland
RCAN3	19.0188	peripheral_blood_lymphocytes
RCAN3	22.4792	pituitary_gland
RCAN3	26.5493	trachea
RCAN3	28.336	thymus
RCAN3	3.09585	testis
RCAN3	4.49375	uterus
RCAN3	5.98633	small_intestine
RCAN3	9.8607	Monocytes
CDKN2AIP	18.1284	pituitary_gland
CDKN2AIP	3.86981	testis
CDKN2AIP	73.2473	adrenal_gland
CIRBP	106.066	peripheral_blood_lymphocytes
CIRBP	152.354	adrenal_gland
CIRBP	192.482	uterus
CIRBP	21.4811	salivary_gland
CIRBP	21.7878	skeletal_muscle
CIRBP	26.548	Monocytes

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RBPs	Tag frequency	Tissue
CIRBP	34.9076	kidney
CIRBP	37.4296	bladder
CIRBP	38.0518	bone_marrow
CIRBP	505.42	pituitary_gland
CIRBP	66.366	thymus
CIRBP	68.8427	small_intestine
CIRBP	71.2045	testis
CIRBP	82.6821	trachea
CNOT4	12.2012	skeletal_muscle
CNOT4	17.959	small_intestine
CNOT4	28.2803	pituitary_gland
CNOT4	36.4104	trachea
CNOT4	36.5386	thymus
CNOT4	53.176	uterus
CNOT4	58.8211	testis
CNOT4	6.59226	adrenal_gland
CNOT4	9.10218	Monocytes
CNOT4	9.7317	bladder
CPEB2	217.483	testis
CPEB2	3.0342	trachea
CPEB2	5.22908	skeletal_muscle
CPEB4	103.279	adrenal_gland
CPEB4	11.1853	thymus
CPEB4	11.2344	uterus
CPEB4	11.3777	Monocytes
CPEB4	14.8157	skeletal_muscle
CPEB4	16.4271	kidney
CPEB4	46.4087	pituitary_gland
CPEB4	5.92583	salivary_gland
CPEB4	6.82696	trachea
CPEB4	6.9185	bone_marrow
CPEB4	7.73962	testis
CPSF5	106.776	kidney
CPSF5	113.783	trachea
CPSF5	212.52	thymus
CPSF5	248.533	bladder
CPSF5	248.654	uterus
CPSF5	31.4541	peripheral_blood_lymphocytes
CPSF5	428.555	pituitary_gland
CPSF5	43.703	salivary_gland
CPSF5	45.6621	bone_marrow
CPSF5	50.5478	skeletal_muscle
CPSF5	61.5277	adrenal_gland
CPSF5	74.8291	small_intestine
CPSF5	85.9098	testis
CPSF5	89.5048	Monocytes
CPSF6	12.5924	salivary_gland
CPSF6	19.1733	skeletal_muscle

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RBPs	Tag frequency	Tissue
CPSF6	20.5092	adrenal_gland
CPSF6	23.9562	kidney
CPSF6	29.2094	uterus
CPSF6	35.9762	bone_marrow
CPSF6	37.1502	testis
CPSF6	47.0279	Monocytes
CPSF6	47.9099	bladder
CPSF6	50.7595	pituitary_gland
CPSF6	50.8229	trachea
CPSF6	54.6252	small_intestine
CPSF6	65.6203	thymus
CSTF2	5.30986	trachea
CSTF2	5.85978	adrenal_gland
CSTF2	6.1917	testis
CSTF2	7.97649	pituitary_gland
CSTF2	8.94822	thymus
CSTF3	3.74479	uterus
CSTF3	5.98633	small_intestine
CSTF2T	11.97748	bladder
CSTF2T	12.38339	testis
CSTF2T	12.5924	salivary_gland
CSTF2T	16.6881	trachea
CSTF2T	23.9295	pituitary_gland
CSTF2T	28.33605	thymus
CSTF2T	31.42816	small_intestine
CSTF2T	32.9613	adrenal_gland
CSTF2T	37.6454	kidney
CSTF2T	5.22908	skeletal_muscle
CSTF2T	6.06812	Monocytes
CSTF2T	6.9185	bone_marrow
CSTF2T	9.50939	peripheral_blood_lymphocytes
CSTF2T	9.73646	uterus
CUGBP1	4.55109	Monocytes
CUGBP1	9.28754	testis
CUGBP2	104.80278	bladder
CUGBP2	109.61553	thymus
CUGBP2	12.3203	kidney
CUGBP2	14.4118	Monocytes
CUGBP2	19.34908	testis
CUGBP2	20.0637	bone_marrow
CUGBP2	21.2133	peripheral_blood_lymphocytes
CUGBP2	31.3745	skeletal_muscle
CUGBP2	33.6938	adrenal_gland
CUGBP2	3.70365	salivary_gland
CUGBP2	40.443758	uterus
CUGBP2	43.4009	small_intestine
CUGBP2	44.7545	trachea
CUGBP2	69.613	pituitary_gland

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RBPs	Tag frequency	Tissue
DDX17	17.2962	bone_marrow
DDX17	18.6421	thymus
DDX17	3.09585	testis
DDX17	5.23804	small_intestine
DDX17	5.30986	trachea
DDX17	6.74062	uterus
DDX17	7.48592	bladder
DDX17	8.78968	adrenal_gland
DDX17	9.62948	salivary_gland
DDX53	14.7053	testis
DGCR8	3.72842	thymus
DHX9	104.054	bladder
DHX9	106.776	kidney
DHX9	113.019	Monocytes
DHX9	119.19	testis
DHX9	121.331	uterus
DHX9	148.056	bone_marrow
DHX9	152.469	trachea
DHX9	159.679	adrenal_gland
DHX9	165.542	thymus
DHX9	16.5588	skeletal_muscle
DHX9	201.29	small_intestine
DHX9	26.6663	salivary_gland
DHX9	355.317	pituitary_gland
DHX9	51.9359	peripheral_blood_lymphocytes
DICER	117.818	thymus
DICER	127.261	bladder
DICER	143.565	adrenal_gland
DICER	147.545	uterus
DICER	17.0272	testis
DICER	240.02	pituitary_gland
DICER	285.202	Monocytes
DICER	30.4414	bone_marrow
DICER	40.7401	salivary_gland
DICER	45.6457	small_intestine
DICER	54.8619	peripheral_blood_lymphocytes
DICER	63.6205	skeletal_muscle
DICER	69.1307	kidney
DICER	94.0603	trachea
DNAJC17	12.3273	pituitary_gland
DNAJC17	4.49375	uterus
DNAJC17	4.55131	trachea
DNAJC17	4.84295	bone_marrow
DNAJC17	5.21979	thymus
DNAJC17	55.6679	adrenal_gland
DKC1	128.706	small_intestine
DKC1	13.9442	skeletal_muscle
DKC1	19.0188	peripheral_blood_lymphocytes

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RBPs	Tag frequency	Tissue
DKC1	19.259	salivary_gland
DKC1	20.5339	kidney
DKC1	25.4521	bladder
DKC1	26.5493	trachea
DKC1	35.6502	Monocytes
DKC1	47.7377	bone_marrow
DKC1	57.2732	testis
DKC1	61.6365	pituitary_gland
DKC1	6.74062	uterus
DKC1	70.8401	thymus
DKC1	76.1772	adrenal_gland
DKFZp686F02.	119.965	peripheral_blood_lymphocytes
DKFZp686F02.	13.4223	thymus
DKFZp686F02.	145.168	small_intestine
DKFZp686F02.	35.1838	bladder
DKFZp686F02.	3.62568	pituitary_gland
DKFZp686F02.	4.1511	bone_marrow
DKFZp686F02.	44.8898	testis
DKFZp686F02.	504.449	kidney
DKFZp686F02.	6.74062	uterus
DKFZp686F02.	742.586	Monocytes
DKFZp686F02.	752.982	adrenal_gland
DKFZp686F02.	96.336	trachea
DKFZp686F02.	98.517	salivary_gland
DND1	3.42231	kidney
DND1	75.0743	testis
DRB1	3.62568	pituitary_gland
DRB1	5.41773	testis
DSRAD	12.2012	skeletal_muscle
DSRAD	13.1668	peripheral_blood_lymphocytes
DSRAD	22.7565	trachea
DSRAD	30.7639	adrenal_gland
DSRAD	31.8576	Monocytes
DSRAD	34.3015	thymus
DSRAD	37.4145	small_intestine
DSRAD	4.44438	salivary_gland
DSRAD	45.6638	testis
DSRAD	50.5051	bone_marrow
DSRAD	74.689	pituitary_gland
DSRAD	88.3771	uterus
DSRAD	9.7317	bladder
DUSP11	14.9792	uterus
DUSP11	3.09585	testis
DUSP11	3.65746	peripheral_blood_lymphocytes
DUSP11	3.72842	thymus
DUSP11	4.35757	skeletal_muscle
DUSP11	5.85978	adrenal_gland
DUS2L	11.9833	uterus

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RBPs	Tag frequency	Tissue
DUS2L	12.7209	small_intestine
DUS2L	16.469	bladder
DUS2L	19.7214	Monocytes
DUS2L	7.25136	pituitary_gland
DUS2L	7.52908	kidney
DUS2L	88.2317	testis
DUS2L	9.6859	bone_marrow
DUS2L	9.6939	thymus
E2AK2	3.03406	Monocytes
E2AK2	3.66237	adrenal_gland
ELAV1	14.6298	peripheral_blood_lymphocytes
ELAV1	15.0582	kidney
ELAV1	18.8535	pituitary_gland
ELAV1	19.7223	trachea
ELAV1	25.4419	small_intestine
ELAV1	26.369	adrenal_gland
ELAV1	32.9542	uterus
ELAV1	4.84295	bone_marrow
ELAV1	6.71116	thymus
ELAV1	7.58515	Monocytes
ELAV1	7.84362	skeletal_muscle
ELAV1	9.28754	testis
ELAVL3	50.0344	pituitary_gland
ELAVL4	17.4033	pituitary_gland
ELAVL4	5.23804	small_intestine
ELAVL4	7.32473	adrenal_gland
EPRS	16.4771	uterus
EPRS	18.2052	trachea
EPRS	20.0448	skeletal_muscle
EPRS	24.6936	small_intestine
EPRS	24.8707	peripheral_blood_lymphocytes
EPRS	3.70365	salivary_gland
EPRS	4.47411	thymus
EPRS	6.59226	adrenal_gland
EPRS	76.1393	pituitary_gland
ERAL1	12.6766	thymus
ERAL1	14.6495	adrenal_gland
ERAL1	15.171	trachea
ERAL1	15.2279	pituitary_gland
ERAL1	18.7148	bladder
ERAL1	19.3718	bone_marrow
ERAL1	20.0448	skeletal_muscle
ERAL1	26.2135	uterus
ERAL1	43.3419	testis
ERAL1	46.394	small_intestine
ERAL1	6.58342	peripheral_blood_lymphocytes
ERAL1	71.3004	Monocytes
EXOS9	106.633	thymus

Supplementary file 8

RBPs	Tag frequency	Tissue
EXOS9	112.224	testis
EXOS9	14.0739	salivary_gland
EXOS9	150.945	Monocytes
EXOS9	18.7148	bladder
EXOS9	23.197	small_intestine
EXOS9	31.4853	kidney
EXOS9	3.45925	bone_marrow
EXOS9	40.9611	skeletal_muscle
EXOS9	41.695	peripheral_blood_lymphocytes
EXOS9	43.9484	adrenal_gland
EXOS9	62.3617	pituitary_gland
EXOS9	83.1344	uterus
EXOS9	96.336	trachea
FBL	110.677	testis
FBL	124.354	peripheral_blood_lymphocytes
FBL	157.281	uterus
FBL	18.4805	kidney
FBL	27.407	salivary_gland
FBL	37.7071	pituitary_gland
FBL	44.7524	Monocytes
FBL	5.22908	skeletal_muscle
FBL	56.1328	trachea
FBL	57.1329	adrenal_gland
FBL	58.3902	bladder
FBL	59.115	small_intestine
FBL	74.5685	thymus
FBL	96.859	bone_marrow
FMR1	10.0615	testis
FMR1	13.0727	skeletal_muscle
FMR1	14.9658	small_intestine
FMR1	23.514	Monocytes
FMR1	25.0322	trachea
FMR1	32.2288	adrenal_gland
FMR1	35.201	uterus
FMR1	3.70365	salivary_gland
FMR1	44.2333	pituitary_gland
FMR1	52.1979	thymus
FMR1	9.50939	peripheral_blood_lymphocytes
FUBP1	3.0342	trachea
FUBP1	3.74479	uterus
FUBP2	105.551	bladder
FUBP2	10.9871	adrenal_gland
FUBP2	127.992	bone_marrow
FUBP2	165.628	testis
FUBP2	203.282	Monocytes
FUBP2	26.9625	uterus
FUBP2	28.435	small_intestine
FUBP2	29.9911	peripheral_blood_lymphocytes

Supplementary file 8

RBPs	Tag frequency	Tissue
FUBP2	32.6177	trachea
FUBP2	33.989	skeletal_muscle
FUBP2	43.9954	thymus
FUBP2	52.0191	kidney
FUBP2	57.2857	pituitary_gland
FUS	121.15	kidney
FUS	177.75195	peripheral_blood_lymphocytes
FUS	183.3312	small_intestine
FUS	200.2572	trachea
FUS	219.00989	adrenal_gland
FUS	240.2977	bladder
FUS	358.34454	testis
FUS	369.81954	pituitary_gland
FUS	378.973	uterus
FUS	43.5757	skeletal_muscle
FUS	449.041	Monocytes
FUS	495.3648	bone_marrow
FUS	785.95153	thymus
FUS	91.10969	salivary_gland
FXR1	38.3466	skeletal_muscle
FXR1	8.34406	trachea
G3BP2	10.4582	skeletal_muscle
G3BP2	13.9313	testis
G3BP2	3.0342	trachea
G3BP2	3.74296	bladder
GRSF1	44.4472	skeletal_muscle
HNRPH3	124.3962	Monocytes
HNRPH3	130.7596	bone_marrow
HNRPH3	164.95508	kidney
HNRPH3	174.14154	testis
HNRPH3	188.1205	trachea
HNRPH3	206.6112	bladder
HNRPH3	217.0046	small_intestine
HNRPH3	247.9053	uterus
HNRPH3	256.22514	skeletal_muscle
HNRPH3	353.4543	thymus
HNRPH3	40.232	peripheral_blood_lymphocytes
HNRPH3	45.18446	salivary_gland
HNRPH3	82.7695	adrenal_gland
HNRPH3	84.1158	pituitary_gland
HNRPUL1	129.252	testis
HNRPUL1	14.8157	skeletal_muscle
HNRPUL1	21.9028	kidney
HNRPUL1	21.998	trachea
HNRPUL1	30.6799	small_intestine
HNRPUL1	39.1573	pituitary_gland
HNRPUL1	40.232	peripheral_blood_lymphocytes
HNRPUL1	44.2784	bone_marrow

Supplementary file 8

RBPs	Tag frequency	Tissue
HNRPUL1	44.6809	adrenal_gland
HNRPUL1	52.3376	Monocytes
HNRPUL1	54.6472	bladder
HNRPUL1	65.1594	uterus
HNRPUL1	82.0253	thymus
HNRPUL1	8.88875	salivary_gland
HNRPLL	14.2175	small_intestine
HNRPLL	15.3819	adrenal_gland
HNRPLL	16.469	bladder
HNRPLL	17.8964	thymus
HNRPLL	21.7198	uterus
HNRPLL	24.87075	peripheral_blood_lymphocytes
HNRPLL	29.5835	trachea
HNRPLL	38.6843	Monocytes
HNRPLL	54.1773	testis
HNRPLL	6.1006	skeletal_muscle
HNRPA0	117.52914	bladder
HNRPA0	119.781	kidney
HNRPA0	132.14355	bone_marrow
HNRPA0	174.49051	thymus
HNRPA0	218.27652	adrenal_gland
HNRPA0	221.69141	uterus
HNRPA0	292.02806	Monocytes
HNRPA0	35.55498	salivary_gland
HNRPA0	70.95465	peripheral_blood_lymphocytes
HNRPA0	75.09656	trachea
HNRPA0	76.622262	testis
HNRPA0	97.6095	skeletal_muscle
HNRPA0	99.34361	pituitary_gland
HNRPA0	99.52267	small_intestine
HNRPCL1	106.951	Monocytes
HNRPCL1	12.5924	salivary_gland
HNRPCL1	13.0727	skeletal_muscle
HNRPCL1	13.4747	bladder
HNRPCL1	13.6539	trachea
HNRPCL1	14.9137	thymus
HNRPCL1	15.2279	pituitary_gland
HNRPCL1	41.1927	uterus
HNRPCL1	45.4133	adrenal_gland
HNRPCL1	47.7377	bone_marrow
HNRPCL1	49.0099	peripheral_blood_lymphocytes
HNRPCL1	55.4414	kidney
HNRPCL1	56.4992	testis
HNRPCL1	6.73462	small_intestine
HNRPD	14.9658	small_intestine
HNRPD	22.4792	pituitary_gland
HNRPD	26.099	thymus
HNRPD	3.0342	trachea

Supplementary file 8

RBPs	Tag frequency	Tissue
HNRPD	3.66237	adrenal_gland
HNRPD	8.23854	uterus
HNRPDL	1028.3	thymus
HNRPDL	112.591	salivary_gland
HNRPDL	149.44	bone_marrow
HNRPDL	291.524	adrenal_gland
HNRPDL	299.11	kidney
HNRPDL	345.991	skeletal_muscle
HNRPDL	356.797	testis
HNRPDL	392.262	bladder
HNRPDL	465.728	Monocytes
HNRPDL	533.258	uterus
HNRPDL	534.28	small_intestine
HNRPDL	685.73	trachea
HNRPDL	685.978	pituitary_gland
HNRPDL	82.6585	peripheral_blood_lymphocytes
HNRPF	12.1362	Monocytes
HNRPF	131.1127	adrenal_gland
HNRPF	152.6347	kidney
HNRPF	160.8823	small_intestine
HNRPF	168.2316	pituitary_gland
HNRPF	174.4903	thymus
HNRPF	226.0745	bladder
HNRPF	231.3577	trachea
HNRPF	37.306	peripheral_blood_lymphocytes
HNRPF	67.1094	bone_marrow
HNRPF	76.3937	uterus
HNRPF	87.1514	skeletal_muscle
HNRPF	92.8755	testis
HNRPG	113.0374	bladder
HNRPG	127.6239	pituitary_gland
HNRPG	144.549	uterus
HNRPG	15.9125	bone_marrow
HNRPG	236.382	thymus
HNRPG	28.14772	salivary_gland
HNRPG	30.8008	kidney
HNRPG	37.1673	Monocytes
HNRPG	43.88951	peripheral_blood_lymphocytes
HNRPG	46.4378	testis
HNRPG	54.9355	adrenal_gland
HNRPG	80.0671	small_intestine
HNRPG	81.165	trachea
HNRPG	88.0229	skeletal_muscle
HNRPGT	115.32	testis
HNRPK	10.6192	Monocytes
HNRPK	15.3613	peripheral_blood_lymphocytes
HNRPK	22.4449	testis
HNRPK	23.5309	skeletal_muscle

Supplementary file 8

RBPs	Tag frequency	Tissue
HNRPK	24.6936	small_intestine
HNRPK	25.3798	pituitary_gland
HNRPK	38.8211	adrenal_gland
HNRPK	50.1802	uterus
HNRPK	5.5348	bone_marrow
HNRPL	15.7204	bladder
HNRPL	17.796	kidney
HNRPL	18.6421	thymus
HNRPL	20.0448	skeletal_muscle
HNRPL	20.5092	adrenal_gland
HNRPL	32.6311	pituitary_gland
HNRPL	37.1502	testis
HNRPL	49.4312	uterus
HNRPL	5.1851	salivary_gland
HNRPL	7.31491	peripheral_blood_lymphocytes
HNRPL	8.34406	trachea
HNRPL	85.3051	small_intestine
HNRPL	9.6859	bone_marrow
SYNCRIP	16.2532	testis
SYNCRIP	19.1733	skeletal_muscle
SYNCRIP	23.2064	bladder
SYNCRIP	24.6076	thymus
SYNCRIP	24.6936	small_intestine
SYNCRIP	25.6366	adrenal_gland
SYNCRIP	3.42231	kidney
SYNCRIP	52.3376	Monocytes
SYNCRIP	5.30986	trachea
SYNCRIP	6.52622	pituitary_gland
HNRPR	11.22888	bladder
HNRPR	11.8517	salivary_gland
HNRPR	13.14515	bone_marrow
HNRPR	15.17102	trachea
HNRPR	19.17325	skeletal_muscle
HNRPR	24.76679	testis
HNRPR	38.1628	small_intestine
HNRPR	4.55109	Monocytes
HNRPR	49.07568	adrenal_gland
HNRPR	58.4188	uterus
HNRPR	81.2797	thymus
HNRPR	95.7179	pituitary_gland
HNRPU	11.0696	bone_marrow
HNRPU	114.09011	thymus
HNRPU	18.96292	Monocytes
HNRPU	19.25892	salivary_gland
HNRPU	24.87067	peripheral_blood_lymphocytes
HNRPU	26.83	pituitary_gland
HNRPU	47.8906	small_intestine
HNRPU	47.9099	bladder

Supplementary file 8

RBPs	Tag frequency	Tissue
HNRPU	68.26955	trachea
HNRPU	74.14691	uterus
HNRPU	74.7122	adrenal_gland
HNRPU	82.135462	kidney
HNRPU	88.0229	skeletal_muscle
HNRPU	94.4234	testis
HTF9C	10.2546	adrenal_gland
HTF9C	12.7323	uterus
HTF9C	3.42231	kidney
HTF9C	3.72842	thymus
HTF9C	48.7596	testis
HTF9C	5.30986	trachea
HTATSF1	10.4582	skeletal_muscle
HTATSF1	14.2302	uterus
HTATSF1	5.12731	adrenal_gland
HTATSF1	5.24014	bladder
HTATSF1	5.96548	thymus
HTATSF1	6.96566	testis
HTATSF1	8.70163	pituitary_gland
IF2B	110.682	skeletal_muscle
IF2B	14.0739	salivary_gland
IF2B	18.2044	Monocytes
IF2B	18.7148	bladder
IF2B	27.7115	uterus
IF2B	42.4265	peripheral_blood_lymphocytes
IF2B	48.3432	adrenal_gland
IF2B	50.8838	small_intestine
IF2B	53.0986	trachea
IF2B	61.143	testis
IF2B	64.8746	thymus
IF2B	79.0398	pituitary_gland
EIF3S4	6.58342	peripheral_blood_lymphocytes
EIF4H	116.733	small_intestine
EIF4H	140.332	trachea
EIF4H	156.532	uterus
EIF4H	188.246	adrenal_gland
EIF4H	211.706	bone_marrow
EIF4H	28.1477	salivary_gland
EIF4H	283.685	Monocytes
EIF4H	41.794	testis
EIF4H	426.38	pituitary_gland
EIF4H	427.446	bladder
EIF4H	84.2624	thymus
EIF4H	88.2956	kidney
EIF4H	99.3526	skeletal_muscle
IGF2BP1	6.96566	testis
IGF2BP3	12.3834	testis
IGF2BP3	3.79275	trachea

Supplementary file 8

RBPs	Tag frequency	Tissue
IGF2BP3	4.84295	bone_marrow
IGF2BP3	6.52622	pituitary_gland
ILF3	101.768	small_intestine
ILF3	109.616	thymus
ILF3	15.3613	peripheral_blood_lymphocytes
ILF3	18.1284	pituitary_gland
ILF3	19.1649	kidney
ILF3	22.7555	Monocytes
ILF3	28.3659	bone_marrow
ILF3	3.48605	skeletal_muscle
ILF3	36.3762	testis
ILF3	42.2216	salivary_gland
ILF3	53.8571	trachea
ILF3	63.6303	bladder
ILF3	67.3875	adrenal_gland
ILF3	80.8875	uterus
IREB1	114.591	uterus
IREB1	125.015	bladder
IREB1	164.624	small_intestine
IREB1	238.054	adrenal_gland
IREB1	26.9822	bone_marrow
IREB1	30.3699	salivary_gland
IREB1	34.1332	Monocytes
IREB1	38.0299	thymus
IREB1	41.8327	skeletal_muscle
IREB1	476.386	kidney
IREB1	52.34	trachea
IREB1	55.8355	pituitary_gland
IREB1	78.9441	testis
IREB2	5.85978	adrenal_gland
KRR1	15.4792	testis
KRR1	17.9662	bladder
KRR1	21.70042	small_intestine
KRR1	33.3562	pituitary_gland
KRR1	33.37625	trachea
KRR1	40.4437	uterus
KRR1	45.4868	thymus
KRR1	7.61035	bone_marrow
KRR1	8.21355	kidney
KRR1	8.343665	Monocytes
KRR1	8.78968	adrenal_gland
LA	10.4854	uterus
LA	11.3777	Monocytes
LA	3.74145	small_intestine
LA	4.38895	peripheral_blood_lymphocytes
LA	4.49155	bladder
LA	5.85978	adrenal_gland
LA	6.1006	skeletal_muscle

Supplementary file 8

RBPs	Tag frequency	Tissue
LA	9.28754	testis
LARP4	15.4792	testis
LARP4	4.55109	Monocytes
LARP4	6.84462	kidney
LARP7	11.2244	small_intestine
LARP7	11.2289	bladder
LARP7	11.3777	Monocytes
LARP7	12.3834	testis
LARP7	12.6766	thymus
LARP7	15.171	trachea
LARP7	18.1284	pituitary_gland
LARP7	18.724	uterus
LARP7	24.9041	adrenal_gland
LARP7	33.989	skeletal_muscle
LRPPRC	10.6192	Monocytes
LRPPRC	13.8983	peripheral_blood_lymphocytes
LRPPRC	27.407	salivary_gland
LRPPRC	32.5169	bone_marrow
LRPPRC	34.2231	kidney
LRPPRC	38.9458	uterus
LRPPRC	42.0579	pituitary_gland
LRPPRC	49.6763	skeletal_muscle
LRPPRC	60.6841	trachea
LRPPRC	62.1081	small_intestine
LRPPRC	67.3733	bladder
LRPPRC	71.5858	thymus
LRPPRC	91.3275	testis
LRPPRC	91.5591	adrenal_gland
LSM5	15.3819	adrenal_gland
LSM5	17.80115	testis
LSM5	23.20636	bladder
LSM5	24.71562	uterus
LSM5	29.5821	Monocytes
LSM5	30.8008	kidney
LSM5	33.67306	small_intestine
LSM5	37.3599	bone_marrow
LSM5	45.35247	peripheral_blood_lymphocytes
LSM5	49.96088	thymus
LSM5	50.54775	skeletal_muscle
LSM5	62.9597	trachea
LSM5	63.7027	salivary_gland
LSM5	66.71254	pituitary_gland
MOV10L1	11.6094	testis
MATR3	12.1368	trachea
MATR3	20.9521	small_intestine
MATR3	4.35757	skeletal_muscle
MATR3	4.64377	testis
MATR3	5.92583	salivary_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
MATR3	5.96548	thymus
MATR3	5.99167	uterus
MATR3	9.7317	bladder
MGC10433	10.4396	thymus
MGC10433	16.6881	trachea
MGC10433	18.5751	testis
MGC10433	21.2417	adrenal_gland
MGC10433	32.9542	uterus
MGC10433	6.9185	bone_marrow
MGC10433	7.31491	peripheral_blood_lymphocytes
MGC10433	8.2312	small_intestine
MGC10433	9.10218	Monocytes
MGC27016	33.2804	testis
MINT	11.1109	salivary_gland
MINT	154.737	Monocytes
MINT	160.322	thymus
MINT	17.5558	peripheral_blood_lymphocytes
MINT	26.1454	skeletal_muscle
MINT	51.2731	adrenal_gland
MINT	68.4462	kidney
MINT	71.9785	testis
MINT	72.8209	trachea
MINT	81.6364	uterus
MINT	81.9403	pituitary_gland
MINT	89.7949	small_intestine
MINT	89.831	bladder
MINT	9.6859	bone_marrow
MKI67IP	3.86981	testis
MKI67IP	6.97211	skeletal_muscle
MKI67IP	8.20253	thymus
MSI2H	19.7768	adrenal_gland
MSI2H	27.7115	uterus
MSI2H	32.0644	thymus
MSI2H	6.52622	pituitary_gland
MSI2H	9.7317	bladder
MYEF2	100.61502	testis
MYEF2	10.9871	adrenal_gland
MYEF2	27.6979	bladder
MYEF2	31.4282	small_intestine
MYEF2	34.30145	thymus
MYEF2	50.03434	pituitary_gland
SLC4A1AP	20.2038	small_intestine
SLC4A1AP	6.74062	uterus
NCBP1	5.21979	thymus
NCBP1	7.73962	testis
NHP2L1	114.3052	kidney
NHP2L1	119.3096	thymus
NHP2L1	145.0294	adrenal_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
NHP2L1	161.5715	trachea
NHP2L1	168.4333	bladder
NHP2L1	172.5596	skeletal_muscle
NHP2L1	211.6257	Monocytes
NHP2L1	222.9909	small_intestine
NHP2L1	228.4173	pituitary_gland
NHP2L1	231.8823	peripheral_blood_lymphocytes
NHP2L1	254.6332	testis
NHP2L1	62.962	salivary_gland
NHP2L1	85.3812	uterus
NHP2L1	92.0161	bone_marrow
NKRF	17.226	uterus
NKRF	8.777901	peripheral_blood_lymphocytes
NOL3	10.3702	salivary_gland
NOL3	11.2289	bladder
NOL3	21.7878	skeletal_muscle
NOL3	25.4646	uterus
NOL3	3.74145	small_intestine
NOL3	6.1917	testis
NOL3	65.1901	adrenal_gland
NOL4	11.9775	bladder
NOL4	32.5064	testis
NOL4	57.2857	pituitary_gland
NOL4	8.20253	thymus
NOL8	3.79275	trachea
NOL8	5.21979	thymus
NOL8	6.06812	Monocytes
NOL8	8.70163	pituitary_gland
NOL8	8.7779	peripheral_blood_lymphocytes
NOL8	9.58665	skeletal_muscle
NONO	10.9514	kidney
NONO	17.226	uterus
NONO	20.1335	thymus
NONO	25.6366	adrenal_gland
NONO	28.0664	trachea
NONO	53.8546	Monocytes
NONO	6.73733	bladder
NONO	7.40729	salivary_gland
NONO	7.73962	testis
NONO	8.3022	bone_marrow
NONO	9.42677	pituitary_gland
NONO	9.58665	skeletal_muscle
NOP56	17.2962	bone_marrow
NOP56	17.44581	Monocytes
NOP56	22.6594	skeletal_muscle
NOP56	27.10152	adrenal_gland
NOP56	29.18337	small_intestine
NOP56	34.1348	trachea

Supplementary file 8

RBPs	Tag frequency	Tissue
NOP56	3.70365	salivary_gland
NOP56	52.2098	pituitary_gland
NOP56	55.4229	uterus
NOP56	58.5193	peripheral_blood_lymphocytes
NOP56	61.891885	thymus
NOVA1	13.4223	thymus
NOVA1	16.4624	small_intestine
NOVA1	21.6709	testis
NOVA1	27.5552	pituitary_gland
NOVA1	3.0342	trachea
NOVA1	6.73733	bladder
NCL	133.121	testis
NCL	13.837	bone_marrow
NCL	169.157	trachea
NCL	178.093	small_intestine
NCL	225.197	thymus
NCL	225.602	adrenal_gland
NCL	246.407	uterus
NCL	26.3337	peripheral_blood_lymphocytes
NCL	39.4428	Monocytes
NCL	54.9054	skeletal_muscle
NCL	55.5547	salivary_gland
NCL	66.7125	pituitary_gland
NCL	85.3395	bladder
NUFIP1	3.86981	testis
NUFIP1	5.23804	small_intestine
NUFIP1	5.2427	uterus
NUFIP1	6.97211	skeletal_muscle
HRB	13.0727	skeletal_muscle
HRB	152.462	Monocytes
HRB	176.463	testis
HRB	26.5493	trachea
HRB	29.2094	uterus
HRB	42.0579	pituitary_gland
HRB	44.8974	small_intestine
HRB	49.0099	peripheral_blood_lymphocytes
HRB	61.1462	thymus
HRB	62.2665	bone_marrow
HRB	66.6247	bladder
HRB	77.6421	adrenal_gland
HRB	8.14802	salivary_gland
HRB	91.7179	kidney
NXF1	109.25	small_intestine
NXF1	140.446	bone_marrow
NXF1	155.179	pituitary_gland
NXF1	190.984	uterus
NXF1	194.938	Monocytes
NXF1	207.3	thymus

Supplementary file 8

RBPs	Tag frequency	Tissue
NXF1	23.4077	peripheral_blood_lymphocytes
NXF1	24.4441	salivary_gland
NXF1	268.085	adrenal_gland
NXF1	31.3745	skeletal_muscle
NXF1	36.2765	kidney
NXF1	82.3451	bladder
NXF1	82.6821	trachea
NXF1	83.5879	testis
NXF2	10.6197	trachea
NXF2	11.6022	pituitary_gland
NXF2	33.2804	testis
NXF2	3.42231	kidney
PABP2	108.62	bone_marrow
PABP2	12.3273	pituitary_gland
PABP2	12.5924	salivary_gland
PABP2	13.8983	peripheral_blood_lymphocytes
PABP2	19.3878	thymus
PABP2	30.342	trachea
PABP2	3.74296	bladder
PABP2	47.1423	small_intestine
PABP2	62.9927	adrenal_gland
PABP2	69.6531	uterus
PABP2	70.4305	testis
PABP2	8.89801	kidney
PABP2	9.10218	Monocytes
PABP4	12.1368	trachea
PABP4	184.761	skeletal_muscle
PABP4	26.9625	uterus
PABP4	28.5282	peripheral_blood_lymphocytes
PABP4	3.79258	Monocytes
PABP4	44.8898	testis
PABP4	49.8082	adrenal_gland
PABP4	5.4757	kidney
PABP4	68.8427	small_intestine
PABP4	8.94822	thymus
PAIP1	116.094	testis
PAIP1	129.648	adrenal_gland
PAIP1	15.9288	Monocytes
PAIP1	161.775	uterus
PAIP1	25.9255	salivary_gland
PAIP1	29.0054	pituitary_gland
PAIP1	36.6662	small_intestine
PAIP1	426.17	skeletal_muscle
PAIP1	70.8401	thymus
PAIP1	8.04641	peripheral_blood_lymphocytes
PAIP1	88.7505	trachea
C20orf119	11.2244	small_intestine
C20orf119	11.3783	trachea

Supplementary file 8

RBPs	Tag frequency	Tissue
C20orf119	13.0524	pituitary_gland
C20orf119	41.0127	thymus
C20orf119	4.64377	testis
C20orf119	4.79124	kidney
C20orf119	5.24271	uterus
C20orf119	5.5348	bone_marrow
C20orf119	5.92583	salivary_gland
C20orf119	6.59226	adrenal_gland
PAPOLA	22.7555	Monocytes
PAPOLA	27.407	salivary_gland
PAPOLA	28.5282	peripheral_blood_lymphocytes
PAPOLA	36.681	bladder
PAPOLA	38.9458	uterus
PAPOLA	42.4834	adrenal_gland
PAPOLA	46.394	small_intestine
PAPOLA	52.7036	kidney
PAPOLA	53.0986	trachea
PAPOLA	62.749	skeletal_muscle
PAPOLA	76.8055	thymus
PAPOLA	80.4901	pituitary_gland
PAPOLA	99.8411	testis
PAPOLB	10.0615	testis
PAPOLB	5.96548	thymus
PAPOLG	10.4803	bladder
PAPOLG	13.0524	pituitary_gland
PAPOLG	18.7073	small_intestine
PAPOLG	26.8447	thymus
PAPOLG	8.34406	trachea
PCBP2	101.01	bone_marrow
PCBP2	101.096	skeletal_muscle
PCBP2	146.047	uterus
PCBP2	150.149	testis
PCBP2	172.176	bladder
PCBP2	176.734	Monocytes
PCBP2	186.324	small_intestine
PCBP2	209.537	thymus
PCBP2	47.0302	trachea
PCBP2	48.1474	salivary_gland
PCBP2	54.3852	pituitary_gland
PCBP2	60.2327	kidney
PCBP2	62.2602	adrenal_gland
PCBP2	88.5105	peripheral_blood_lymphocytes
POLDIP3	4.55131	trachea
POLDIP3	5.24271	uterus
SF3B14	106.325	skeletal_muscle
SF3B14	117.196	adrenal_gland
SF3B14	123.998	pituitary_gland
SF3B14	13.1452	bone_marrow

Supplementary file 8

RBPs	Tag frequency	Tissue
SF3B14	138.808	Monocytes
SF3B14	140.934	thymus
SF3B14	190.814	small_intestine
SF3B14	25.7907	trachea
SF3B14	58.5193	peripheral_blood_lymphocytes
SF3B14	67.7618	kidney
SF3B14	68.8826	testis
SF3B14	7.40729	salivary_gland
SF3B14	89.875	uterus
SF3B14	95.8198	bladder
PNPT1	15.4792	testis
PNPT1	20.0448	skeletal_muscle
PNPT1	21.9447	peripheral_blood_lymphocytes
PNPT1	7.25136	pituitary_gland
PNPT1	7.45685	thymus
PNPT1	9.52215	adrenal_gland
PPP1R8	13.1574	testis
PPP1R8	3.48605	skeletal_muscle
PPP1R8	4.35081	pituitary_gland
PPP1R8	8.9875	uterus
PPIE	10.2546	adrenal_gland
PPIE	3.09585	testis
PPIE	3.74145	small_intestine
PPIE	3.79275	trachea
PPIE	8.9875	uterus
PPIE	9.6939	thymus
PPIL4	10.877	pituitary_gland
PPIL4	10.9871	adrenal_gland
PPIL4	15.47923	testis
PPIL4	22.75544	Monocytes
PPIL4	3.48605	skeletal_muscle
PPIL4	5.24271	uterus
PPIL4	5.98874	bladder
PPIL4	6.16016	kidney
PPIL4	6.71116	thymus
PPRC1	10.4803	bladder
PPRC1	11.60943	testis
PPRC1	12.73229	uterus
PPRC1	16.60445	bone_marrow
PPRC1	23.861885	thymus
PPRC1	26.369	adrenal_gland
PPRC1	3.48605	skeletal_muscle
PPRC1	5.30986	trachea
PPRC1	6.06812	Monocytes
PPRC1	8.21355	kidney
PPRC1	8.2312	small_intestine
PPRC1	8.70163	pituitary_gland
PRKRA	15.4792	testis

Supplementary file 8

RBPs	Tag frequency	Tissue
PRKRA	17.4303	skeletal_muscle
PRKRA	4.49375	uterus
PRKRA	5.12731	adrenal_gland
PRKRA	5.80109	pituitary_gland
PSPC1	23.2189	testis
PSPC1	4.35757	skeletal_muscle
PSPC1	5.80109	pituitary_gland
PSPC1	5.85978	adrenal_gland
PSPC1	6.71116	thymus
PTBP1	108.406	adrenal_gland
PTBP1	108.502	small_intestine
PTBP1	116.053	Monocytes
PTBP1	119.96373	testis
PTBP1	120.3817	bone_marrow
PTBP1	134.223685	thymus
PTBP1	14.8157	skeletal_muscle
PTBP1	18.4805	kidney
PTBP1	211.95487	uterus
PTBP1	37.7071	pituitary_gland
PTBP1	50.0644	trachea
PTBP1	62.1331	bladder
PTBP1	80.7395	salivary_gland
PTBP1	82.6585	peripheral_blood_lymphocytes
PTBP2	11.8517	salivary_gland
PTBP2	121.331	uterus
PTBP2	131.229	trachea
PTBP2	14.8157	skeletal_muscle
PTBP2	168.724	testis
PTBP2	177.658	pituitary_gland
PTBP2	20.4799	Monocytes
PTBP2	246.822	thymus
PTBP2	34.9076	kidney
PTBP2	35.2844	bone_marrow
PTBP2	39.5535	adrenal_gland
PTBP2	90.5796	bladder
PTBP2	96.5295	small_intestine
RAD51AP1	13.1845	adrenal_gland
RAD51AP1	16.6044	bone_marrow
RAD51AP1	3.0342	trachea
RAD51AP1	41.156	small_intestine
RAD51AP1	57.4177	thymus
RAD51AP1	5.99167	uterus
RAE1	31.7324	testis
RAE1	3.48605	skeletal_muscle
RAE1	4.49375	uterus
RAE1	6.73462	small_intestine
RAVER2	15.7141	small_intestine
RAVER2	22.4792	pituitary_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
RAVER2	32.6177	trachea
RAVER2	3.48605	skeletal_muscle
RAVER2	5.12731	adrenal_gland
RAVER2	6.96566	testis
RBM12B	3.86981	testis
RBM12B	8.94822	thymus
RBM15B	16.4624	small_intestine
RBM15B	3.09585	testis
RBM15B	32.9171	peripheral_blood_lymphocytes
RBM15B	4.49375	uterus
RBM15B	5.30986	trachea
RBM15B	8.14802	salivary_gland
RBM15B	8.94822	thymus
RBM15B	9.10218	Monocytes
RBM35A	11.2344	uterus
RBM35A	11.9727	small_intestine
RBM35A	5.30986	trachea
RBM35B	11.6359	kidney
RBM35B	13.15736	testis
RBM35B	16.6881	trachea
RBM35B	4.489741	small_intestine
RBM35B	7.48592	bladder
RBM10	11.9775	bladder
RBM10	22.2219	salivary_gland
RBM10	22.4449	testis
RBM10	28.336	thymus
RBM10	37.7071	pituitary_gland
RBM10	39.4446	trachea
RBM10	54.9355	adrenal_gland
RBM10	55.4414	kidney
RBM10	64.353	small_intestine
RBM10	67.5079	Monocytes
RBM10	74.6121	peripheral_blood_lymphocytes
RBM10	89.9405	bone_marrow
RBM10	9.58665	skeletal_muscle
RBM10	99.6114	uterus
RBM12	102.22059	salivary_gland
RBM12	109.9899	trachea
RBM12	112.2885	bladder
RBM12	114.2658	adrenal_gland
RBM12	141.40149	pituitary_gland
RBM12	214.2021	uterus
RBM12	235.2287	bone_marrow
RBM12	266.209485	thymus
RBM12	314.02515	Monocytes
RBM12	36.2765	kidney
RBM12	67.1066	skeletal_muscle
RBM12	69.6566	testis

Supplementary file 8

RBPs	Tag frequency	Tissue
RBM12	79.001	peripheral_blood_lymphocytes
RBM12	96.52952	small_intestine
RBM14	11.931	thymus
RBM14	18.9638	trachea
RBM14	3.48605	skeletal_muscle
RBM14	45.6865	uterus
RBM14	4.64377	testis
RBM15	4.64377	testis
RBM16	6.1006	skeletal_muscle
RBM16	9.28754	testis
RBM18	6.1917	testis
RBM19	15.5553	salivary_gland
RBM22	10.4582	skeletal_muscle
RBM22	15.3819	adrenal_gland
RBM22	20.897	testis
RBM22	23.2717	kidney
RBM22	4.38895	peripheral_blood_lymphocytes
RBM22	5.96548	thymus
RBM22	8.34406	trachea
RBM23	10.6192	Monocytes
RBM23	15.5553	salivary_gland
RBM23	18.9638	trachea
RBM23	20.4818	peripheral_blood_lymphocytes
RBM23	22.6594	skeletal_muscle
RBM23	32.1895	bladder
RBM23	39.1573	pituitary_gland
RBM23	41.0127	thymus
RBM23	47.1844	uterus
RBM23	4.79124	kidney
RBM23	48.7596	testis
RBM23	56.8701	small_intestine
RBM23	62.9927	adrenal_gland
RBM23	8.99405	bone_marrow
RBM24	107.797	bladder
RBM24	11.6094	testis
RBM24	12.8954	trachea
RBM24	29.1833	small_intestine
RBM24	29.7495	bone_marrow
RBM24	30.7073	uterus
RBM24	31.1808	pituitary_gland
RBM24	848.854	skeletal_muscle
RBM25	12.1362	Monocytes
RBM25	13.0048	kidney
RBM25	21.9742	adrenal_gland
RBM25	22.8311	bone_marrow
RBM25	26.1454	skeletal_muscle
RBM25	33.3762	trachea
RBM25	41.02	testis

Supplementary file 8

RBPs	Tag frequency	Tissue
RBM25	41.1927	uterus
RBM25	41.3327	pituitary_gland
RBM25	41.7584	thymus
RBM25	8.88875	salivary_gland
RBM25	8.9831	bladder
RBM25	95.0329	small_intestine
RBM26	12.8954	trachea
RBM26	17.226	uterus
RBM26	18.5751	testis
RBM26	20.5092	adrenal_gland
RBM26	26.9385	small_intestine
RBM26	3.70365	salivary_gland
RBM26	41.0127	thymus
RBM26	4.49155	bladder
RBM26	71.0633	pituitary_gland
RBM27	12.73229	uterus
RBM27	13.42234	thymus
RBM27	14.5289	bone_marrow
RBM27	15.9288	Monocytes
RBM27	20.95211	small_intestine
RBM27	3.79275	trachea
RBM27	50.4729	peripheral_blood_lymphocytes
RBM27	51.8555	testis
RBM27	5.4757	kidney
RBM27	6.1006	skeletal_muscle
RBM3	129.628	salivary_gland
RBM3	131.113	adrenal_gland
RBM3	142.409	testis
RBM3	158.111	kidney
RBM3	169.862	small_intestine
RBM3	225.326	bladder
RBM3	232.614	peripheral_blood_lymphocytes
RBM33	15.3613	peripheral_blood_lymphocytes
RBM33	16.296	salivary_gland
RBM33	24.9041	adrenal_gland
RBM33	30.3406	Monocytes
RBM33	313.933	thymus
RBM33	33.6866	bladder
RBM3	338.298	Monocytes
RBM33	54.674	uterus
RBM33	55.7253	testis
RBM33	56.1328	trachea
RBM33	6.1006	skeletal_muscle
RBM3	366.131	thymus
RBM33	80.0671	small_intestine
RBM33	8.99405	bone_marrow
RBM33	93.5425	pituitary_gland
RBM34	12.1368	trachea

Supplementary file 8

RBPs	Tag frequency	Tissue
RBM34	18.6421	thymus
RBM34	20.2038	small_intestine
RBM34	27.0887	testis
RBM34	28.4604	uterus
RBM34	3.70365	salivary_gland
RBM34	3.79258	Monocytes
RBM34	39.1573	pituitary_gland
RBM34	4.79124	kidney
RBM34	6.1006	skeletal_muscle
RBM34	6.58342	peripheral_blood_lymphocytes
RBM34	9.52215	adrenal_gland
RBM3	49.6763	skeletal_muscle
RBM3	533.261	trachea
RBM3	68.4931	bone_marrow
RBM39	109.9847	Monocytes
RBM39	133.341514	skeletal_muscle
RBM39	139.0619	bone_marrow
RBM39	257.51637	bladder
RBM3	92.8708	uterus
RBM39	307.3237	kidney
RBM39	326.6118	testis
RBM39	391.14	adrenal_gland
RBM39	48.1474	salivary_gland
RBM39	512.579	small_intestine
RBM39	539.501	pituitary_gland
RBM39	650.982	thymus
RBM39	725.934	trachea
RBM39	793.894958	uterus
RBM39	90.705	peripheral_blood_lymphocytes
RBM4	129.376	bone_marrow
RBM41	3.09585	testis
RBM4	135.022	trachea
RBM41	3.79275	trachea
RBM4	143.053	kidney
RBM4	156.594	thymus
RBM4	160.199	bladder
RBM4	176.526	adrenal_gland
RBM4	182.734	pituitary_gland
RBM4	208.773	small_intestine
RBM4	216.935	Monocytes
RBM4	247.668	testis
RBM4	310.069	uterus
RBM4	65.8342	peripheral_blood_lymphocytes
RBM4	83.6653	skeletal_muscle
RBM4	85.1839	salivary_gland
RBM5	15.171	trachea
RBM5	19.4556	small_intestine
RBM5	64.5371	pituitary_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
RBM5	8.23854	uterus
RBM5	8.94822	thymus
RBM6	11.3297	skeletal_muscle
RBM6	11.3777	Monocytes
RBM6	18.3118	adrenal_gland
RBM6	19.1649	kidney
RBM6	32.5169	bone_marrow
RBM6	35.1116	peripheral_blood_lymphocytes
RBM6	3.70365	salivary_gland
RBM6	47.2117	testis
RBM6	74.338	trachea
RBM6	80.1385	uterus
RBM6	89.0466	small_intestine
RBM6	90.2279	thymus
RBM7	11.8517	salivary_gland
RBM7	22.4687	uterus
RBM7	29.5821	Monocytes
RBM7	31.8251	bone_marrow
RBM7	32.9171	peripheral_blood_lymphocytes
RBM7	36.2765	kidney
RBM7	45.4133	adrenal_gland
RBM7	47.1338	pituitary_gland
RBM7	55.3958	bladder
RBM7	57.2732	testis
RBM7	71.0876	small_intestine
RBM7	71.3038	trachea
RBM7	85.0081	thymus
RBM7	95.8665	skeletal_muscle
RBM8A	10.1519	pituitary_gland
RBM8A	18.64214	thymus
RBM8A	20.9163	skeletal_muscle
RBM8A	22.7555	Monocytes
RBM8A	3.034201	trachea
RBM8A	3.74296	bladder
RBM8A	4.10677	kidney
RBM8A	4.1511	bone_marrow
RBM8A	4.44438	salivary_gland
RBM8A	5.127313	adrenal_gland
RBM8A	5.99167	uterus
RBM8A	6.191692	testis
RBM8A	6.73462	small_intestine
RBMS2	18.5751	testis
RBMS2	3.0342	trachea
RBMS2	5.23804	small_intestine
RBMS3	13.0524	pituitary_gland
RBMS3	13.1845	adrenal_gland
RBMS3	14.9658	small_intestine
RBMS3	17.1508	thymus

Supplementary file 8

RBPs	Tag frequency	Tissue
RBMS3	18.2052	trachea
RBMS3	30.7073	uterus
RBMS3	3.09585	testis
RBMS3	37.4296	bladder
RBMS3	4.79124	kidney
RBMS3	8.99405	bone_marrow
RBMS3	9.58665	skeletal_muscle
RBP56	10.8355	testis
RBP56	10.877	pituitary_gland
RBP56	18.7073	small_intestine
RBP56	21.2417	adrenal_gland
RBP56	32.9381	bladder
RBP56	4.55131	trachea
RBP56	7.45685	thymus
RBP56	8.9875	uterus
RBP56	9.10218	Monocytes
RBP56	9.58665	skeletal_muscle
RBPMS	13.917	adrenal_gland
RBPMS	14.2232	bladder
RBPMS	18.7073	small_intestine
RBPMS	3.09585	testis
RBPMS	3.70365	salivary_gland
RBPMS	3.72842	thymus
RBPMS	41.9417	uterus
RIB1	10.4396	thymus
RIB1	113.772	testis
RIB1	155.904	pituitary_gland
RIB1	19.8494	kidney
RIB1	20.0448	skeletal_muscle
RIB1	27.674	bone_marrow
RIB1	31.4541	peripheral_blood_lymphocytes
RIB1	32.5921	salivary_gland
RIB1	51.5815	trachea
RIB1	7.58515	Monocytes
RIB1	86.1302	uterus
RIB1	8.9831	bladder
RIB1	91.2915	small_intestine
RIB1	95.2215	adrenal_gland
RKHD3	125.3818	testis
RPL12	113.037	bladder
RPL12	28.4604	uterus
RPL12	35.918	small_intestine
RPL12	40.9611	skeletal_muscle
RPL12	43.2354	Monocytes
RPL12	470.349	peripheral_blood_lymphocytes
RPL12	49.2152	thymus
RPL12	52.0056	adrenal_gland
RPL12	53.66	pituitary_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
RPL12	8.21355	kidney
RPL12	83.4406	trachea
RPL12	94.0916	bone_marrow
RPL13	1073.8	small_intestine
RPL13	1075.83	bone_marrow
RPL13	1123	thymus
RPL13	1477.4	adrenal_gland
RPL13	147.91	Monocytes
RPL13	157.953	bladder
RPL13	1890.37	uterus
RPL13	270.363	kidney
RPL13	3353.89	peripheral_blood_lymphocytes
RPL13	535.15	pituitary_gland
RPL13	578.685	skeletal_muscle
RPL13	628.839	trachea
RPL13	643.936	testis
RPL13	959.985	salivary_gland
RPL18	1146.09	bladder
RPL18	1415.02	small_intestine
RPL18	1427.53	Monocytes
RPL18	1529.68	bone_marrow
RPL18	1787.23	adrenal_gland
RPL18	1926.32	uterus
RPL18	3339.26	peripheral_blood_lymphocytes
RPL18	386.952	skeletal_muscle
RPL18	613.752	testis
RPL18	685.174	salivary_gland
RPL18	792.686	trachea
RPL18	808.526	pituitary_gland
RPL18	82.8199	kidney
RPL18	829.202	thymus
RPL22	1327.66	peripheral_blood_lymphocytes
RPL22	292.558	testis
RPL22	303.722	bone_marrow
RPL22	354.809	salivary_gland
RPL22	399.738	Monocytes
RPL22	453.401	adrenal_gland
RPL22	613.963	kidney
RPL22	662.914	thymus
RPL22	682.395	skeletal_muscle
RPL22	702.928	bladder
RPL22	707.007	pituitary_gland
RPL22	797.996	trachea
RPL22	904.683	small_intestine
RPL22	930.9552	uterus
RPL28	115.039514	skeletal_muscle
RPL28	131.9877	trachea
RPL28	1356.9213	peripheral_blood_lymphocytes

Supplementary file 8

RBPs	Tag frequency	Tissue
RPL28	137.2059	thymus
RPL28	143.359515	Monocytes
RPL28	21.02897	pituitary_gland
RPL28	21.2183	kidney
RPL28	267.35273	adrenal_gland
RPL28	470.67491	small_intestine
RPL28	560.2205	uterus
RPL28	571.46855	bone_marrow
RPL28	91.32826	bladder
RPL28	99.84113	testis
RPL28	99.99846	salivary_gland
RPL34	1051.84	salivary_gland
RPL34	1158.24	skeletal_muscle
RPL34	1300.85	Monocytes
RPL34	1311.75	bone_marrow
RPL34	1593.86	adrenal_gland
RPL34	1706.36	kidney
RPL34	1962.76316	small_intestine
RPL34	2173.67	thymus
RPL34	2195.62	bladder
RPL34	2349.44	pituitary_gland
RPL34	2352.48	uterus
RPL34	2473.6297	trachea
RPL34	4860.03	peripheral_blood_lymphocytes
RPL34	636.197	testis
RPL3L	27.8884	skeletal_muscle
RPLP2	113.846	pituitary_gland
RPLP2	131.986	thymus
RPLP2	143.701	salivary_gland
RPLP2	155.644	small_intestine
RPLP2	160.805	Monocytes
RPLP2	162.585	bone_marrow
RPLP2	220.579	testis
RPLP2	2272.01	peripheral_blood_lymphocytes
RPLP2	260.028	adrenal_gland
RPLP2	354.706	skeletal_muscle
RPLP2	584.187	uterus
RPLP2	65.8761	bladder
RPLP2	726.692	trachea
RPLP2	79.3976	kidney
MRPL3	38.4322	pituitary_gland
MRPL23	5.98874	bladder
RNASEL	4.84295	bone_marrow
RNASEL	6.74062	uterus
RNASE1	13.4692	small_intestine
RNASE1	6.74062	uterus
RNASE1	8.78968	adrenal_gland
RNH1	3.72842	thymus

Supplementary file 8

RBPs	Tag frequency	Tissue
RO60	3.62568	pituitary_gland
RO60	3.66237	adrenal_gland
RO60	4.64377	testis
HNRPA2B1	1064.2	testis
HNRPA2B1	1174.72	pituitary_gland
HNRPA2B1	194.071	salivary_gland
HNRPA2B1	2176.65	thymus
HNRPA2B1	314.541	peripheral_blood_lymphocytes
HNRPA2B1	337.276	skeletal_muscle
HNRPA2B1	411.362	kidney
HNRPA2B1	433.79	bone_marrow
HNRPA2B1	483.933	Monocytes
HNRPA2B1	664.001	bladder
HNRPA2B1	825.352	uterus
HNRPA2B1	930.973	adrenal_gland
HNRPA2B1	948.189	trachea
HNRPA2B1	981.009	small_intestine
HNRPA3	147.159	kidney
HNRPA3	183.560515	Monocytes
HNRPA3	195.794	bone_marrow
HNRPA3	209.5644	pituitary_gland
HNRPA3	211.9553	uterus
HNRPA3	222.99078	small_intestine
HNRPA3	235.15065	trachea
HNRPA3	243.2927	bladder
HNRPA3	262.9575	adrenal_gland
HNRPA3	338.54079	thymus
HNRPA3	520.8764	testis
HNRPA3	59.2508	peripheral_blood_lymphocytes
HNRPA3	74.81365	salivary_gland
HNRPA3	85.40837	skeletal_muscle
ROD1	3.0342	trachea
ROD1	32.5064	testis
ROD1	6.16016	kidney
ROD1	6.71116	thymus
RPL18A	1054.53	skeletal_muscle
RPL18A	1080.22	bladder
RPL18A	1166.65	salivary_gland
RPL18A	1492.32	bone_marrow
RPL18A	1617.3	adrenal_gland
RPL18A	1637.26	small_intestine
RPL18A	1773.98	thymus
RPL18A	226.796	Monocytes
RPL18A	2336.75	uterus
RPL18A	2386.86	peripheral_blood_lymphocytes
RPL18A	336.755	kidney
RPL18A	627.683	testis
RPL18A	646.821	pituitary_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
RPL18A	912.537	trachea
RPP29	10.2546	adrenal_gland
RPP29	10.3778	bone_marrow
RPP29	10.8355	testis
RPP29	12.2012	skeletal_muscle
RPP29	17.226	uterus
RPP29	17.4467	trachea
RPP29	21.9028	kidney
RPP29	26.3337	peripheral_blood_lymphocytes
RPP29	29.5821	Monocytes
RPP29	8.94822	thymus
RPP29	8.97949	small_intestine
RPS17	1126.14	pituitary_gland
RPS17	1629.02	adrenal_gland
RPS17	1670.09	kidney
RPS17	1673.59	bone_marrow
RPS17	1704.64	thymus
RPS17	1712.12	uterus
RPS17	1772.7	small_intestine
RPS17	2005.51	Monocytes
RPS17	2009.22	bladder
RPS17	2016.99	trachea
RPS17	5841.69	peripheral_blood_lymphocytes
RPS17	720.559	testis
RPS17	882.843	skeletal_muscle
RPS17	888.875	salivary_gland
RPS3	1126.63	bladder
RPS3	1180.52	pituitary_gland
RPS3	131.417	kidney
RPS3	1432.13	bone_marrow
RPS3	1679.16	uterus
RPS3	1828.82	small_intestine
RPS3	1918.65	thymus
RPS3	1950.89	peripheral_blood_lymphocytes
RPS3	290.214	skeletal_muscle
RPS3	562.06	Monocytes
RPS3	612.583	salivary_gland
RPS3	630.005	testis
RPS3	794.203	trachea
RPS3	889.955	adrenal_gland
RPS4Y1	100.124	Monocytes
RPS4Y1	161.705	pituitary_gland
RPS4Y1	168.433	bladder
RPS4Y1	182.386	adrenal_gland
RPS4Y1	194.948	trachea
RPS4Y1	296.783	thymus
RPS4Y1	6.73462	small_intestine
RPS4Y1	68.1471	salivary_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
RPS4Y1	68.8826	testis
RPS4Y1	83.6653	skeletal_muscle
RPS4Y1	86.316	peripheral_blood_lymphocytes
RPS4Y1	91.3242	bone_marrow
RPS5	11.2289	bladder
RPS5	127.45	adrenal_gland
RPS5	179.018	trachea
RPS5	181.086	small_intestine
RPS5	266.683	skeletal_muscle
RPS5	269.92	peripheral_blood_lymphocytes
RPS5	300.332	uterus
RPS5	35.7929	thymus
RPS5	5.80109	pituitary_gland
RPS5	61.143	testis
RPS5	73.3322	salivary_gland
RPS5	86.4813	bone_marrow
RPS5	9.10218	Monocytes
RPS9	1193.441	bone_marrow
RPS9	2441.719	peripheral_blood_lymphocytes
RPS9	281.4769	salivary_gland
RPS9	295.8556	pituitary_gland
RPS9	338.9954	testis
RPS9	385.2094	skeletal_muscle
RPS9	502.103	small_intestine
RPS9	516.084	kidney
RPS9	569.704	thymus
RPS9	632.87	uterus
RPS9	694.0419	Monocytes
RPS9	699.384	trachea
RPS9	776.29	bladder
RPS9	781.548	adrenal_gland
MRPS15	10.1519	pituitary_gland
MRPS15	10.9724	peripheral_blood_lymphocytes
MRPS15	12.4533	bone_marrow
MRPS15	162.102	skeletal_muscle
MRPS15	168.724	testis
MRPS15	18.2044	Monocytes
MRPS15	18.5182	salivary_gland
MRPS15	21.7092	bladder
MRPS15	24.7156	uterus
MRPS15	30.342	trachea
MRPS15	38.0299	thymus
MRPS15	40.4077	small_intestine
MRPS15	43.9484	adrenal_gland
MRPS15	8.21355	kidney
RTCD1	10.877	pituitary_gland
RTCD1	3.48605	skeletal_muscle
RTCD1	3.65746	peripheral_blood_lymphocytes

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RBPs	Tag frequency	Tissue
RTCD1	3.72842	thymus
RTCD1	4.39484	adrenal_gland
RTCD1	4.44438	salivary_gland
RTCD1	4.49375	uterus
RTCD1	9.72778	small_intestine
SNRPB2	104.396	thymus
SNRPB2	129.074	pituitary_gland
SNRPB2	22.6594	skeletal_muscle
SNRPB2	23.9562	kidney
SNRPB2	24.2736	trachea
SNRPB2	24.7668	testis
SNRPB2	25.031	Monocytes
SNRPB2	30.6799	small_intestine
SNRPB2	39.5535	adrenal_gland
SNRPB2	39.9994	salivary_gland
SNRPB2	41.1726	bladder
SNRPB2	43.4396	uterus
SNRPB2	60.7138	peripheral_blood_lymphocytes
SNRPB2	7.61035	bone_marrow
SAFB2	103.1581	Monocytes
SAFB2	119.1902	testis
SAFB2	152.1193	thymus
SAFB2	17.4467	trachea
SAFB2	20.7555	bone_marrow
SAFB2	23.5309	skeletal_muscle
SAFB2	31.4963	adrenal_gland
SAFB2	34.8143	salivary_gland
SAFB2	38.3299	kidney
SAFB2	5.85193	peripheral_blood_lymphocytes
SAFB2	66.5979	small_intestine
SAFB2	66.62472	bladder
SAFB2	71.1511	uterus
SAFB2	89.9168	pituitary_gland
SART3	4.84295	bone_marrow
SCAND1	10.2669	kidney
SCAND1	109.129	testis
SCAND1	14.8157	skeletal_muscle
SCAND1	16.296	salivary_gland
SCAND1	23.9295	pituitary_gland
SCAND1	24.2725	Monocytes
SCAND1	31.3188	thymus
SCAND1	32.9171	peripheral_blood_lymphocytes
SCAND1	4.48974	small_intestine
SCAND1	8.34406	trachea
SCAND1	95.2215	adrenal_gland
SCAND1	97.3646	uterus
SF1	101.809	bladder
SF1	133.895	testis

Supplementary file 8

RBPs	Tag frequency	Tissue
SF1	141.367	adrenal_gland
SF1	15.0582	kidney
SF1	170.61	small_intestine
SF1	189.26	pituitary_gland
SF1	214.012	thymus
SF1	235.922	uterus
SF1	38.5179	salivary_gland
SF1	40.9611	skeletal_muscle
SF1	43.9939	Monocytes
SF1	49.1213	bone_marrow
SF1	67.511	trachea
SF1	90.7049	peripheral_blood_lymphocytes
SFPQ	103.694	pituitary_gland
SFPQ	116.86862	testis
SFPQ	212.52042	thymus
SFPQ	23.4077	peripheral_blood_lymphocytes
SFPQ	25.27387	skeletal_muscle
SFPQ	47.18435	uterus
SFPQ	48.8881	salivary_gland
SFPQ	57.61837	small_intestine
SFPQ	63.655	kidney
SFPQ	68.2664	Monocytes
SFPQ	74.1106	bladder
SFPQ	75.0965	trachea
SFPQ	78.1791	bone_marrow
SFPQ	91.5591	adrenal_gland
SFRS11	108.8295	kidney
SFRS11	118.9782	small_intestine
SFRS11	148.6527	pituitary_gland
SFRS11	153.2444	testis
SFRS11	16.82433	peripheral_blood_lymphocytes
SFRS11	177.1139	bone_marrow
SFRS11	190.2355	uterus
SFRS11	198.377	bladder
SFRS11	20.74039	salivary_gland
SFRS11	284.8516	thymus
SFRS11	390.6353	Monocytes
SFRS11	48.8047	skeletal_muscle
SFRS11	56.4004	adrenal_gland
SFRS11	90.2676	trachea
SFRS12	12.1368	trachea
SFRS12	3.66237	adrenal_gland
SFRS12	4.48974	small_intestine
SFRS12	4.49375	uterus
SFRS12	4.79124	kidney
SFRS12	5.30961	Monocytes
SFRS12	5.41773	testis
SFRS12	6.71116	thymus

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RBPs	Tag frequency	Tissue
SFRS1	106.197	trachea
SFRS1	190.895	thymus
SFRS1	212.417	adrenal_gland
SFRS1	22.7555	Monocytes
SFRS1	286.595	small_intestine
SFRS1	29.2597	peripheral_blood_lymphocytes
SFRS1	295.09	uterus
SFRS1	488.016	pituitary_gland
SFRS1	62.2861	kidney
SFRS1	68.1471	salivary_gland
SFRS1	77.4872	bone_marrow
SFRS1	78.6022	bladder
SFRS1	85.4083	skeletal_muscle
SFRS1	89.0056	testis
SFRPS2	153.591	bone_marrow
SFRPS2	161.775	uterus
SFRPS2	166.873	Monocytes
SFRPS2	177.96	kidney
SFRPS2	236.058	testis
SFRPS2	256.225	skeletal_muscle
SFRPS2	296.581	pituitary_gland
SFRPS2	326.386	bladder
SFRPS2	468.43	small_intestine
SFRPS2	500.355	thymus
SFRPS2	61.4805	salivary_gland
SFRPS2	65.1027	peripheral_blood_lymphocytes
SFRPS2	80.572	adrenal_gland
SFRPS2B	14.8157	skeletal_muscle
SFRPS2B	15.5553	salivary_gland
SFRPS2B	16.0928	peripheral_blood_lymphocytes
SFRPS2B	27.1015	adrenal_gland
SFRPS2B	32.8101	thymus
SFRPS2B	32.9248	small_intestine
SFRPS2B	35.95	uterus
SFRPS2B	4.55109	Monocytes
SFRPS2B	56.1259	kidney
SFRPS2B	6.22665	bone_marrow
SFRPS2B	88.4666	pituitary_gland
SFRPS2B	90.2676	trachea
SFRPS2B	9.28754	testis
SFRPS2B	98.8142	bladder
SFRPS3	100.2143	peripheral_blood_lymphocytes
SFRPS3	100.6151	testis
SFRPS3	139.9308	small_intestine
SFRPS3	153.0869	adrenal_gland
SFRPS3	161.5642	Monocytes
SFRPS3	174.4669	trachea
SFRPS3	175.1706	bladder

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RBPs	Tag frequency	Tissue
SFRPS3	178.2187	thymus
SFRPS3	187.9885	uterus
SFRPS3	204.4885	pituitary_gland
SFRPS3	25.92552	salivary_gland
SFRPS3	78.4362	skeletal_muscle
SFRPS3	88.9801	kidney
SFRPS3	93.3998	bone_marrow
SFRS4	22.4487	small_intestine
SFRS4	3.45925	bone_marrow
SFRS4	4.49375	uterus
SFRPS5	12.8948	Monocytes
SFRPS5	18.9638	trachea
SFRPS5	22.4792	pituitary_gland
SFRPS5	24.7668	testis
SFRPS5	25.4419	small_intestine
SFRPS5	3.48605	skeletal_muscle
SFRPS5	55.9264	thymus
SFRPS5	7.31491	peripheral_blood_lymphocytes
SFRPS5	83.1344	uterus
SFRPS5	8.78968	adrenal_gland
SFRS6	13.4812	uterus
SFRS6	13.6533	Monocytes
SFRS6	18.7148	bladder
SFRS6	27.59037	thymus
SFRS6	28.435	small_intestine
SFRS6	29.6315	skeletal_muscle
SFRS6	30.9585	testis
SFRS6	34.42622	adrenal_gland
SFRS6	49.9657	kidney
SFRS6	58.4084	trachea
SFRS6	6.9185	bone_marrow
SFRS6	87.74139	pituitary_gland
SFRS7	165.317	peripheral_blood_lymphocytes
SFRS7	208.292	skeletal_muscle
SFRS7	215.857	bone_marrow
SFRS7	230.664	kidney
SFRS7	294.945	bladder
SFRS7	295.09	uterus
SFRS7	346.46	adrenal_gland
SFRS7	362.214	testis
SFRS7	368.369	pituitary_gland
SFRS7	43.703	salivary_gland
SFRS7	454.372	trachea
SFRS7	469.521	Monocytes
SFRS7	518.565	small_intestine
SFRS7	675.591	thymus
SFRPS9	194.729	uterus
SFRPS9	199.997	salivary_gland

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RBPs	Tag frequency	Tissue
SFRPS9	219.316	bone_marrow
SFRPS9	237.956	small_intestine
SFRPS9	24.4024	skeletal_muscle
SFRPS9	307.213	trachea
SFRPS9	309.421	peripheral_blood_lymphocytes
SFRPS9	314.168	kidney
SFRPS9	330.878	bladder
SFRPS9	396.704	thymus
SFRPS9	441.681	adrenal_gland
SFRPS9	443.058	pituitary_gland
SFRPS9	513.137	testis
SFRPS9	675.079	Monocytes
SIAHBP1	119.393	adrenal_gland
SIAHBP1	161.775	uterus
SIAHBP1	26.0096	kidney
SIAHBP1	27.8884	skeletal_muscle
SIAHBP1	31.8514	salivary_gland
SIAHBP1	50.8229	trachea
SIAHBP1	5.24014	bladder
SIAHBP1	56.3248	peripheral_blood_lymphocytes
SIAHBP1	58.9091	thymus
SIAHBP1	60.6115	small_intestine
SIAHBP1	68.1628	pituitary_gland
SIAHBP1	79.5628	bone_marrow
SIAHBP1	90.5536	testis
SIAHBP1	9.8607	Monocytes
SLBP	4.48974	small_intestine
SLBP	5.99167	uterus
SNRPA	12.2012	skeletal_muscle
SNRPA	12.4354	peripheral_blood_lymphocytes
SNRPA	20.1335	thymus
SNRPA	32.2052	uterus
SNRPA	6.73462	small_intestine
SNRPA	6.82696	trachea
SNRPA	7.40729	salivary_gland
SNRPA	7.58515	Monocytes
SNRPA	8.78968	adrenal_gland
SON	131.1091	salivary_gland
SON	136.8924	kidney
SON	154.2577	skeletal_muscle
SON	202.0036	testis
SON	246.9358	small_intestine
SON	274.595151	trachea
SON	286.71	bladder
SON	315.434	pituitary_gland
SON	321.55557	adrenal_gland
SON	349.7637	uterus
SON	362.5699	Monocytes

Supplementary file 8

RBPs	Tag frequency	Tissue
SON	709.89237	thymus
SON	71.2606	bone_marrow
SON	96.5569	peripheral_blood_lymphocytes
SPF45	110.9458	pituitary_gland
SPF45	124.6077	testis
SPF45	124.6268	skeletal_muscle
SPF45	135.02186	trachea
SPF45	138.49	bladder
SPF45	143.6715	small_intestine
SPF45	249.05905	thymus
SPF45	28.14776	salivary_gland
SPF45	49.0099	peripheral_blood_lymphocytes
SPF45	50.062	Monocytes
SPF45	59.33028	adrenal_gland
SPF45	77.8917	uterus
SPF45	80.94645	bone_marrow
SPF45	82.81995	kidney
SR140	127.43641	trachea
SR140	133.9978	bladder
SR140	134.2572	Monocytes
SR140	147.6456	thymus
SR140	19.01882	peripheral_blood_lymphocytes
SR140	30.67996	small_intestine
SR140	33.11751	skeletal_muscle
SR140	37.036429	salivary_gland
SR140	40.8192	bone_marrow
SR140	41.0677	kidney
SR140	43.2159	adrenal_gland
SR140	54.17738	testis
SR140	65.90832	uterus
SR140	73.23871	pituitary_gland
SRP14	145.505	testis
SRP14	158.434	bone_marrow
SRP14	201.016	trachea
SRP14	202.895	adrenal_gland
SRP14	230.08	skeletal_muscle
SRP14	233.467	small_intestine
SRP14	247.244	peripheral_blood_lymphocytes
SRP14	271.872	uterus
SRP14	338.364	bladder
SRP14	344.506	thymus
SRP14	465.434	kidney
SRP14	851.813	Monocytes
SRP14	91.1097	salivary_gland
SRP14	94.9928	pituitary_gland
SRP68	4.49375	uterus
SRP72	132.74	Monocytes
SRP72	23.5151	trachea

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RBPs	Tag frequency	Tissue
SRP72	28.8884	salivary_gland
SRP72	34.8605	skeletal_muscle
SRP72	46.394	small_intestine
SRP72	49.4312	uterus
SRP72	49.5336	testis
SRP72	50.0344	pituitary_gland
SRP72	54.8619	peripheral_blood_lymphocytes
SRP72	54.9355	adrenal_gland
SRP72	66.366	thymus
SRP72	83.022	bone_marrow
SRP72	88.9801	kidney
SRP72	99.5627	bladder
SRrp35	5.80109	pituitary_gland
STAU1	11.7039	peripheral_blood_lymphocytes
STAU1	133.4699	kidney
STAU1	136.244	bladder
STAU1	25.35329	thymus
STAU1	28.82356	Monocytes
STAU1	30.95848	testis
STAU1	3.70365	salivary_gland
STAU1	38.0518	bone_marrow
STAU1	41.941658	uterus
STAU1	42.783	pituitary_gland
STAU1	48.8048	skeletal_muscle
STAU1	57.6499	trachea
STAU1	68.0944	small_intestine
STAU1	75.4447	adrenal_gland
STAU2	101.096	skeletal_muscle
STAU2	11.3783	trachea
STAU2	13.4692	small_intestine
STAU2	15.953	pituitary_gland
STAU2	20.212	bladder
STAU2	20.5092	adrenal_gland
STAU2	4.44438	salivary_gland
STAU2	6.84462	kidney
STAU2	8.23854	uterus
STRBP	10.4854	uterus
STRBP	11.2289	bladder
STRBP	14.9658	small_intestine
STRBP	154.792	testis
STRBP	16.4271	kidney
STRBP	18.68	bone_marrow
STRBP	26.6663	salivary_gland
STRBP	34.1348	trachea
STRBP	42.0579	pituitary_gland
STRBP	42.504	thymus
STRBP	5.12044	peripheral_blood_lymphocytes
STRBP	6.59226	adrenal_gland

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RBPs	Tag frequency	Tissue
SUPV3L1	19.1733	skeletal_muscle
SUPV3L1	3.0342	trachea
SUPV3L1	3.62568	pituitary_gland
SUPV3L1	6.1917	testis
SUPV3L1	6.59226	adrenal_gland
SUPV3L1	8.2312	small_intestine
SUPV3L1	8.94822	thymus
SUPV3L1	9.73646	uterus
GARS	128.478	testis
GARS	14.0739	salivary_gland
GARS	202.313	pituitary_gland
GARS	28.7599	skeletal_muscle
GARS	29.0577	bone_marrow
GARS	40.424	bladder
GARS	41.695	peripheral_blood_lymphocytes
GARS	45.1745	kidney
GARS	58.5978	adrenal_gland
GARS	60.6115	small_intestine
GARS	66.366	thymus
GARS	67.5079	Monocytes
GARS	68.1552	uterus
GARS	85.7163	trachea
HARS	11.2344	uterus
HARS	4.38895	peripheral_blood_lymphocytes
HARS	5.98633	small_intestine
HARS	6.1006	skeletal_muscle
MARS	12.3203	kidney
MARS	140.055	uterus
MARS	160.322	thymus
MARS	19.1733	skeletal_muscle
MARS	211.292	testis
MARS	27.407	salivary_gland
MARS	39.8825	pituitary_gland
MARS	40.4077	small_intestine
MARS	44.1669	bladder
MARS	44.621	peripheral_blood_lymphocytes
MARS	63.7153	Monocytes
MARS	64.4576	adrenal_gland
MARS	65.2354	trachea
MARS	96.859	bone_marrow
SYNJ2	7.48958	uterus
SYWC	11.2244	small_intestine
SYWC	16.6881	trachea
SYWC	19.3491	testis
SYWC	3.62568	pituitary_gland
SYWC	4.39484	adrenal_gland
SYWC	4.49375	uterus
SYWC	5.85193	peripheral_blood_lymphocytes

Supplementary file 8

RBPs	Tag frequency	Tissue
SYWC	5.96548	thymus
TACC1	18.8535	pituitary_gland
TACC1	20.0448	skeletal_muscle
TACC1	250.778	bladder
TACC1	253.671	small_intestine
TACC1	28.3659	bone_marrow
TACC1	36.4087	Monocytes
TACC1	38.5179	salivary_gland
TACC1	41.695	peripheral_blood_lymphocytes
TACC1	422.412	uterus
TACC1	43.1211	kidney
TACC1	56.4004	adrenal_gland
TACC1	62.2012	trachea
TACC1	70.8401	thymus
TACC1	99.0671	testis
TARDBP	4.39484	adrenal_gland
TARDBP	7.97649	pituitary_gland
TARB1	10.2669	kidney
TARB1	12.1368	trachea
TARB1	13.1845	adrenal_gland
TARB1	14.9792	uterus
TARB1	17.8011	testis
TARB1	208.114	pituitary_gland
TARB1	21.4811	salivary_gland
TARB1	21.7878	skeletal_muscle
TARB1	44.1492	small_intestine
TARB1	54.435	thymus
TDRKH	37.92417	testis
Tenr	431.871	testis
TIA1	11.0696	bone_marrow
TIA1	15.2279	pituitary_gland
TIA1	17.4459	Monocytes
TIA1	17.80111	testis
TIA1	18.70726	small_intestine
TIA1	27.7115	uterus
TIA1	43.99542	thymus
TIA1	8.23451	bladder
TIA1	8.34406	trachea
TIA1	9.58247	kidney
TIA1	9.58665	skeletal_muscle
TIAR	13.94425	skeletal_muscle
TIAR	15.74262	kidney
TIAR	18.71483	bladder
TIAR	18.96286	Monocytes
TIAR	24.139191	peripheral_blood_lymphocytes
TIAR	29.58351	trachea
TIAR	33.69378	adrenal_gland
TIAR	3.70365	salivary_gland

Supplementary file 8

RBPs	Tag frequency	Tissue
TIAR	47.89064	small_intestine
TIAR	48.7596	testis
TIAR	56.6721	thymus
TIAR	60.9114	pituitary_gland
TIAR	63.6614	uterus
TNRC6C	11.931	thymus
TNRC6C	3.09585	testis
TRA2A	114.266	adrenal_gland
TRA2A	118.23	small_intestine
TRA2A	12.5924	salivary_gland
TRA2A	13.1452	bone_marrow
TRA2A	146.796	uterus
TRA2A	201.335	thymus
TRA2A	24.1392	peripheral_blood_lymphocytes
TRA2A	24.6406	kidney
TRA2A	25.7895	Monocytes
TRA2A	46.4127	bladder
TRA2A	48.8048	skeletal_muscle
TRA2A	68.1087	testis
TRA2A	78.3147	pituitary_gland
TRA2A	81.165	trachea
TRA2B	100.069	pituitary_gland
TRA2B	105.509	small_intestine
TRA2B	109.226	Monocytes
TRA2B	12.3203	kidney
TRA2B	177.473	thymus
TRA2B	22.9626	salivary_gland
TRA2B	23.5309	skeletal_muscle
TRA2B	55.6679	adrenal_gland
TRA2B	63.6303	bladder
TRA2B	73.5794	trachea
TRA2B	77.5381	peripheral_blood_lymphocytes
TRA2B	83.7139	bone_marrow
TRA2B	92.8708	uterus
TRA2B	92.8755	testis
TRBP2	15.3819	adrenal_gland
TRBP2	21.2384	Monocytes
TRBP2	22.4487	small_intestine
TRBP2	24.9066	bone_marrow
TRBP2	25.5407	testis
TRBP2	3.48605	skeletal_muscle
TRBP2	41.7584	thymus
TRBP2	4.79124	kidney
TRBP2	5.07595	pituitary_gland
TRBP2	61.4805	salivary_gland
TRBP2	9.10261	trachea
TRBP2	9.7317	bladder
TRBP2	9.73646	uterus

Supplementary file 8

RBPs	Tag frequency	Tissue
TRSPAP1	11.7614	bone_marrow
TRSPAP1	14.7053	testis
TRSPAP1	15.7141	small_intestine
TRSPAP1	20.212	bladder
TRSPAP1	21.2384	Monocytes
TRSPAP1	23.2043	pituitary_gland
TRSPAP1	39.5213	thymus
TRSPAP1	4.44438	salivary_gland
TRSPAP1	6.58342	peripheral_blood_lymphocytes
TRSPAP1	7.48958	uterus
TRSPAP1	9.10261	trachea
TUT1	10.6197	trachea
TUT1	11.2289	bladder
TUT1	121.512	testis
TUT1	14.4118	Monocytes
TUT1	20.2219	uterus
TUT1	22.2219	salivary_gland
TUT1	23.2043	pituitary_gland
TUT1	32.8101	thymus
TUT1	4.10677	kidney
TUT1	5.23804	small_intestine
TUT1	52.5806	bone_marrow
TUT1	7.32473	adrenal_gland
U2AF1	129.712	trachea
U2AF1	130.203	small_intestine
U2AF1	131.845	adrenal_gland
U2AF1	145.79	kidney
U2AF1	153.536	uterus
U2AF1	171.169	peripheral_blood_lymphocytes
U2AF1	205.114	bladder
U2AF1	229.093	testis
U2AF1	293.344	bone_marrow
U2AF1	300.511	thymus
U2AF1	329.212	pituitary_gland
U2AF1	363.329	Monocytes
U2AF1	36.6036	skeletal_muscle
U2AF1	77.0358	salivary_gland
HDLBP	3.65746	peripheral_blood_lymphocytes
YBOX1	101.47992	salivary_gland
YBOX1	114.8471	bone_marrow
YBOX1	1401.3963	skeletal_muscle
YBOX1	15.22787	pituitary_gland
YBOX1	161.6303	small_intestine
YBOX1	1705.039	testis
YBOX1	181.159	bladder
YBOX1	290.4017	peripheral_blood_lymphocytes
YBOX1	296.7824	thymus
YBOX1	299.628	trachea

Supplementary file 8

RBPs	Tag frequency	Tissue
YBOX1	387.6018	Monocytes
YBOX1	416.7773	adrenal_gland
YBOX1	42.4367	kidney
YBOX1	492.8142	uterus
YBOX2	10.877	pituitary_gland
YBOX2	3187.95	testis
YBOX2	43.4009	small_intestine
YBOX2	54.814	salivary_gland
YBOX2	6.73733	bladder
ZCRB1	17.7775	salivary_gland
ZCRB1	20.2219	uterus
ZCRB1	21.2384	Monocytes
ZCRB1	21.998	trachea
ZCRB1	25.4419	small_intestine
ZCRB1	40.267	thymus
ZCRB1	43.1211	kidney
ZCRB1	45.4133	adrenal_gland
ZCRB1	50.0344	pituitary_gland
ZCRB1	54.6472	bladder
ZCRB1	6.58342	peripheral_blood_lymphocytes
ZCRB1	81.266	testis
ZCRB1	88.0229	skeletal_muscle
ZNF638	10.9871	adrenal_gland
ZNF638	13.0727	skeletal_muscle
ZNF638	19.5787	pituitary_gland
ZNF638	24.7156	uterus
ZNF638	34.3015	thymus
ZNF638	4.44438	salivary_gland
ZNF638	6.96566	testis
ZNF638	7.58515	Monocytes
ZNF638	8.97949	small_intestine

Supplementary file 9

Supplementary data. List of RBPs that are expressed in only one tissue

IREB2

KHDRBS2

LARP5

LOC161931

Q8TC92

RED2

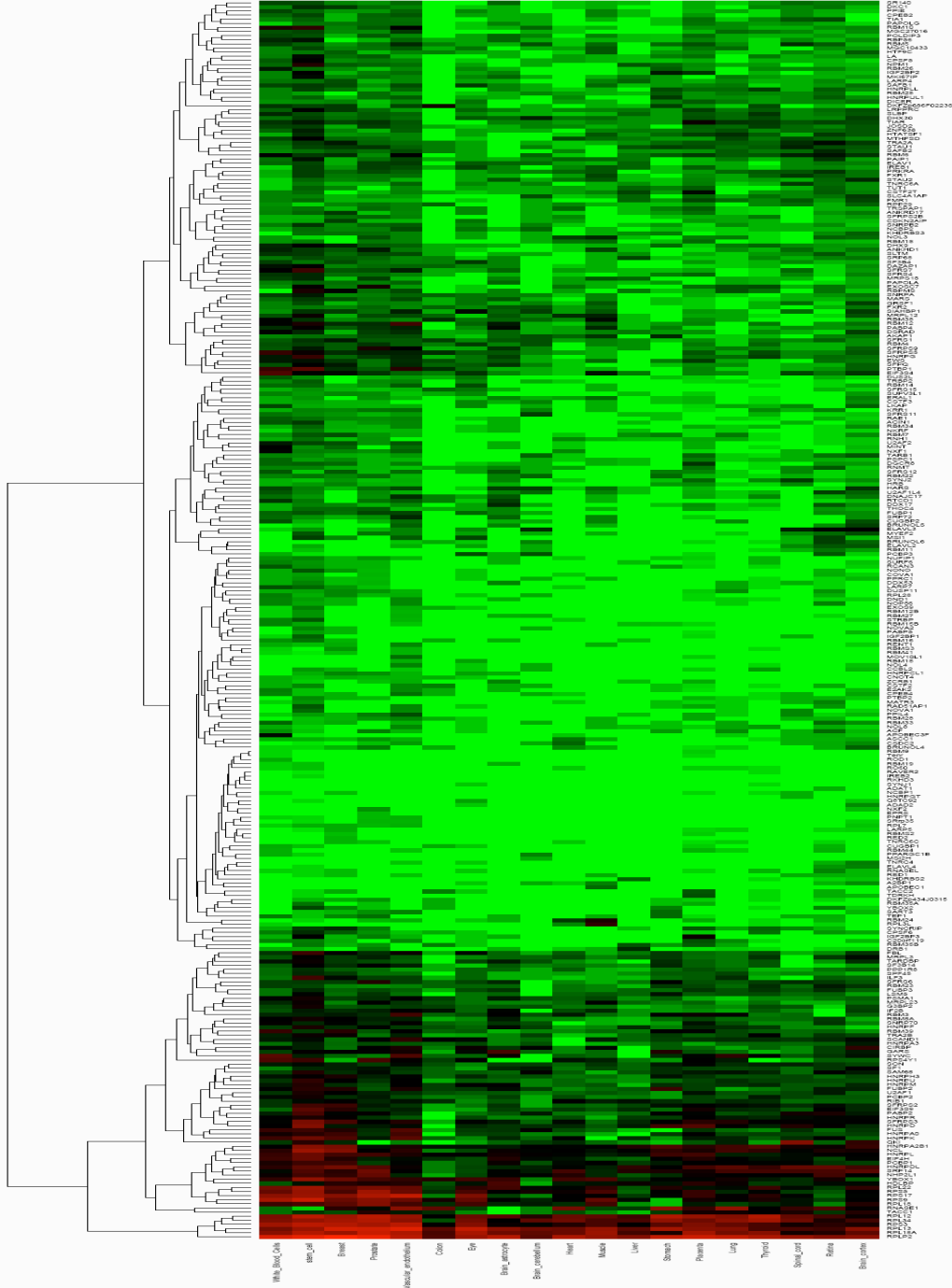
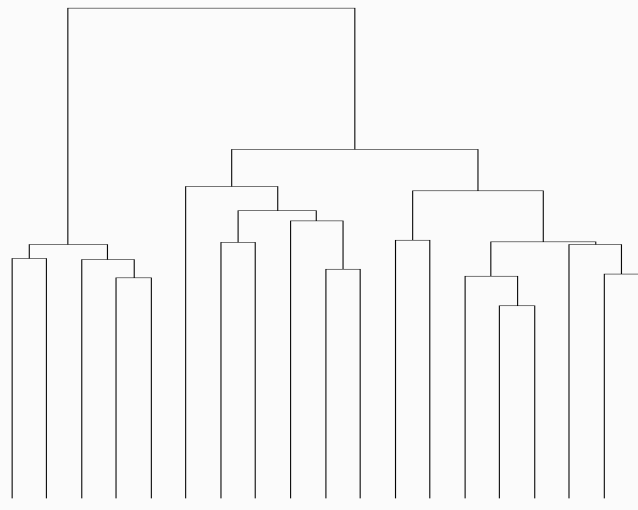
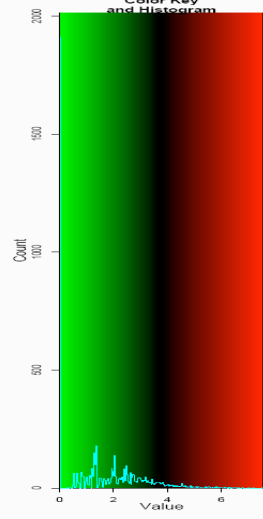
RKHD3

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Supplementary fig 10



Supplementary file 11

UNDER 10X

Tissue	Gene	Ratio [Normal / Tumor]
Brain_cortex	BRUNOL5	19.2151
Colon	RNASE1	38.13329921
Liver	IREB1	12.0649
Liver	MRPS15	12.0649
Lung	EXOSC7	15.7821
Lung	FMR1	11.2729
Lung	HTATSF1	24.8005
Lung	LSM5	15.7821
Lung	MGC27016	11.2729
Lung	PPP1R8	11.2729
Lung	RBPMS	15.7821
Lung	RPS4Y1	31.5642
Lung	ZNF638	15.7821
Thyroid	CPSF6	10.3504
Vascular_endothelium	BRUNOL5	12.8602
Vascular_endothelium	DKFZp686F02235	18.63321
Vascular_endothelium	IGF2BP2	10.4683
Vascular_endothelium	KRR1	10.4683
Vascular_endothelium	MOV10L1	10.4683
Vascular_endothelium	MSI1	13.9577
Vascular_endothelium	NCBP2	17.4471
Vascular_endothelium	NPM1	10.4683
Vascular_endothelium	PAPOLG	10.88753
Vascular_endothelium	RBP56	11.6185
Vascular_endothelium	RPS4Y1	50.3466
Vascular_endothelium	RTCD1	13.9577
Vascular_endothelium	SR140	11.23507
Vascular_endothelium	SRP72	27.1097
White_Blood_Cells	APOBEC3F	37.1575
White_Blood_Cells	ACIN1	15.488
White_Blood_Cells	RCAN3	14.863
White_Blood_Cells	CPEB2	12.14256
White_Blood_Cells	CUGBP2	15.7471
White_Blood_Cells	DHX30	10.1188
White_Blood_Cells	DHX9	25.7131
White_Blood_Cells	DSRAD	13.01117714
White_Blood_Cells	DUS2L	15.488
White_Blood_Cells	HNRPD	15.37399412
White_Blood_Cells	HNRPL	27.04828207
White_Blood_Cells	LARP7	16.1901
White_Blood_Cells	LKAP	18.007

Supplementary file 11

Tissue	Gene	Ratio [Normal / Tumor]
White_Blood_Cells	LRPPRC	13.41787
White_Blood_Cells	MINT	44.0796
White_Blood_Cells	NPM1	11.0199
White_Blood_Cells	PCBP2	12.15422998
White_Blood_Cells	POLDIP3	18.3665
White_Blood_Cells	SF3B14	14.0055
White_Blood_Cells	PPP1R8	11.1472
White_Blood_Cells	PPIE	18.5787
White_Blood_Cells	PSPC1	11.616
White_Blood_Cells	RAE1	14.1663
White_Blood_Cells	RBM10	10.40891787
White_Blood_Cells	RBM26	30.9759
White_Blood_Cells	RBM3	27.526
White_Blood_Cells	RBM4	16.1901
White_Blood_Cells	RBM6	14.73168908
White_Blood_Cells	RBP56	27.1039
White_Blood_Cells	MRPL12	15.7291
White_Blood_Cells	RPS4Y1	111.472
White_Blood_Cells	RTCD1	22.2945
White_Blood_Cells	SF3B4	20.2376
White_Blood_Cells	SFRS12	15.17547
White_Blood_Cells	SFRS7	45.7322
White_Blood_Cells	SLBP	25.7131
White_Blood_Cells	SNRPA	19.36
White_Blood_Cells	SRP14	11.53048732
White_Blood_Cells	SRP72	11.0199
White_Blood_Cells	TACC1	10.0039
White_Blood_Cells	TARDBP	11.37540367
White_Blood_Cells	ZNF638	11.8548

Supplementary file 12

GENES	White Blood Cells	Vascular endothelium	Thyroid	Lung	Prostate	Breast	Liver	Brain cortex	Brain astrocyte	Stomach	Colon	Brain cerebellum
FUBP2	-1.128216331	-1.208860052	-2.154699745	1.595363396	0.061282693	-0.530322388	0.81231804	0.907716332	0.304652522	-0.455177416	1.367162708	1.20686575
RPS17	0.943810252	-1.168746962	0.125883545	0.737910841	-0.398092752	1.67291522	1.576152867	1.725879847	2.588401811	1.282520736	3.208622381	3.275212766
RL3L	0	0	0	0	0	1.815677798	0	1.207947893	0	0	0	0
RB15B	-2.005726377	0	0	0	0	1.196482824	-1.104007707	2.47443535	1.76766143	2.047554769	1.279396553	1.940218267
SFRS6	-1.86131343	0.159273477	-0.273238439	0.978429127	0.210637108	1.74867926	1.192493347	1.348822639	0.392101607	0.215499382	1.506266115	4.154449834
MK67I	-2.054672301	-0.823773395	-1.643874121	-2.199261391	0.857237685	1.017910898	-2.202618891	1.991588174	0.586296421	1.635115297	2.666527961	1.401195607
DKFZp434.	0	0	0	0	0	-0.550132401	0.788296398	0	1.690792989	2.222739616	0	-0.201455943
SLBP	-3.24700059	-0.234875614	-1.538903958	0.203063761	0.176730348	1.155546408	-0.097828911	0.625665469	1.384657214	1.374812209	2.711318189	2.171032589
RAVR2	0	0	0	0	0	1.248238463	0	0	1.362196322	0	0	1.247072523
RM03	-0.940918417	0.534111423	-0.239623618	0.580358803	-0.211154104	0.676382218	0.694535373	0.337983396	1.257602703	0.087085491	4.09761255	1.07242198
FLJ40411	-1.312580542	0	0	0	0	1.25440733	0	0	0	1.125085429	1.185019316	0
HNRPU	0.519189095	-0.125978966	-0.391020248	-0.074569515	-0.187122299	0.417415869	1.140817507	1.229555738	0.892979086	0.995953344	2.550654457	2.157298253
SRP68	-2.237550449	-0.823773395	-0.360250147	0	-0.654431877	-0.064467227	-0.87408176	0.577217612	0.882568016	2.220189954	0.659837951	3.031671165
RBM18	-1.770326414	-2.047128813	0	0	0.014942316	2.996200164	2.415722689	1.502511079	2.172442238	1.952360728	2.601361396	0.927077203
SIAHBP1	0.717609734	-0.976210389	1.571308124	1.394690701	0.990661463	1.410502311	2.3924715	1.262534147	1.246255241	3.427670522	2.450219584	0.313048879
SUPV3L1	-2.046915767	2.358095483	0	1.526855984	-0.047489806	0.939816172	0.224527766	1.525320647	2.389789785	2.047554769	0.983881802	2.027892952
TUT1	0	-0.277933952	2.082132917	1.802193182	2.206760081	0.942450473	0.59531827	1.11569372	1.567759465	0.602336242	2.888837248	1.609568905
DGCR8	-1.34638518	-0.378365341	0	2.495342012	-1.21609602	0.44418865	-1.104007707	0.939091036	1.064881409	2.276367393	2.020682288	1.137955501
TACC2	1.105998278	2.217723041	0	0	0	0.513523864	0	1.860156163	1.387553568	1.153993383	-0.72652886	0
SFRS7	-3.822802645	-2.223709355	1.165840938	1.932321092	0.206906087	-0.108334533	0.883000513	2.066584656	0.435184918	2.837513497	2.069888935	3.724998969
RL18	-0.375994305	0.18742626	0.586679918	1.276908946	0.296706118	-0.134389785	1.894602085	0.743742687	1.65409318	4.311510282	2.329867075	1.575256707
Q8TC92	0	0	0	1.396729723	1.108146691	1.19394065	0	1.343446879	1.315426519	1.223648953	1.972543734	1.10388503
RBM9	0	0	0	0	0	0	1.328535473	0	0	0	1.279396553	1.212375403
KHDR3	0	0.027100398	1.165840938	1.396729723	-1.206564581	-0.564265367	-0.468617757	0.730612584	3.527477548	0.438640473	0	2.104588499
DJC17	-1.208396801	-1.125232206	-0.765715255	2.913149259	1.447580257	2.7335035	2.021682653	0.3316921	-0.73482434	1.354408879	0.911729869	2.230450864
DUS2L	-2.740065531	-1.637850427	0	0	-0.04749136	2.597237814	1.328535473	1.958499145	1.735719841	1.223648953	1.317192076	1.940218267
ACINU	-2.740065531	-1.948009373	0	0	-0.608082513	1.228025435	0.224527766	1.240186428	-1.357104966	0	2.378007915	1.196434462
PRGC1	0	1.119110752	0	0	0	0	0	0	0	0	0	0
LOC16193	0	0	0	0	0	0	0	-0.506637697	0	0	0	0
HNRPD	-2.732677389	0.310966907	0.313096335	-0.23708632	0.62800911	0.596340305	0.777639692	1.086225491	1.211083152	0.150958479	4.712678394	1.885563581
RS5	-0.925702509	0.026278187	0.480818218	1.625210063	-0.030383122	0.6542796	2.016288451	1.707008757	1.814127104	1.277612348	2.54020618	1.364604926
RBMS3	0	-1.254863618	0	0	1.16860466	-1.757271539	0	1.255581849	0	0	0	0
RBM15	0	-1.353982924	0	0	0	0	0	0	0	0	0	0
RBM41	0	-1.249738399	0	0	1.861753395	0	0	1.718068077	0	0	1.185019316	1.099628439
HNRPCL1	1.105998278	-1.254863618	0	2.220003164	0.435542532	-0.038298949	0	2.449054922	-1.357104966	1.451217964	2.571313678	2.198239618
IF39	-1.34638518	-0.130627847	-0.146671034	-1.207999861	0.386773952	0.136079105	1.329289273	1.4275994	1.257858362	1.597875107	2.524328168	2.192832382
SLTM	-2.011077795	-1.05289033	0.774730676	2.220003164	-1.007704686	1.163420829	0.18985499	1.506955311	1.617653385	2.627680718	1.250982869	1.45022233
RNH1	-1.386684285	0	0	-1.506115319	-0.723528762	2.309560707	0.189853332	1.948729029	1.355319557	2.252605672	1.59916533	1.334745771
SFR12	-2.719680308	-0.683401449	0	2.089875666	-0.030381581	0.157615399	0	1.95723506	-0.154249678	0.768072574	3.233557914	1.952219988
SFRS2	-1.366732916	0.616241952	0.384193626	-0.266939575	0.013947638	-0.299265854	0.477533614	1.031130577	0.73151017	0.361172357	1.909789146	3.718591552
PRKRA	-0.280686007	-0.812826221	0.150576723	-1.025675441	-0.245977729	1.044632336	1.293861039	1.156668799	1.162550876	2.763431296	2.528996898	1.620794939
RPL7	0	-1.353982924	0	0	0.030076388	0.52798596	1.328535473	1.350444361	0	1.354408879	1.612706565	2.70963021
RED1	0	0	0	0	0	1.831098765	0	0.423333088	1.542487953	0	0	1.766209156
RBM12	-0.479558328	-1.81751317	-0.583393699	3.211387054	-0.070978754	1.05684602	1.993304143	1.906349308	-0.13828098	1.408041451	4.449593807	1.670377835
MATR3	1.105998278	0	0	0	0	1.500763021	0	2.168432936	1.831966875	1.153993383	0	2.171674264
PAIP1	-0.515247966	-0.002830549	-0.072568075	0.243884216	0.196269677	1.954052269	-0.908757852	1.941067941	1.470311018	2.767258919	1.439157817	1.443188289
BRUNOL6	0	0	0	0	1.16860466	2.30193288	0	-0.403507083	1.728100561	0	-0.03338292	1.001584313
IGF2BP3	-1.312580542	-1.249738399	1.165840938	0	-2.12420217	-1.234759008	2.021682653	2.383943857	0.550629981	0	1.88581735	3.444295185
PCBP3	-1.312580542	0	0	0	0	0	0	1.265351965	0.698239817	1.223648953	0	-0.984154737
MINT	-3.78599709	0.579873703	0	1.396729723	1.199814169	0.727252162	0	0.801308601	1.09682688	0	2.794460286	1.792775619
IF2B2	-2.067266475	-2.348351643	-0.072568075	0	-2.12420217	0.92848674	2.863584339	1.179369805	0.417101344	0.318012304	3.226264462	3.336157401
RU2B	-2.091251528	-0.971081133	0.843723904	0.178296718	-0.66307084	0.308656616	0.917674946	3.259595414	2.718666976	0.844105581	2.292313521	5.23685231
TIA1	-2.046915767	-1.234367055	-0.36024807	2.604922746	-1.209560418	1.965075475	1.293861039	3.516424264	1.440137344	2.097131645	2.053890373	2.920547849
RBM34	-1.37411803	-0.823773395	0.332899111	1.526855984	0.676865329	1.148139896	0	0.955141694	0.915843262	2.223696636	1.972543734	2.202498424
CSDC2	0	-1.637850427	0	0	0	1.372372906	0	-0.493741634	2.067223454	1.818230986	0	0.710260529
PCBP1	-0.23637071	0.601852959	0.014445387	-2.17648342	0.04620427	-0.252943659	-0.36975924	1.148546222	-0.416790467	0.643433761	1.670347241	1.376363428

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GENES	White	Blood Cells	Vascular endothelium	Thyroid	Lung	Prostate	Breast	Liver	Brain cortex	Brain astrocyte	Stomach	Colon	Brain cerebellum
TIAR	-1.99833949		-0.378364395	0.15057852	-0.895549623	-1.240772866	0.679182841	0.18985499	0.421060051	1.426900833	1.757545359	2.761305935	1.511759509
PABP4	0.062199609		-0.600626577	0.332899111	2.443146716	-0.091411009	0.60491742	0.883000513	1.207821738	-0.45612277	-0.254505131	4.193957593	1.804702374
U2AF2	-1.63406237		-0.518739675	0	1.396729723	-0.631861284	0.437081467	0	2.221797942	0.185382987	2.144363973	2.308368338	1.499933266
SAFB1	0.668889716		0.051697864	0	0.583760219	0.435542532	0.932791187	0.917674946	0.702918359	0.651471417	0.743528189	2.985459691	1.69002308
NH2L1	-1.047327218		0.667642583	0.11667269	-1.131936277	-0.747572034	0.219613616	1.91092834	0.724070215	0.492973725	0.748613991	3.968085349	0.657249872
PNPT1	0		-1.353982924	0	0	0.030076388	1.210946432	2.021682653	1.363340424	1.310964489	1.223648953	1.185019316	1.499933266
HTSF1	-1.386684285		0.027100398	-0.765715146	-3.210863814	-0.107947678	1.528193204	-1.79715323	0.520492382	3.441638504	-0.341776666	1.360743193	2.996147187
NOL4	0		-2.043315535	0	0	0	0	1.198668763	0	0	0	0	-0.131552504
SRR35	0		-0.130627647	0	0	-1.21609602	0	0	1.267372865	0	0	0	1.10388503
RO60	0		0	0	1.526855984	1.199814169	1.19394065	0	1.901095074	1.355319557	1.223648953	0	1.544400067
EXOS9	0		-1.353982924	0	0	1.108146691	2.04035502	0	1.97847414	0	1.847140564	-1.394126113	2.198239618
RLA2	0.126358409		0.479478487	0.235155884	-0.651352388	-0.343280741	-0.524883495	2.066790222	1.626275589	0.569935048	0.486164981	2.250706074	2.055342646
RBM26	-3.433209483		0.41008611	-0.765715255	1.802193182	1.817831618	0.858852228	0.224527766	1.342724305	0.997797397	-1.258062439	1.878168026	3.004131898
SRP14	-2.444994599		0.438800733	-0.036198443	-1.684007909	-0.937273477	0.780405439	0.27686607	-0.040710221	1.409006575	0.303995292	0.707188233	1.496965027
PTBP2	0		1.119110752	0	1.396729723	0.66276725	0.913024896	0	2.651649147	2.15397487	1.818230986	2.528995303	2.805872375
FMR1	-2.005726377		-1.249738399	0.487050384	-2.422401615	-0.107949329	1.328852827	0.189857732	0.028447202	2.716547331	0.150959977	2.571313678	1.625387868
LKAP	-2.890760571		-1.353982924	-1.238409013	0.020740665	0.061285896	1.36031432	1.328535473	0.107342154	2.609944785	2.001407025	1.19267017	3.0057584129
PABP2	0.198883396		0.220922893	0.525266818	-0.826555295	-0.700855223	0.932513648	0.447670326	0.115419877	1.686851932	0.710573939	3.152500952	1.106758181
SYHC	-1.816017485		-1.745190778	1.165840938	0	0.240187749	0.638252362	-1.104007707	0.064592607	-0.611637941	1.746561595	3.152500952	-0.785564515
NONO	-0.263223675		0	0	0	0.232325122	-0.256508517	1.293861039	2.691798861	1.137021775	2.223696636	2.794460286	1.940218267
RBM8A	-0.815869545		-0.144046091	-0.072570901	0.947694772	0.024531849	0.860840195	1.83396303	2.325404164	0.076468692	1.046150583	3.129492745	1.941910775
HNRPD	-0.301835923		0.867552035	-0.404701092	-0.913567861	-0.887013858	0.16138832	1.060778576	0.386917252	1.783000091	0.557086803	1.184841175	2.180050992
A2BP1	0		0	0	0	0	0	0	-1.857082058	1.426502567	0	0	0.331767108
SFPQ	-1.649514222		-1.582338145	0	0.868041245	0.723223568	-0.380158904	2.427145113	1.497824913	-0.623752838	1.818230986	1.090181712	1.254742398
CSTF2	-1.301090441		-1.942884147	0	1.932321092	1.16860466	1.146662497	0	1.87495106	0.185382987	1.153993383	-0.72652886	0.387584129
RL34	-0.719618019		0.492062823	-0.35420443	-0.069410913	-0.042493144	0.026780641	0.940157976	1.296044058	2.144510494	0.4727776481	2.057776582	1.446826436
PAPOA	-2.091251528		0.458272655	-1.238409013	0.713887845	-0.262448775	0.705563225	-1.104007707	1.207976745	0.011635378	0.844105581	1.616828534	1.942768075
PP1R8	-2.411188345		-0.518739675	1.431510014	-2.422401615	-0.144669843	0.126332923	0.189853332	0.620530121	0.57928295	0.150954733	0.491691237	1.997022041
YBOX1	0.244378169		0.145374222	-0.07256768	0.10572699	-0.344179212	-0.231998693	1.002742493	3.002853395	0.254422375	1.255034329	0.972645278	3.36351141
EPRS	0		0	0	0	-1.138528273	0	0	1.265351965	0	0	0	1.10388503
TEP1	-2.079831466		1.664950194	0	1.396729723	-0.496879655	0.507555769	1.198668763	2.279630644	1.09682688	-0.542190871	1.360743193	2.433358937
RBM7	-0.686151115		0.458275009	1.858989677	-2.199261391	0.101602414	-0.55413936	-0.468617757	0.562147645	1.792369283	0.050380427	2.308368338	2.459358046
ELAV1	0.263052733		-0.541219606	0	0.682909062	0.136352996	2.543546164	-0.503292191	1.177195133	0.802718457	2.290824204	2.399513962	0.661492476
FBRL	-0.388903821		-0.423430385	-0.988857248	1.500053425	0.658683446	-0.0238113	1.576149255	1.374154559	0.939149277	0.844108469	2.821268734	2.864178333
RB35B	0		0	0	0.713887845	1.015383664	1.439314625	-1.384906831	0	0	0.448183387	0.372085993	0
NOL8	-1.37411803		-1.948009373	-1.643874121	-1.506115319	1.124682813	1.887087831	0.224527766	2.354190339	2.485514798	1.153993383	1.972543734	3.004131898
CIRBP	-1.816018522		-1.359524696	0.060966532	1.970053472	0.887530871	1.316426483	1.293861039	0.171589343	0.864267406	0.220612605	1.870335986	-0.324877616
NOP56	-2.046915767		-0.82889862	0	0	0	0.579719611	0	1.788146283	0	1.223648953	0	1.792775619
SART3	0		-2.047128813	0	2.220003164	-0.663070793	-0.040432722	0	0.794564268	1.661758978	-0.542190871	2.802111116	2.89564671
PSA1	-0.973833187		-0.928017403	0.150576723	3.878230413	0.836162093	0.664247367	0.629991883	1.763648811	-0.507765045	0.709837966	1.366878221	1.047021833
SYNJ2	-0.195092162		0.458275009	0.620580664	1.526855984	1.124682813	0.433921552	2.021682653	0.685782131	0.867446151	2.609942579	1.88581735	1.334745771
FUBP3	-0.085797598		0.20942332	-0.334933231	-0.826555295	-1.024235446	0.933451371	0.189857732	1.37151238	0.508069315	0.815193913	1.758374839	3.812021438
CPEB2	-2.496716637		-0.924201543	1.571306046	0	-0.24546838	2.946771916	0.189853332	3.765831699	1.661233844	2.540287744	1.61521718	2.08051102
ADAT1	0		1.119110752	0	0	0	1.248238463	0	0	0	0	0	0
PAPOG	1.105998278		-2.387618098	0.215115556	-1.911580427	-0.609962793	0.309879306	1.958855989	2.373005345	0.646752611	2.361794125	1.852155045	1.97698067
G3BP2	-0.705916683		-0.668912565	-0.177927205	0.737912264	1.07733847	0.409386267	0.917674946	0.225076504	0.323575818	0.562796342	-0.201452909	0.919813494
SR140	-1.823473459		-2.419040136	0.438258796	2.58394623	0.976150752	1.054655976	0.189853332	1.543680896	0.181931941	3.14816319	3.711568598	2.969798892
LARP7	-2.784399944		0	1.571308124	-1.506115319	1.124682813	-0.502864209	0	2.65367145	1.985167949	1.354408879	0	1.247072523
NUPL	-1.312580542		-0.130627647	0	1.396729723	-0.091413207	0.240307137	0	0.892552727	0.473065594	-0.542187203	1.279396553	2.403542931
MGC10433	0.137345194		-0.669862643	1.165840938	0.766085103	0.0612869	0.933454317	2.021682653	1.712093958	2.701723563	3.015407442	1.590302515	1.486195278
SFRS2B	-1.353768587		-0.382178619	0.263906833	0.426205773	0.284427846	0.357327331	2.021682653	0.55559652	2.303194907	1.354408879	1.360743193	1.220041763
TNRC4	0		0	0	0	0	0	0	-2.138841885	1.237684138	0	0	2.257719518
NOVA1	0		-1.948009373	0	0	1.16860466	1.25440733	0	2.651649147	2.414037018	0	0	1.544400067
RL13	2.214950206		-0.736000049	0.405466043	1.499566924	0.555271669	0.962215769	1.256781938	1.151707741	1.382865894	0.560266277	2.289291369	1.989343643
HNRPR	-1.259368146		0.93957527	-0.008027995	-1.232020403	0.32223738	0.907808649	0.323384513	0.643597603	2.484049124	0.507633873	3.677292585	2.517150947
SFRS1	0.681240494		-0.283059645	0.802899624	3.606296439	0.206858736	0.020007877	0.629991883	1.737679006	0.563525292	3.250638023	0.674014263	1.61765088
TACC1	-2.302975017		0.464155165	-1.288961412	-1.2914438	-1.295382764	0.634954661	-1.104007707	-0.392504106	4.16875606	1.162560946	0.204008154	1.281051683

Supplementary file 12

GENES	White Blood Cells	Vascular endothelium	Thyroid	Lung	Prostate	Breast	Liver	Brain cortex	Brain astrocyte	Stomach	Colon	Brain cerebellum
AKAP1	-2.005726377	-1.052043597	1.180197565	2.313020455	-0.091410243	1.258849438	1.328535473	0.512076983	0.292776412	0.581008747	1.177187565	0.418411792
ZCRB1	-1.353768587	-1.353982924	1.165840938	0	0	1.923655776	-1.104007707	1.752021956	0.012369989	1.916796134	0	1.915221166
RBM39	0.066186534	0.419530106	0.332893096	0.632947331	0.681779816	0.018521604	-0.180936237	2.401135059	1.69214343	0.262389232	1.672863721	1.8744467
PDIP3	-2.910528353	-1.942884147	1.431510014	0.426205773	-0.379097161	1.21960604	0.883000513	2.303836453	0.410554468	1.916796134	2.665692305	0.961753134
RBM14	-1.305190067	0.557395533	1.571308124	1.802193182	0.723223568	1.713151394	1.328535473	0.849782185	2.067223454	2.439708776	2.999001922	2.633362574
DDX53	-0.280686007	0	0	-1.506115319	-1.21609602	1.146662497	0	2.383943857	-0.694932589	0	0	0
RBM10	-2.342662927	-1.707499514	0.332899111	0.531567767	-0.395633265	2.011167847	1.293861039	2.530939338	1.384657214	2.763428142	3.950541549	0.008687164
RBM16	-1.774686214	2.217723041	1.165840938	1.526855984	1.124682813	0.985183651	-1.104007707	2.160981205	-0.041678447	1.746561595	-0.114729559	1.099628439
DND1	-2.005726377	-0.130626214	0	0	1.16860466	-0.05489633	0	1.890012285	0.666296507	1.451217964	1.59916533	1.905522584
ANKRD17	-0.957510762	-0.135752865	2.418605688	1.526855984	1.108146691	-0.012406708	1.813304541	0.796038452	2.955045687	1.418735963	1.973380781	1.650454495
STAU2	1.105998278	0.310969163	-0.478033183	-0.042207352	0.061620492	0.609670665	0.094661056	-0.34504368	0.1182707	2.306120835	1.810419287	2.032380431
SFR15	-1.353768587	0.557394315	0	0	1.199814169	0.969763752	0	1.890012285	-0.001785409	2.144363973	-0.03376221	1.905522584
TRA2B	-0.895431115	0.313450345	-0.254889008	-0.058580256	1.240131246	-0.104231063	-0.063153478	1.03354383	1.337091041	0.562419708	1.877032816	0.856801397
RBM33	-1.366734571	-1.353982924	0	0.020740665	1.124682813	-0.179624828	0	0.305944372	2.048466737	1.223648953	1.279396553	0.858921153
THOC4	-0.247770308	0.415211795	0	0	-0.83122621	-0.213855914	1.328535473	1.005017247	-0.041678447	2.223696636	2.082494091	3.402213532
FXR2	-2.067266475	-1.948009373	-1.238409013	2.220003164	0.030074786	0.66424993	0	1.230347817	-0.743802246	2.367511039	0.659766685	0.505371703
PPIL4	0	-1.740062693	0	0	-2.12420217	0.762156537	-1.104007707	1.494249081	1.985167949	1.832791042	1.985167949	1.212375403
BRUNOL4	0	0	0	0	-2.12420217	-1.251379154	1.293861039	-0.787258859	1.384715616	1.223648953	-1.394126113	1.942768075
RL22	-0.4611176379	-0.208943521	-0.22338995	1.032341299	-0.272777166	0.267725985	2.341616734	0.826454028	0.624356432	0.675569196	2.293742308	1.433724977
PCBP2	-2.497677255	1.003001536	0.502796984	0.531569246	-0.091411434	0.460161187	1.037153081	1.508772017	0.818260507	1.606907702	1.779236716	3.346574559
CNOT4	0.498056673	0.557394315	0	1.396729723	0.645657375	0.933453054	0	1.96629104	1.542487953	1.223648953	0.65976426	1.940218267
LSM5	-1.184410989	0.569359628	0.022741538	-2.758876386	0.399873001	2.13567698	0.749471561	2.060263872	0.664276163	0.231002316	3.265422331	3.448131857
PP1E	-2.922015763	-1.333487903	2.082132917	1.119350057	0.869879326	0.537824582	-1.104007707	2.956944667	1.558782914	2.047554769	3.545075479	2.433358937
SAFB2	-1.99423626	-1.74416555	-2.154699745	-0.042205166	0.207429342	0.694923757	-0.503292191	1.070160877	3.88627844	1.746046768	0.710253154	3.090962905
SF3B4	-3.00754226	-0.928018061	-1.238409013	1.802193182	0.723223568	0.291119491	1.98700822	1.889327262	0.02979921	2.511378167	0.222705647	1.970036748
LPPRC	-2.596587401	0.10346451	0.150576926	-2.199262499	-0.160402319	0.958631663	1.393828338	0.542140524	0.915990038	1.081574144	3.140739133	2.2906728361
BOLL	0	0	0	0	1.199814169	0	0	0	0	2.322261242	0	1.645642543
RL28	-2.079831466	-0.234872172	0	0	-0.030381581	-0.141731315	0	1.587267965	2.222739616	2.047554769	2.713826818	0.304795421
RBM27	0	-1.353982924	0	0	1.89296135	1.857687649	0	2.541391735	0.404126565	1.798114235	1.930120431	0.38673383
MSI2H	0	0	0	0	0	0	0	0	0	0	0	-0.274739997
PABP5	0	1.119110752	0	0	1.124682813	0.393945072	1.328535473	0.101417088	1.80574291	0	0	0
SPF45	-1.293704451	-1.211886855	0.438258796	-0.112788378	-0.047493288	0.543079006	0.224529423	1.437620832	1.56329474	0.304374662	2.851931185	1.584527235
RBM24	0	-1.353982924	0	0	0.754433077	2.16440405	0	1.762967256	1.794823038	0	1.185019316	2.790066345
PAP1L	-0.899728099	-0.135751444	1.165840938	-0.384722965	1.124682813	1.679448923	1.328535473	1.808840901	2.91588788	2.540286168	2.053890373	1.905522584
CCBL2	0	1.119110752	-1.643874121	0	0	-0.040432722	0	0.872657878	2.080051356	1.746561595	0	0.706003042
RNAS1	1.511075668	0.484932337	0.174293561	-1.000909539	0.075642607	0.570778551	-1.158099931	1.781563283	4.784537383	-0.843289824	-3.641087895	2.311678424
RM23	-1.447853476	-0.094498869	-0.072570901	2.372115095	-0.229813411	1.701719564	-0.503292191	1.742193939	0.87014756	1.102879653	1.383694213	1.430010561
LARP4	-2.285239525	0.176515412	0.620580664	-1.911580427	-0.091413207	0.166249384	-1.1964402	2.783367299	0.964234166	0.71512527	1.360743193	2.352125452
RB12B	0	-1.249738399	0	0	1.100813198	0	0	1.165186237	0	0	0	0
RBM23	-1.872659044	0.870582457	-2.018476568	-0.109381161	0.676865329	0.800960831	0.224527766	0.780006694	0.540623975	-0.571095864	0.08623178	3.550964181
RPP29	-1.803569456	0.02328712	-0.072566517	0	0.27383986	0.652711479	0.787809744	0.314793645	1.792369283	1.354408879	-0.474563259	-0.131552504
CCP3	-2.698874903	0	1.165840938	0	-0.016281851	0.647792724	1.98700822	1.668936253	1.426502567	-1.696180587	1.185019316	1.792775619
IF2B	-0.555803251	-2.02663078	-0.206098245	2.323325758	0.968348445	0.239848181	0.917673955	1.207819656	-0.556966101	1.925364441	1.153090013	3.183327401
ELAV2	0	-1.353982924	0	0	0	0	0	-0.880771174	2.490449593	0	0	3.735306654
JOS2	-1.79618127	-0.570739195	-1.371851089	0.227086835	-0.704283069	2.394865482	0.883004912	0.636564293	1.403354241	1.354408879	2.456332079	0.872415368
TNR6C	0	0	0	0	0	-1.251379154	0	1.207947893	0	0	0	0
HNRPGT	0	0	0	0	0	0	0	1.18180388	1.730805627	0	0	1.173555864
CSTF3	-1.37411803	0	0	0	1.561946553	-0.023812576	0	0.520596803	0.473065594	2.144363973	1.616828534	0.414151852
HNRH3	-1.045873982	0.496807858	-0.988858408	0.288133748	-0.591239854	-0.555505953	0.224527766	0.939771177	1.07175965	-0.067068729	1.089248492	1.632841648
NXF2	0	0	0	0	-1.831675453	0	0	-1.163934877	0	0	0	0
HNRPM	-0.568365395	-0.536092955	-0.701175176	-0.109385596	0.645658785	0.093100233	1.323137406	0.726076627	-0.024260641	0.409730842	1.109048516	1.740250491
RBM4	-2.784399944	-0.565615394	-0.22671606	1.932321092	-0.091411009	0.094297876	-0.468617757	1.443148844	0.762085996	2.734523341	0.626554765	0.996882003
PSPC1	-2.452383458	-1.254863618	-1.238409013	0	-0.016281851	0.71279265	0.224527766	0.814539263	1.831966875	1.746561595	1.19267017	1.037601203
ANKHD1	-1.568106647	0.015337791	1.431510707	0.531567767	-0.036829655	0.709196246	0.836536643	1.258807615	3.353026925	3.092440113	1.586485924	3.336777254
ASCC1	-1.301090441	-1.353982924	0	0	-0.518922892	1.109647839	1.98700822	0.520596336	2.996342088	1.223648953	1.19267017	1.766209156
NADAP	-2.046915767	0.174407506	-1.171184228	-1.506115319	-0.030379931	1.34585866	-0.063155298	-0.410682078	2.483330408	2.252605672	1.88581735	0.484233326
ELAV3	0	-2.043315535	0	0	-1.21609602	1.25440733	0	-0.460336871	3.18006015	0	0	1.29178729

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GENES	White	Blood	Cells	Vascular	endothelium	Thyroid	Lung	Prostate	Breast	Liver	Brain	cortex	Brain	astrocyte	Stomach	Colon	Brain	cerebellum
RBM3	-3.315131013			-2.040170617		1.536871084	-0.826555476	0.455126665	0.534933439	0.59531827	1.935713362	-0.400605541	0.402272657		0.889511325		0.784742007	
ROD1	-1.369221953			0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
RB35A	0			0	0	0	0	1.124682813	2.236562805	1.98700822	0	0	0	1.354408879	0.205039217	0.253990204		
HNRPF	-1.985435008			-0.260584197		-0.360248489	-0.89784336	-0.301525021	0.316243039	-1.854912118	4.108866934	0.223546166	1.617297289	0.4971692754	1.041692754	0.253990204		
CPSF6	-0.98525325			1.119110752		-2.337025166	1.932321092	0.375084627	-0.125107873	1.328535473	1.711433368	2.080051356	1.354408879	1.177187565	2.856510435			
PAPOB	0			0	0	0	0	0	0	1.328535473	0	0	0	0	0	0	0	0
SF01	-0.280684352			0.375079471		-0.524552635	0.243885325	-0.482366635	0.387048723	-0.063155298	-0.030529203	0.200657992	-0.283413633	1.065233218	0.425384738			
EWS	-0.268119946			-0.012672759		2.082132917	0.989228342	-1.037326392	0.495761316	1.293861039	1.631634119	-0.138858215	0.122050941	1.035051542	1.66020505			
NOVA2	0			1.119110752		0	0	0	-1.251379154	0	1.99865439	1.308259844	0	0	-1.378625027			
IF34	-0.73393241			0.310961936		0	1.883044698	0.896789307	0.864919894	0.512208295	0.871348163	1.09682688	3.070992796	1.805798484	2.009544527			
CPSF5	-1.544869038			-2.047130104		0.332899111	0	0.363452772	0.967817894	0.189853332	3.480562937	2.422477902	2.5369455	1.805798484	2.009544527			
MTHFS	-1.816017485			0.716431778		0.620579106	-1.090215389	-1.206564581	0.572003148	-0.180936237	2.106431178	3.464156531	1.249573317	2.531503935	3.291817689			
SON	-1.646149826			-0.522824571		0.038656852	-0.137517935	0.352250269	0.327658943	0.59531827	1.553867406	0.984970674	0.106491846	0.802179837	1.130137998			
RTC1	-3.104340011			-2.636031327		0	1.396729723	-0.030380481	1.906227386	0	1.24728038	-1.242271264	1.818230986	0.372085993	1.537303683			
YBOX2	0			0	0	0	0	1.16860466	0.510978411	0	1.27314526	0	-0.747236816	0.897156345	1.334745771			
TRBP2	-0.888237982			-2.047128813		0	2.779620193	0.357974784	2.445819261	0	2.616103027	1.71191327	2.252605672	2.865469818	1.512762881			
BRUNOL5	0			-2.554137271		0	0	1.124682813	0	0	-2.955696428	-0.119420828	0	0	0	0	0	0
RBM11	0			0	0	0	0	1.809596113	1.206024698	0	0.28722343	2.06438374	0	0	0.888046837			
RIB1	0.144083473			-0.050515884		-0.313730088	0.204021805	0.416017177	0.891531589	0.224527525	1.358692566	0.710104591	0.940884323	0.850307433	2.587232541			
NKRF	0			-0.130627647		1.858989677	0	-1.138528273	0.999213141	0.836536643	2.485719653	3.388559953	1.223648953	2.447367455	2.088885556			
RS3	-0.474060256			-0.497048859		0.354175422	1.050785832	0.063693053	-0.14880804	2.139346547	2.534841241	0.929510798	0.666123095	2.014462634	1.853654103			
RKHD3	0			0	0	0	0	1.199814169	1.825938649	0	0	0	0	0	0	0	0	0
U2AF1	-0.940918969			0.07874186		0.24588364	0.024145223	-0.281693999	-0.309229952	1.000786425	2.130765573	0.611256151	1.655038425	0.696199729	1.660869378			
RPL18A	0.229655305			-0.101096069		-0.421446238	0.818879282	0.152209885	1.181119717	1.543359718	1.751069436	1.16236549	1.703836978	2.038125961	1.800836829			
VIGLN	0.816510554			-0.442803451		0.661403313	-0.070230591	0.68178234	0.928532009	-0.043736637	1.139786218	-0.913228015	0.499265875	1.662696556	3.000644059			
NPM1	-2.399702729			-2.348351643		0.620580145	0.713887845	1.282837755	0.770502341	-0.033284162	1.645648525	2.200841362	0.556422469	3.582915979	1.804702374			
LA	-2.079831466			1.22725611		0.397437113	1.526855984	0.297516117	0.369095698	0.224527766	1.610633135	1.729285776	0.969347101	3.272113228	2.89564671			
NXF1	-1.484657708			0.510883164		0	0	-0.631861284	1.023319529	0	1.890012285	1.426502567	1.223648953	1.61521718	2.202498424			
ERAL	-1.398104348			-1.353982924		0	0	0.662768901	2.292551929	1.328535473	0.56559513	-0.001785409	2.609944785	3.272113228	0.414150593			
SFRS9	0.445373462			-0.522552295		0.332898072	0.580358803	0.192763291	2.109814733	0.917673955	1.115044326	-0.228465587	2.763428142	2.283166472	0.996877159			
DUS	-0.253100111			-0.717671883		-0.36024807	1.669403057	0.147857822	0.470348077	1.288463793	1.052614685	0.153704412	2.453020308	2.688916489	-0.274736107			
KFC1	-0.940917489			-0.382178619		0.332898072	1.932321092	0.550989543	-0.429280664	3.239771972	2.026664922	1.344618598	2.223696636	3.487604409	2.963152262			
MYEF2	0			-0.378363449		1.165840938	0	-0.401917258	2.793458182	0	0.480822137	3.241540236	0	1.279396553	2.266640518			
RN5A	0			0		-1.238409013	0	0	1.663137908	1.198668763	1.115168669	1.913158183	2.037040176	0	2.334251382			
ROA3	-0.510697234			0.027097875		-0.919863858	0.581040287	-0.969434615	0.871633209	0.207340831	4.08946304	1.228660834	0.270462289	1.052557686	1.864522142			
NCBP2	-0.280686007			-2.859173446		0.620580664	2.779620193	0.656347662	1.223864789	1.293861039	1.755798922	1.639237122	1.856682292	2.571313678	0.568300573			
E2AK2	-2.091251528			-1.942884147		0	0	1.904093613	0	0	0.121392082	0.710118488	2.453020308	-0.03338292	0			
RBP56	-3.299677629			-2.452598655		1.571308124	0.426205773	0.601735597	2.470985194	0.224527766	0.79456709	0.461976941	2.740704531	0.624044895	1.330437995			
RBM25	-2.046915767			-1.637850427		0.620580664	-2.199261391	-1.19002846	1.793962042	0.224527766	2.606098683	1.225910973	-0.765330755	1.360743193	1.220046741			
RS9	-0.978570347			-0.010407312		-0.101553849	1.427726151	0.657843602	0.719080214	0.538160625	1.717577061	1.206119189	4.686611915	2.533499536	2.760199549			
SFR11	-2.290409268			0.579872614		-1.238409013	0	-0.016281851	0.082539727	0.094661056	2.551482157	1.831966875	0	1.88581735	-0.615755907			
CPEB4	0.445380181			-0.541219874		0	1.526855984	1.16860466	1.502147791	0	0.818102782	2.235635133	1.818230986	0.218580452	-0.043879256			
SRP72	-2.399702729			-3.299891597		0	0	0.88590915	0.696175356	0	0.305944372	-1.100398556	1.818230986	2.802111116	2.490184365			
HNRPA0	0.631662915			-0.439720971		-0.391021027	0.395226978	0.229907324	0.362643259	3.377703892	1.451524238	0.946151387	0.766823913	3.516390399	2.571402953			
TRA2A	0.486572711			1.068218038		-0.632180648	-0.672403559	0.797084599	0.182139277	0.629991883	0.378387698	3.153957498	0.160496475	0.911729869	2.417066389			
RU17	-1.71410687			-0.690590541		-0.36024807	0.37741751	-0.706990994	0.025682531	0.224527766	2.057711732	0.494530275	1.319226855	1.089247379	0.510960395			
TNR6A	0			-0.787648461		0.620580664	0	-0.0479136	1.439314625	-0.598484467	0.749016625	1.614529269	1.847140564	1.972543734	1.175688447			
STRBP	-2.005726377			-1.254863618		1.165840938	1.932321092	2.298427462	2.452082104	0	1.87495106	1.330255614	1.847140564	1.279396553	1.940218267			
HNRPQ	-0.263223675			1.664950194		-0.22671606	0.020740665	-0.999519356	-0.199503321	1.288463793	2.493957236	2.222739616	1.916796134	1.612706565	2.43653015			
ILF3	-2.276729705			-1.151163936		1.026045253	-1.684007831	-0.474793755	0.403807322	0.224527766	0.672602162	1.973899539	0.924881842	1.303283679	2.618651578			
SFRS5	-1.955777055			1.072350745		0.957051788	2.10018112	0.429011744	0.701716147	2.937970737	1.250399398	0.799915034	4.192056401	0.572753117	1.330442584			
MS1H	0			-2.636031327		0	0	0	-0.597318496	0	2.100590948	2.3906326	0	2.459358046	3.41620093			
RBM22	-2.005726377			-0.130627647		0	1.396729723	-1.206564581	-0.142178206	0	1.38754728	0.773772048	0.150954733	2.053890373	0.798640368			
SNRPA	-2.963209082			-0.382182722		0	2.089875666	1.123844033	1.118818584	2.427145113	0.593624145	0.760276421	2.322260261	0.889511325	0.418413417			
IREB1	0			-0.788757513		4.02804271	0.937031397	-1.146100119	0.944741776	-2.490300411	1.141150629	0.958574027	2.833083924	1.758374839	2.643013778			
SYWC	-1.217198733			0.857651824		0.995273586	0.001839129	1.089628952	0.977839741	2.706271286	0.787828816	-0.368061994	2.08382951	2.308955373	3.24638943			
RAE1L	-2.650865905			-0.928018061		0	0	-0.74063854	-0.038298949	0	2.449054922	1.049768068	1.223648953	2.802111116	2.198239618			

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GENES	White	Blood	Cells	Vascular	endothelium	Thyroid	Lung	Prostate	Breast	Liver	Brain	cortex	Brain	astrocyte	Stomach	Colon	Brain	cerebellum
ROA2	-1.257182245	0	0	0.149703954	-0.47803011	-0.444621541	0.439217336	-0.072681474	0.189853283	0.913443035	1.405237403	0.263435099	1.051307459	1.852579056				
NCBP1	0	0	0	0	0	0	0	1.347207839	0	0	0	0	0	1.196434462				
RM12	-2.7555125	-0.277934411	1.165840938	1.273504875	0.299675114	1.228025435	1.288463793	1.457116824	-0.289466194	3.690033788	2.010343391	2.205911321	2.205911321					
DHX30	-2.31439508	0.004617676	-1.276538801	0.478404405	1.3172738	1.272863816	0.224529423	0.487084258	0.724342037	1.136903337	3.627607592	0.146424329	0.146424329					
TADBP	-2.431453452	-0.483721338	-0.478030856	0.924757672	0.369812928	0.629667341	2.021682653	0.712527074	-0.389019677	0.239077006	3.241782649	1.119411172	1.119411172					
HNRL1	-0.206579264	-0.565615394	0.438258796	-1.911578948	0.271397258	2.038424735	1.293861039	1.350333813	2.098388032	-0.472529246	2.578966048	3.126506118	3.126506118					
QKI	-1.292286173	1.632725032	0.215115556	-1.851059124	2.223296184	0.190281704	-1.79715323	0.927516768	3.196939475	2.367511039	0	0.835423891	0.835423891					
KHDRBS2	0	0	0	0	0	0	0	1.210946432	-1.291631648	0	0	0	2.931924152					
TARB1	-2.046915767	-1.637850427	1.165840938	2.443146716	1.007200164	2.075706795	0	-0.133004349	3.066945345	2.252605672	2.283632624	1.008272256	1.008272256					
DICER	-0.673585703	0.815551218	-0.478033183	0.243885325	1.199814169	1.35739711	1.328535473	1.229443018	2.748494524	0.042718279	1.88581735	3.644966751	3.644966751					
HNRPG	-1.616030422	-1.20214624	-0.273238439	0.284940122	0.082449522	1.64376847	2.404165573	1.199316402	0.270432407	3.038421484	2.227833053	2.575218003	2.575218003					
ABC3G	-3.615165635	0	0	2.089875666	0	1.831098765	0	2.088492954	1.306447259	0	1.185019316	1.196434462	1.196434462					
HTF9C	-2.069667683	-0.928019352	1.165840938	3.298069926	0.709649336	0.217945391	2.404165573	0.433861193	1.32057812	1.134182228	0.974722731	2.908002093	2.908002093					
RS4Y1	-4.713773438	-3.91893109	0	-3.452023567	0.132138577	-1.251379154	-2.202618891	2.169084925	4.302180887	2.057622471	1.071831228	4.357569994	4.357569994					
R51A1	0	-1.942884147	0	0	1.861753395	1.210946432	1.328535473	1.960521454	2.414037018	2.662850583	1.972543734	2.310989675	2.310989675					
IF4H	-0.665092451	0.334251172	-0.126633383	-0.141777754	0.496983397	0.658908418	1.106145722	1.168209538	0.173008466	0.351374613	1.540333412	0.8086918	0.8086918					
CUGBP2	-2.756656221	-1.211886855	0	-1.911578948	-0.016281851	1.709835131	1.293861039	-0.148965324	1.202104875	0	2.014043985	0.649267925	0.649267925					
RBM5	-1.857210804	-1.229475664	2.418606578	0.868041245	0.419409453	1.406860836	1.293861039	2.029951291	1.73235897	0.374096707	-0.201454703	0.231231578	0.231231578					
KRR1	-1.312580542	-2.348351643	-1.931556194	1.932321092	-0.419308944	-0.018466186	0	3.01867995	1.716038659	2.479574124	2.747040118	0.168212751	0.168212751					
CDKN2AIP	-0.247770308	0.557394315	-1.238490913	-1.506115319	-0.04749136	-0.596993807	0	1.285120045	1.847140564	1.847140564	1.775212952	1.775212952	1.775212952					
RBM28	1.105998278	-2.047128813	0	0	-0.550132401	0.652714458	1.328535473	2.736741313	1.384715616	1.354408879	0.578417621	1.889581643	1.889581643					
HNRL	-0.238802939	-0.094496977	-0.861023135	0.020741552	1.073774281	1.09444719	1.328535473	1.204664863	0.617026754	0.126928421	2.861617854	2.011572073	2.011572073					
DSRAD	-2.565808768	-0.983049911	-0.765715255	3.030935311	-0.091407984	2.041144823	-0.908757852	0.26335938	0.8011546	0.844105581	1.964713373	1.512762881	1.512762881					
U2AF1L4	-1.592876625	-0.234872172	1.431510014	1.526855984	-1.909243201	2.582781719	0.189853332	-0.137547895	0.440197446	1.916796134	0	0.156126938	0.156126938					
HNRL	-3.297623494	-0.223867428	0.461517457	0.451525969	-0.204417269	0.202823245	0.62517263	0.703672023	0.27069618	0.056310976	1.640124704	1.11853434	1.11853434					
ACF	-0.697571178	-1.249738399	0	-1.911578948	-2.121694911	1.831098765	1.328535473	0.999091553	1.355319557	-1.696180587	1.279396553	1.792775619	1.792775619					
SYMC	-1.293704451	-1.046915513	0	2.779620193	0.0612869	0.039177486	1.328535473	1.029677629	-0.289466194	2.047554769	2.429950533	1.111559338	1.111559338					
EXOS7	-1.803569456	-1.637850427	0.022741538	-2.758876386	-1.481496701	1.614272725	3.08561868	0.961064966	2.819502126	2.252605672	2.020682288	3.476140957	3.476140957					
ABEC1	0	0	0	0	0	1.196482824	1.198668763	0	0	1.746561595	0	1.173555864	1.173555864					
RBMS2	0	0	0	0	0	0	1.25440733	1.198668763	0	0	0	0	0					
DAZP1	-1.00601907	-1.333487903	-1.238490913	2.443146716	0.399877494	1.148137153	1.98700822	1.900968918	-0.663347971	2.511378167	1.400336269	2.202498424	2.202498424					
RBM19	0	0	0	1.396729723	1.16860466	1.347207839	0	1.788146283	-1.357104966	0	1.19267017	0	0					
TDRKH	0	0	0	0	0	0	0	0	-1.163934877	0	0	0	0					
RENT1	-0.280686007	-1.948009373	0	1.526855984	0	1.020004213	0	1.937837654	0	1.223648953	2.022292981	0	0					
STAU1	-0.98525325	-0.741879338	0.332894545	-0.490080523	-0.427884173	0.501418105	-0.320970804	0.742463158	3.446514856	0.402274409	0.714839063	1.111469409	1.111469409					
PTBP1	-1.113643768	-0.210109454	1.165840938	2.913149259	0.486012816	1.183338256	1.98700822	2.011146089	0.946894127	2.54982791	1.350355445	1.517019472	1.517019472					
DKFZp686l	-0.401038746	-2.924945473	-0.295708909	3.298069926	1.943152313	0.575283093	0.094661056	2.393608826	2.380748189	0.249381284	-1.744442998	0.743478416	0.743478416					
SFRS3	-0.573500262	-0.006424735	-0.111785849	-0.895549343	-0.053533541	0.406649698	0.326477289	0.858395512	0.865586348	0.393476409	2.227188913	2.477201724	2.477201724					
DUS11	-1.369221953	0.410086577	0	-1.911578948	-1.718737061	0.198561596	1.328535473	2.263797486	2.354902363	1.746561595	1.185019316	2.248649363	2.248649363					
CSTFT	-2.07042224	-0.928018061	0.438258796	1.007200164	2.930942521	-1.60190448	1.084149313	1.538561639	0.756839721	-0.03338292	1.102240134	1.102240134	1.102240134					
MCES	-2.046915767	0	1.165840938	0	-0.016281851	0.860455153	0.860455153	2.427145113	2.633075187	0.532763793	2.740704531	1.66020505	1.66020505					
MGC27016	0.267509428	-0.318068305	-0.583391621	-2.422401615	-0.091411583	0.501292926	0	0.231410896	2.389789785	1.125085429	1.19267017	1.565561289	1.565561289					
SFRS4	-0.653234669	0.458274667	1.571308124	2.913149259	-1.114897104	0.590617257	-1.104007707	0.90692633	1.066847841	0.853649927	1.13486919	2.376569486	2.376569486					
LARP5	0	1.119110752	0	0	0	1.221130162	0	1.858331858	0	1.847140564	1.360743193	1.334745771	1.334745771					
SAM68	-1.091613458	1.266479734	-0.215665562	-1.077870145	-0.186723656	0.66441452	0.978313011	0.582766109	1.280173736	0.122049671	0.771726591	0.701294123	0.701294123					
RT15	-1.17951021	-0.234872172	-0.072568075	0.583760348	0.907661919	2.747386292	-2.490300411	1.441064982	-0.220081586	2.662850583	3.582915979	1.940218267	1.940218267					
FUBP1	-0.686151127	-1.112684856	0	2.124028839	0.105866739	0.743827512	1.293861039	1.471748911	0.802306396	2.845173302	3.582916257	2.034968208	2.034968208					
IGF2BP1	0	0	0	0	0	1.248238463	0	1.255581849	0	1.88581735	0	1.099628439	1.099628439					
M10L1	0	-2.348351643	0	0	0	0	0	0	0	0	0	0	0					
DRBP1	-1.398104348	0	1.165840938	0	-2.121694911	2.35302152	-1.279547052	2.036592755	-0.031763561	0	0	1.940218267	1.940218267					
IREB2	0	0	0	0	0	0	0	1.165186237	0	1.125085429	0	1.196434462	1.196434462					
GRSF1	-2.067266475	0.02328712	-1.238490913	1.526855984	0.83616196	0.596339757	1.293861039	0.294853338	1.033527634	2.252605672	1.400334174	0.418413417	0.418413417					
CUGB1	-1.37411803	1.119110752	0	0	0	1.124682813	0	1.948729029	2.048466737	0	-0.03338292	1.099628439	1.099628439					
SCAND1	-0.150334612	-0.855287481	0.232815559	0.98922569	0.803268539	2.130755017	0.412998541	0.918023117	1.280075604	0.812520594	0.360853529	2.306214618	2.306214618					
FXR1	-1.353768587	0.569359628	-0.22671606	-1.506115319	0.27383986	0.584649645	1.799291244	0.89255358	1.544355089	2.322260261	2.528995303	3.615187165	3.615187165					
RBPMS	1.10598278	0.652703604	-0.765717042	-2.758876386	-1.19002846	0.091573129	-0.908757852	2.449054922	-2.050252146	2.511378167	1.88581735	2.295046751	2.295046751					
COVA1	-1.369221953	1.119110752	0	0	-1.21609602	1.210946432	1.198668763	1.726157271	1.740127521	0	1.19267017	1.970036748	1.970036748					

Supplementary file 12

GENES	White Blood Cells	Vascular endothelium	Thyroid	Lung	Prostate	Breast	Liver	Brain cortex	Brain astrocyte	Stomach	Colon	Brain cerebellum
NUCL	-1.175158395	0.042683921	0.146122684	-0.975101663	-0.003238422	-0.065824646	1.214358403	1.345172755	0.563667737	0.447224721	3.469019185	1.734390898
TRSPAP1	-0.195089147	-0.928018061	1.571308124	1.932321092	0.101605835	0.985183651	0.917674946	1.228157349	2.008573699	0.150959977	2.571313678	1.220041763
RBM38	-1.857206566	-0.13575144	0	0.426205773	-1.114897104	1.076934477	0.224527766	1.748191464	0.237022154	2.734523341	-0.470442774	-0.38835008
SURF6	-2.046915767	0	-1.238409013	0	-1.21609602	0.249383867	0.189853332	1.343446879	0.005091356	0	1.972543734	1.212375403
ELAV4	0	0	0	0	0	1.137951585	0	-0.780603887	1.643149246	0	0	3.291817689
RED2	0	0	0	0	0	-1.251379154	0	0	0	0	0	0
PM14	-2.63945011	-0.971083025	0.12810463	0.583760219	-0.030379303	0.367983135	1.477293527	2.024779698	0.517939395	0.246269297	1.677028982	1.486196537
RBM6	-2.690000893	-0.712895012	0.032795017	0.020740726	0.601737221	0.606981523	0.224526222	0.337000204	3.559422606	1.096083304	2.018171008	1.160095228
PPRC1	-0.263223675	0	0	0	0.754433077	2.508826606	0	1.722824809	1.355319557	1.847140564	3.557967485	0.823873397
PRGC2	-1.353768587	0	0	0	0	1.210946432	0	1.27314526	1.355319557	0	1.19267017	-0.967885918
NUFP1	-1.803569456	-1.249738399	-1.238409013	0	0.085066319	0.120993751	1.198668763	1.99865439	-1.357104966	1.223648953	1.612706565	1.905522584
HNRPK	-1.763706946	-1.051705672	-0.206098245	-0.598294914	0.184389174	-0.253157762	-0.097828911	2.178534493	1.280578318	0.220615547	2.298027124	1.831097364
DHX9	-3.24700059	-1.435026519	-0.22671606	-0.266940299	-0.395633265	-0.401111426	0.883000513	2.247526413	1.416096069	0.22061335	1.12133399	3.270287312
NOL3	-1.72978477	-2.047130104	1.858989677	0.937032506	0.309936716	3.33014167	0.094661056	1.256844117	3.151924154	2.486342285	3.293994429	1.633431282
RL12	-1.029889223	0.423180785	-0.068896819	-0.028049711	0.085064424	0.986229388	2.657952915	2.13907224	1.808659904	0.33728876	2.907141829	2.404354277
DDX17	0.486566571	-0.82889862	-1.238409013	2.913149259	1.278348334	1.335109002	0	0.599552139	-0.542644113	3.204525348	2.053890373	1.216831152
SYG	0.088700291	-1.804602648	1.026045253	1.416674282	-0.056933922	0.828609874	2.714827185	1.520198244	-1.54237649	-0.159451011	3.217473361	0.875586995
ZNF638	-2.472732849	-1.435026519	0.332898072	-2.758876386	0.585199503	1.324455847	0.735351846	0.82397791	2.365487272	-0.254505131	1.360743193	2.667343036

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Supplementary file 13

Name	# of Studies	Study Details
TNRC6C	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 4.672 Pvalue: 5.9E5
RPS4Y1	2	
		Study: Finak_Breast Experiment Type: mRNA Analysis: Breast Stroma Type Class 1: Normal Breast (6) Class 2: Breast Carcinoma (53) Ttest: 14.179 Pvalue: 1.1E6
		Study: Karnoub_Breast Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (15) Class 2: Invasive Ductal Breast Carcinoma (7) Ttest: 3.045 Pvalue: 0.007

Name	# of Studies	Study Details
DRB1	1	
		Study: Karnoub_Breast Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (15) Class 2: Invasive Ductal Breast Carcinoma (7) Ttest: 7.112 Pvalue: 3.3E6
ASCC1	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 5.029 Pvalue: 4.8E4
ZNF638	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 5.642 Pvalue: 1.1E
RNASEL	2	
		Study: Finak_Breast Experiment Type: mRNA Analysis: Breast Stroma Type Class 1: Normal Breast (6) Class 2: Breast Carcinoma (53) Ttest: 10.924 Pvalue: 1.4E6
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 3.643 Pvalue: 0.00
COVA1	1	

		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 5.197 Pvalue: 6.3E5
CUGBP1	1	
		Study: Finak_Breast Experiment Type: mRNA Analysis: Breast Stroma Type Class 1: Normal Breast (6) Class 2: Breast Carcinoma (53) Ttest: 5.121 Pvalue: 0.002
HTATSF1	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 5.686 Pvalue: 7E6
RBM27	2	
		Study: Finak_Breast Experiment Type: mRNA Analysis: Breast Stroma Type Class 1: Normal Breast (6) Class 2: Breast Carcinoma (53) Ttest: 6.856 Pvalue: 6.7E4
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 3.612 Pvalue: 0.004
FMR1	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 4.948 Pvalue: 1.4E5
ELAVL4	1	

		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 4.722 Pvalue: 2.4E5
MYEF2	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 3.437 Pvalue: 0.002

NCBP1	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 6.689 Pvalue: 2.6E6
SLC4A1A P	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 5.296 Pvalue: 1.1E
ZCRB1	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 6.609 Pvalue: 3.1E
RBM35B	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 7.038 Pvalue: 2.7E8
DSRAD	2	

		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 4.496 Pvalue: 7.8E4
		Study: Karnoub_Breast Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (15) Class 2: Invasive Ductal Breast Carcinoma (7) Ttest: 3.954 Pvalue: 0.001
JOSD2	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 3.698 Pvalue: 0.001
DDX53	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 3.255 Pvalue: 0.009
TIA1	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 6.903 Pvalue: 1.2E7
POLDIP3	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 3.252 Pvalue: 0.005
APOBEC 1	1	

		Study: Karnoub_Breast Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (15) Class 2: Invasive Ductal Breast Carcinoma (7) Ttest: 3.284 Pvalue: 0.004
ELAVL3	1	
		Study: Finak_Breast Experiment Type: mRNA Analysis: Breast Stroma Type Class 1: Normal Breast (6) Class 2: Breast Carcinoma (53) Ttest: 4.311 Pvalue: 0.007
IGF2BP1	1	
		Study: Turashvili_Breast Experiment Type: mRNA Analysis: Breast Ductal Tissue Type Class 1: Adjacent Normal Ductal (5) Class 2: Invasive Ductal Carcinoma (5) Ttest: 3.604 Pvalue: 0.01
NCBP2	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 5.524 Pvalue: 1.5E5
RBM35A	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 6.38 Pvalue: 8.2E6
RBP56	1	
		Study: Karnoub_Breast Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (15) Class 2: Invasive Ductal Breast Carcinoma (7) Ttest: 2.968 Pvalue: 0.01
PPARGC 1B	1	

		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 3.548 Pvalue: 0.001
PNPT1	1	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 8.435 Pvalue: 5.2E7
RTCD1	3	
		Study: Richardson_Breast_2 Experiment Type: mRNA Analysis: Breast Type Class 1: Normal Breast (7) Class 2: Breast Carcinoma (40) Ttest: 6.031 Pvalue: 1.2E5
		Study: Finak_Breast Experiment Type: mRNA Analysis: Breast Stroma Type Class 1: Normal Breast (6) Class 2: Breast Carcinoma (53) Ttest: 5.569 Pvalue: 5.2E4
		Study: Sorlie_Breast Experiment Type: mRNA Analysis: Breast Type Class 1: Benign Breast (7) Class 2: Breast Carcinoma (78) Ttest: 3.609 Pvalue: 0.008

Name	# of Studies	Study Details
SYNCRIP	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 4.18 Pvalue: 2.8E4
TRBP2	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.202 Pvalue: 0.002
SR140	2	
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 4.802 Pvalue: 3.3E5
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 2.882 Pvalue: 0.008
CPSF5	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.738 Pvalue: 3.3E9
FUBP2	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.718 Pvalue: 4.8E6

HNRPDL	1	
		Study: Hong_Colon Experiment Type: mRNA Analysis: Colonic Mucosa Type Class 1: Normal (10) Class 2: Adjacent NonTumoral (12) Ttest: 5.691 Pvalue: 2E5
PAIP1	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.81 Pvalue: 1.8E9
PPIE	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 5.685 Pvalue: 2.2E7
PNPT1	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.627 Pvalue: 0.003
HRB	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.887 Pvalue: 5.3E6
NOL8	2	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 5.773 Pvalue: 4.2E6

		Study: Zou_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colonic Epithelium (8) Class 2: Invasive Colon Carcinoma (9) Ttest: 3.273 Pvalue: 0.007
NCBP2	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 5.431 Pvalue: 6E7
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 4.782 Pvalue: 1.9E5
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.284 Pvalue: 0.003
RPL22	2	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.213 Pvalue: 0.003
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 2.908 Pvalue: 0.00
PSMA1	3	

		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.964 Pvalue: 8.8E10
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 5.387 Pvalue: 5.7E6
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 3.159 Pvalue: 0.003
FXR1	2	
		Study: Hong_Colon Experiment Type: mRNA Analysis: Colonic Mucosa Type Class 1: Normal (10) Class 2: Adjacent NonTumoral (12) Ttest: 3.376 Pvalue: 0.004
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 2.934 Pvalue: 0.004
PTBP1	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.135 Pvalue: 5E8

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.284 Pvalue: 0.003
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.12 Pvalue: 0.004
CSTF3	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.093 Pvalue: 4.2E8
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.149 Pvalue: 0.004
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 2.942 Pvalue: 0.006
SRP72	2	
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 5.124 Pvalue: 2.2E5

		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 2.734 Pvalue: 0.008
FUS	1	

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.606 Pvalue: 0.001
KRR1	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.555 Pvalue: 6.5E4
HNRPM	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 7.052 Pvalue: 6.9E10
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.639 Pvalue: 9E4
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 2.698 Pvalue: 0.01
DHX9	1	

		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.193 Pvalue: 7.3E5
PABP4	3	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 4.028 Pvalue: 5E4
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 3.484 Pvalue: 0.001
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.409 Pvalue: 0.002
ILF3	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 9.196 Pvalue: 8.1E14
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 4.259 Pvalue: 1.5E4

		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 3.592 Pvalue: 7.4E4
SNRPB2	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.676 Pvalue: 4.2E9
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 4.078 Pvalue: 1.6E4
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 2.98 Pvalue: 0.005
DKC1	2	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 9.694 Pvalue: 6.8E15
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 8.545 Pvalue: 3.8E
NUFIP1	1	

		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 12.808 Pvalue: 5.7E21
RBM9	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.746 Pvalue: 8.7E4

HNRPUL1	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.194 Pvalue: 7.2E5
FBL	4	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 8.417 Pvalue: 4.2E1
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 6.844 Pvalue: 7E7
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 5.473 Pvalue: 4.2E6

		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 4.682 Pvalue: 4.1E5
EXOSC7	1	
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 4.221 Pvalue: 2E4
RBM8A	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 7.731 Pvalue: 4.4E11
NOL3	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.699 Pvalue: 0.001
FUBP1	3	
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 4.69 Pvalue: 4.3E5
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.031 Pvalue: 1.5E4

		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 3.509 Pvalue: 0.001
HNRPG	3	
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 6.324 Pvalue: 3.4E7
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 4.065 Pvalue: 3.6E4
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 2.765 Pvalue: 0.008
HARS	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.586 Pvalue: 5.8E4
HNPRU	2	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.83 Pvalue: 7.6E4

		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.605 Pvalue: 0.001
EIF3S9	2	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 6.376 Pvalue: 1.1E6

		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.698 Pvalue: 5.2E4
EIF3S4	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.78 Pvalue: 9.7E6
RPL12	2	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 7.558 Pvalue: 3.4E8
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 5.724 Pvalue: 1.8E7
NKRF	1	

		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.707 Pvalue: 3.9E4
NCL	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 8.445 Pvalue: 1.5E12
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.927 Pvalue: 0.001
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 3.372 Pvalue: 0.001
LSM5	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.044 Pvalue: 0.003
SAFB1	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.815 Pvalue: 7E6
DSRAD	1	

		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 2.822 Pvalue: 0.006
NHP2L1	2	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.25 Pvalue: 0.003
		Study: Hong_Colon Experiment Type: mRNA Analysis: Colonic Mucosa Type Class 1: Normal (10) Class 2: Adjacent NonTumoral (12) Ttest: 3.117 Pvalue: 0.006
NPM1	4	
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 9.858 Pvalue: 1.8E11
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.77 Pvalue: 2.1E9
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 5.672 Pvalue: 5.3E7

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 5.075 Pvalue: 2.9E5
MARS	3	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 7.295 Pvalue: 2.1E10

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 5.509 Pvalue: 7.7E6
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.909 Pvalue: 4.4E4
MRPL23	2	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.933 Pvalue: 2E4
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 4.006 Pvalue: 4.2E4
HNRPL	1	

		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.045 Pvalue: 0.004
NONO	4	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 7.505 Pvalue: 8.1E11
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 6.206 Pvalue: 1.2E6
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 6.099 Pvalue: 1.8E6
		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 3.885 Pvalue: 3.1E4
LA	2	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 7.96 Pvalue: 1.1E11

		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 4.376 Pvalue: 1.3E4
TIAR	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 5.328 Pvalue: 9.6E7
RPL34	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 4.604 Pvalue: 8.5E5
MRPL3	2	
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 4.567 Pvalue: 1.6E4
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.126 Pvalue: 0.004
RBM26	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.378 Pvalue: 0.00
RPL13	2	

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 6.19 Pvalue: 1.1E6
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.615 Pvalue: 9.7E4
RPL18	1	

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.77 Pvalue: 8E4
SLBP	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 3.999 Pvalue: 1.4E4
RBPMS	1	
		Study: Hong_Colon Experiment Type: mRNA Analysis: Colonic Mucosa Type Class 1: Normal (10) Class 2: Adjacent NonTumoral (12) Ttest: 4.42 Pvalue: 4.9E4
RPLP2	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 2.863 Pvalue: 0.008
RPL7	1	

		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.015 Pvalue: 0.005
RPS5	2	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 5.467 Pvalue: 8.3E6
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 2.958 Pvalue: 0.004
SIAHBP1	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.019 Pvalue: 0.009
RPL28		
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.952 Pvalue: 4.1E6
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 2.935 Pvalue: 0.007
RPS17	1	

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.212 Pvalue: 0.004
RBM19	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 3.874 Pvalue: 7E4
PCBP1	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 7.708 Pvalue: 3.2E11
PABP2	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 2.699 Pvalue: 0.009
RBM14	1	
		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 4.354 Pvalue: 2.8E4
RBM10	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 4.784 Pvalue: 7.9E6
RPS3	3	

		Study: Graudens_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (12) Class 2: Colorectal Carcinoma (18) Ttest: 4.974 Pvalue: 3.3E5
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		Study: Alon_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (22) Class 2: Colon Adenocarcinoma (40) Ttest: 4.086 Pvalue: 2.4E4
		Study: Notterman_Colon Experiment Type: mRNA Analysis: Colon Type Class 1: Normal Colon (18) Class 2: Colon Adenocarcinoma (18) Ttest: 3.688 Pvalue: 9E4
ELAV1	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.61 Pvalue: 6.4E9

Name	# of Studies	Study Details
DKFZp686F02235	1	
		Study: Hong_Colon Experiment Type: mRNA Analysis: Colonic Mucosa Type Class 1: Normal (10) Class 2: Adjacent NonTumoral (12) Ttest: 4.005 Pvalue: 8.3E4
RNASE1	1	
		Study: Ki_Colon Experiment Type: mRNA Analysis: Tissue Type Class 1: Adjacent Normal Colon Mucosa (28) Class 2: Primary Colon Carcinoma (53) Ttest: 6.219 Pvalue: 5.4E8

Name	# of Studies	Study Details
NKRF	1	
		Study: Hippo_Gastric Experiment Type: mRNA Analysis: Gastric Tissue Type Class 1: Normal (8) Class 2: Gastric Carcinoma (22) Ttest: 3.227 Pvalue: 0.01 Y07707_at Reporter:
FUBP1	1	
		Study: Hippo_Gastric Experiment Type: mRNA Analysis: Gastric Tissue Type Class 1: Normal (8) Class 2: Gastric Carcinoma (22) Ttest: 3.992 Pvalue: 0.002 U05040_at Reporter:
CSTF2	1	
		Study: Hippo_Gastric Experiment Type: mRNA Analysis: Gastric Tissue Type Class 1: Normal (8) Class 2: Gastric Carcinoma (22) Ttest: 3.902 Pvalue: 0.004 M85085_at Reporter:
FXR1	1	
		Study: Hippo_Gastric Experiment Type: mRNA Analysis: Gastric Tissue Type Class 1: Normal (8) Class 2: Gastric Carcinoma (22) Ttest: 3.795 Pvalue: 0.002 U25165_at Reporter:
NONO	1	
		Study: Hippo_Gastric Experiment Type: mRNA Analysis: Gastric Tissue Type Class 1: Normal (8) Class 2: Gastric Carcinoma (22) Ttest: 3.865 Pvalue: 9E4
EIF3S9	1	

		Study: Hippo_Gastric Experiment Type: mRNA Analysis: Gastric Tissue Type Class 1: Normal (8) Class 2: Gastric Carcinoma (22) Ttest: 3.865 Pvalue: 9E4
RO60	1	
		Study: Hippo_Gastric Experiment Type: mRNA Analysis: Gastric Tissue Type Class 1: Normal (8) Class 2: Gastric Carcinoma (22) Ttest: 4.128 Pvalue: 0.002

Name	# of Studies	Study Details
RPL7	1	

		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.778 Pvalue: 0.001
SIAHBP1	3	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.722 Pvalue: 4.9E6
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 4.539 Pvalue: 5E5
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Benign Liver Disease (7) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.377 Pvalue: 9.5E4
HNRPG	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.957 Pvalue: 1.1E4
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.878 Pvalue: 0.001
HNRPLL	1	

		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 5.503 Pvalue: 1.4E7
SNRPA	2	
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 5.169 Pvalue: 3.5E5
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 2.663 Pvalue: 0.009
NCBP2	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.355 Pvalue: 9.7E4
PTBP1	1	
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.339 Pvalue: 0.005
SF3B14	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.406 Pvalue: 8.2E4
SF3B4	2	

		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 9.428 Pvalue: 3.1E17
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 5.941 Pvalue: 4.8E6
RPS5	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 6.007 Pvalue: 1.3E8
EIF3S9	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 5.002 Pvalue: 1.4E6
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.406 Pvalue: 0.003
PAPOLG	1	
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 4.392 Pvalue: 2.3E4
FUBP1	1	

		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.75 Pvalue: 2.4E4
DKC1	1	
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.097 Pvalue: 0.007
MGC104 33	1	
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 4.062 Pvalue: 5.9E4
IGF2BP2	1	
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 8.36 Pvalue: 3.7E10
NUFIP1	1	
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.634 Pvalue: 0.002
TIA1	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 2.803 Pvalue: 0.006
PNPT1	1	

		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.224 Pvalue: 0.006
RNMT	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 5.015 Pvalue: 1.5E6
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 5.312 Pvalue: 3.9E6
RCAN3	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.914 Pvalue: 1.3E4
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.348 Pvalue: 0.002
IGF2BP3	2	
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 5.735 Pvalue: 1.6E6

		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.961 Pvalue: 1.1E4
HNRPUL 1	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 6.618 Pvalue: 4.7E10
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 2.989 Pvalue: 0.005
RBM9	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.079 Pvalue: 6.8E5
CPSF6	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.514 Pvalue: 1.2E5
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 4.717 Pvalue: 4.3E4

RPL18	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.168 Pvalue: 0.002
GARS	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 5.811 Pvalue: 3E8
RBM8A	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.311 Pvalue: 0.00
APOBEC 1	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Benign Liver Disease (7) Class 2: Hepatocellular Carcinoma (104) Ttest: 6.621 Pvalue: 2.8E9
RBM28	1	
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.967 Pvalue: 5E4
C20orf11 9	1	
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.315 Pvalue: 0.002
ANKRD1 7	1	
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 4.161 Pvalue: 1.5E4
NONO	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.604 Pvalue: 7.9E6
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.336 Pvalue: 0.003
MARS	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.234 Pvalue: 0.001
TARDBP	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.049 Pvalue: 7.7E5
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 2.747 Pvalue: 0.009
SYNJ2	1	
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.883 Pvalue: 7.7E4
LRPPRC	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 4.132 Pvalue: 5.8E5
RBM10	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.066 Pvalue: 0.003
		Study: Wurnbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.087 Pvalue: 0.007
AKAP1	1	

		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 6.448 Pvalue: 1.1E9
FXR1	3	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 5.523 Pvalue: 1.2E7

Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.651 Pvalue: 8.2E4
Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Benign Liver Disease (7) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.655 Pvalue: 0.007

Name	# of Studies	Study Details
MRPS15	1	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.886 Pvalue: 1.5E4
PPIE	1	
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 2.966 Pvalue: 0.005
PAPOLA	1	
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 5.697 Pvalue: 6.3E6
LARP4	3	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 11.805 Pvalue: 4E24
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Benign Liver Disease (7) Class 2: Hepatocellular Carcinoma (104) Ttest: 5.518 Pvalue: 6E4
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 2.987 Pvalue: 0.006
IREB1	2	

		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 5.367 Pvalue: 2.6E7
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 4.268 Pvalue: 5.5E4
RBM35B	2	
		Study: Chen_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Nontumor Liver (76) Class 2: Hepatocellular Carcinoma (104) Ttest: 3.959 Pvalue: 1.1E4
		Study: Wurmbach_Liver Experiment Type: mRNA Analysis: Liver Type Class 1: Normal Liver (10) Class 2: Hepatocellular Carcinoma (35) Ttest: 3.18 Pvalue: 0.003

Name	# of Studies	Study Details
RPL18	4	
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 6.379 Pvalue: 4.5E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.229 Pvalue: 0.003
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 3.258 Pvalue: 0.003
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 2.79 Pvalue: 0.009
RTCD1	5	

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.237 Pvalue: 4.8E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.124 Pvalue: 7.3E4

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 3.932 Pvalue: 8.4E4
		Study: Powell_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: NonMalignant Lung (11) Class 2: Lung Adenocarcinoma (11) Ttest: 3.462 Pvalue: 0.006
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Large Cell Lung Cancer (4) Ttest: 4.408 Pvalue: 0.01
PABP4	2	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 7.14 Pvalue: 4.7E8
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 2.813 Pvalue: 0.007
PSMA1	5	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 5.168 Pvalue: 4.5E5

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.306 Pvalue: 6.7E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 3.482 Pvalue: 0.002
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.51 Pvalue: 0.006
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 2.753 Pvalue: 0.008
MRPL23	3	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 5.237 Pvalue: 1.2E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 4.013 Pvalue: 3.3E4

		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 3.615 Pvalue: 0.001
CNOT4	2	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 6.328 Pvalue: 3E7
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.196 Pvalue: 2.9E4
CSTF2	3	
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 4.666 Pvalue: 2.4E5
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 4.556 Pvalue: 0.003
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 2.946 Pvalue: 0.006

PTBP1	5	
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 6.157 Pvalue: 1.6E6
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.031 Pvalue: 3.6E4
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 4.573 Pvalue: 0.002
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 2.9 Pvalue: 0.005
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Small Cell Lung Cancer (4) Ttest: 3.505 Pvalue: 0.01
RBM14	2	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.204 Pvalue: 1.1E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 3.979 Pvalue: 0.002
SYNJ2	9	
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 5.125 Pvalue: 1E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.45 Pvalue: 4.9E5
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 5.631 Pvalue: 5.8E5
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.84 Pvalue: 3.5E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.433 Pvalue: 3.6E4
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Small Cell Lung Cancer (4) Ttest: 6.412 Pvalue: 6.2E4
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 4.597 Pvalue: 0.002
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.565 Pvalue: 0.003
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 3.007 Pvalue: 0.008
TRBP2	4	
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 5.183 Pvalue: 2E4
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 3.52 Pvalue: 0.001
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 4.306 Pvalue: 0.002
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 3.942 Pvalue: 0.009
RBM34	8	

		Study: Yamagata_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (3) Class 2: Squamous Cell Lung Carcinoma (11) Ttest: 7.044 Pvalue: 1.6E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.824 Pvalue: 1.8E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.118 Pvalue: 5.5E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 3.798 Pvalue: 0.001

		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 4.797 Pvalue: 0.001
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 3.524 Pvalue: 0.002
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 3.348 Pvalue: 0.004
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 2.75 Pvalue: 0.008
SFRS1	6	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.219 Pvalue: 1.8E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 5.304 Pvalue: 2.8E5

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 5.765 Pvalue: 2.9E5
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Small Cell Lung Cancer (4) Ttest: 4.92 Pvalue: 0.002
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.957 Pvalue: 0.003
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 2.981 Pvalue: 0.009
SART3	7	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 7.1 Pvalue: 3.5E8
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 4.973 Pvalue: 8.9E5

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.936 Pvalue: 7.7E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.431 Pvalue: 0.003
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 3.339 Pvalue: 0.004
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 2.944 Pvalue: 0.006

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 2.81 Pvalue: 0.009
DKC1	9	
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 7.492 Pvalue: 1.4E7

		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 5.124 Pvalue: 4.3E6
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 5.669 Pvalue: 1E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 3.802 Pvalue: 5.5E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 3.761 Pvalue: 6.3E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.804 Pvalue: 8.9E4
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 3.997 Pvalue: 0.002
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Large Cell Lung Cancer (4) Ttest: 3.721 Pvalue: 0.006

		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.209 Pvalue: 0.009
TUT1	1	
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.798 Pvalue: 0.006
RBM16	3	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 6.539 Pvalue: 1.3E6
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.924 Pvalue: 9.1E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 3.558 Pvalue: 0.005
MARS	9	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 8.176 Pvalue: 3.5E9

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 6.646 Pvalue: 5.7E7
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 4.682 Pvalue: 1.7E5
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 4.782 Pvalue: 3.2E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 5.349 Pvalue: 2.8E4
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.414 Pvalue: 0.001

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 3.276 Pvalue: 0.002
		Study: Powell_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: NonMalignant Lung (11) Class 2: Lung Adenocarcinoma (11) Ttest: 3.26 Pvalue: 0.004

		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 4.845 Pvalue: 0.005
RBM12	1	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 3.65 Pvalue: 0.005
U2AF2	5	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 7.54 Pvalue: 2.1E8
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 7.066 Pvalue: 1.8E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 6.33 Pvalue: 2.4E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.245 Pvalue: 1.9E4

		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 2.944 Pvalue: 0.005
RENT1	2	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.156 Pvalue: 2.6E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.086 Pvalue: 8.3E4
RBM4	5	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 6.033 Pvalue: 7.1E7
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 5.288 Pvalue: 1.8E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 4.933 Pvalue: 1.9E5

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 5.934 Pvalue: 3.1E4
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 4.368 Pvalue: 0.001
DAZAP1	1	
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 2.879 Pvalue: 0.006
MRPL12	4	
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 5.248 Pvalue: 2.7E6
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 4.891 Pvalue: 4.6E6

		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 5.477 Pvalue: 1.2E5
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		Study: Powell_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: NonMalignant Lung (11) Class 2: Lung Adenocarcinoma (11) Ttest: 3.15 Pvalue: 0.005
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SIAHBP1	6	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.903 Pvalue: 1.1E6

		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 5.083 Pvalue: 1.4E5
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		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.992 Pvalue: 2.3E5
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		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.974 Pvalue: 2E4
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		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 4.729 Pvalue: 2.1E4
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		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 3.861 Pvalue: 0.009
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AKAP1	7	
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.933 Pvalue: 2.2E4

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.348 Pvalue: 4.7E4
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		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 4.021 Pvalue: 0.001
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		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Small Cell Lung Cancer (4) Ttest: 5.152 Pvalue: 0.002
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		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.366 Pvalue: 0.005
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		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.29 Pvalue: 0.005
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		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 2.803 Pvalue: 0.008
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CPSF6	5	
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 4.827 Pvalue: 7.7E4

		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 5.028 Pvalue: 0.001
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		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.09 Pvalue: 0.003
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		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Small Cell Lung Cancer (4) Ttest: 4.71 Pvalue: 0.004
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		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 3.291 Pvalue: 0.004
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LA	5	
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		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.377 Pvalue: 8.2E6
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 5.235 Pvalue: 2.6E5
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 4.463 Pvalue: 1.6E4
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 3.788 Pvalue: 0.002
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 2.956 Pvalue: 0.006
NCBP2	10	

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 6.508 Pvalue: 1.9E6
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 5.589 Pvalue: 4.2E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 5.489 Pvalue: 6.1E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 5.314 Pvalue: 6.6E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 4.65 Pvalue: 8.3E5
		Study: Yamagata_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (3) Class 2: Squamous Cell Lung Carcinoma (11) Ttest: 3.553 Pvalue: 0.004
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 2.993 Pvalue: 0.004

		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.683 Pvalue: 0.004
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 4.563 Pvalue: 0.008
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 2.942 Pvalue: 0.009
DSRAD	3	
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 5.976 Pvalue: 3E5
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 3.752 Pvalue: 7.5E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.088 Pvalue: 0.005
EIF3S4	1	

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.27 Pvalue: 0.003
GARS	8	
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 8.578 Pvalue: 5.5E10

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 7.415 Pvalue: 7.9E8
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 6.521 Pvalue: 2.3E7
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 6.767 Pvalue: 2.5E7
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 5.262 Pvalue: 2.4E6

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 4.517 Pvalue: 6.6E5
		Study: Powell_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: NonMalignant Lung (11) Class 2: Lung Adenocarcinoma (11) Ttest: 3.469 Pvalue: 0.003
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.063 Pvalue: 0.003
FUBP1	6	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 7.214 Pvalue: 1.4E7
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 5.456 Pvalue: 9.8E6
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.26 Pvalue: 3.1E4

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.689 Pvalue: 3.1E4
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 4.462 Pvalue: 4.1E4
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.737 Pvalue: 0.007
FBL	4	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 7.149 Pvalue: 9.9E9
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 5.525 Pvalue: 4.8E6
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.117 Pvalue: 1.4E5

		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 3.948 Pvalue: 5.9E4
FUBP2	4	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 4.107 Pvalue: 2.4E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.811 Pvalue: 6.9E4
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Small Cell Lung Cancer (4) Ttest: 4.837 Pvalue: 0.003

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.576 Pvalue: 0.004
HTF9C	2	
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.346 Pvalue: 0.001

		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Large Cell Lung Cancer (4) Ttest: 4.436 Pvalue: 0.007
FUS	5	
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.543 Pvalue: 3.3E6
		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 4.279 Pvalue: 1.6E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 4.244 Pvalue: 1.9E4
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.302 Pvalue: 0.002
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 3.705 Pvalue: 0.002
GRSF1	4	

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.413 Pvalue: 1.7E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.445 Pvalue: 0.003
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 3.365 Pvalue: 0.003
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 3.553 Pvalue: 0.009
STRBP	2	
		Study: Yamagata_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (3) Class 2: Squamous Cell Lung Carcinoma (11) Ttest: 4.867 Pvalue: 3.9E4
		Study: Yamagata_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (3) Class 2: Lung Adenocarcinoma (9) Ttest: 3.806 Pvalue: 0.003
PPIE	5	

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 5.245 Pvalue: 1.4E5
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 4.724 Pvalue: 1.5E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 3.897 Pvalue: 0.001
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.627 Pvalue: 0.002
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 4.763 Pvalue: 0.006
ANKRD17	3	

		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 7.269 Pvalue: 4.3E7
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Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 3.99 Pvalue: 9.2E4 Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 3.231 Pvalue: 0.007

Name	# of Studies	Study Details
PAPOLG	1	
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 3.225 Pvalue: 0.002
CDKN2AIP	1	
		Study: Yamagata_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (3) Class 2: Squamous Cell Lung Carcinoma (11) Ttest: 3.535 Pvalue: 0.004
CUGBP2	6	
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 9.412 Pvalue: 1.3E13
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 14.48 Pvalue: 5.1E7
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 4.917 Pvalue: 9.4E6

		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 6.302 Pvalue: 1E5
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 3.629 Pvalue: 0.002
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 3.401 Pvalue: 0.002
RNH1	1	
		Study: Yu_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (23) Class 2: Prostate Carcinoma (64) Ttest: 4.45 Pvalue: 6.5E5
RBM25	1	
		Study: Beer_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (10) Class 2: Lung Adenocarcinoma (86) Ttest: 3.117 Pvalue: 0.005
TACC1	11	
		Study: Su_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (31) Class 2: Lung Adenocarcinoma (31) Ttest: 9.908 Pvalue: 8E13

		Study: Stearman_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (19) Class 2: Lung Adenocarcinoma (20) Ttest: 10.271 Pvalue: 1.7E10
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Squamous Cell Lung Carcinoma (21) Ttest: 8.606 Pvalue: 4.2E10
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Lung Adenocarcinoma (139) Ttest: 7.729 Pvalue: 1.3E7
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Lung Adenocarcinoma (40) Ttest: 6.378 Pvalue: 6.1E7
		Study: Wachi_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (5) Class 2: Squamous Cell Lung Carcinoma (5) Ttest: 8.85 Pvalue: 8.9E5

		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Squamous Cell Lung Carcinoma (13) Ttest: 4.604 Pvalue: 3.5E4
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Small Cell Lung Cancer (6) Ttest: 4.54 Pvalue: 0.002

		Study: Yamagata_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (3) Class 2: Lung Adenocarcinoma (9) Ttest: 3.582 Pvalue: 0.007
		Study: Garber_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (6) Class 2: Small Cell Lung Cancer (4) Ttest: 4.267 Pvalue: 0.009
		Study: Bhattacharjee_Lung Experiment Type: mRNA Analysis: Lung Type Class 1: Normal Lung (17) Class 2: Carcinoid (20) Ttest: 2.741 Pvalue: 0.01

Name	# of Studies	Study Details
ANKRD17	3	
		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 5.72 Pvalue: 1.5E7
		Study: Vanaja_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (8) Class 2: Prostate Adenocarcinoma (27) Ttest: 4.321 Pvalue: 0.002
		Study: Dhanasekaran_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (6), Benign Prostatic Hyperplasia (16) Class 2: Primary Prostate Carcinoma (59) Ttest: 3.064 Pvalue: 0.004
DICER	1	
		Study: Tomlins_Prostate Experiment Type: mRNA Analysis: Prostate Epithelium Type Class 1: Benign Prostate (22) Class 2: Prostate Carcinoma (30) Ttest: 5.412 Pvalue: 2.3E6
SNRPA	2	
		Study: Dhanasekaran_Prostate_2 Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Adjacent Prostate (2), Normal Adult Prostate (7), Normal Pubertal Prostate (3) Class 2: Prostate Carcinoma (25) Ttest: 6.255 Pvalue: 5.4E7
		Study: Dhanasekaran_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (6), Benign Prostatic Hyperplasia (16) Class 2: Primary Prostate Carcinoma (59) Ttest: 3.846 Pvalue: 4E4
CSTF3	4	

		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 3.582 Pvalue: 5.3E4
		Study: Yu_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (23) Class 2: Prostate Carcinoma (64) Ttest: 3.285 Pvalue: 0.002
		Study: Dhanasekaran_Prostate_2 Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Adjacent Prostate (2), Normal Adult Prostate (7), Normal Pubertal Prostate (3) Class 2: Prostate Carcinoma (25) Ttest: 3.529 Pvalue: 0.002
		Study: Tomlins_Prostate Experiment Type: mRNA Analysis: Prostate Epithelium Type Class 1: Benign Prostate (22) Class 2: Prostate Carcinoma (30) Ttest: 3.146 Pvalue: 0.003
NOL8	2	
		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 3.988 Pvalue: 1.5E4
		Study: Dhanasekaran_Prostate_2 Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Adjacent Prostate (2), Normal Adult Prostate (7), Normal Pubertal Prostate (3) Class 2: Prostate Carcinoma (25) Ttest: 3.13 Pvalue: 0.004
C20orf119	1	

		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 3.28 Pvalue: 0.001
STRBP	2	
		Study: Vanaja_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (8) Class 2: Prostate Adenocarcinoma (27) Ttest: 5.67 Pvalue: 1E5
		Study: Varambally_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Benign Prostate (6) Class 2: Prostate Carcinoma (7) Ttest: 3.531 Pvalue: 0.005
CSTF2	3	
		Study: Dhanasekaran_Prostate_2 Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Adjacent Prostate (2), Normal Adult Prostate (7), Normal Pubertal Prostate (3) Class 2: Prostate Carcinoma (25) Ttest: 8.195 Pvalue: 1.3E8
		Study: Tomlins_Prostate Experiment Type: mRNA Analysis: Prostate Epithelium Type Class 1: Benign Prostate (22) Class 2: Prostate Carcinoma (30) Ttest: 3.197 Pvalue: 0.002
		Study: Dhanasekaran_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (6), Benign Prostatic Hyperplasia (16) Class 2: Primary Prostate Carcinoma (59) Ttest: 2.991 Pvalue: 0.006
RBM35A	4	

		Study: Vanaja_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (8) Class 2: Prostate Adenocarcinoma (27) Ttest: 9.183 Pvalue: 9E10
		Study: Dhanasekaran_Prostate_2 Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Adjacent Prostate (2), Normal Adult Prostate (7), Normal Pubertal Prostate (3) Class 2: Prostate Carcinoma (25) Ttest: 6.096 Pvalue: 1.9E6
		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 4.95 Pvalue: 3.4E6
		Study: Liu_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Adjacent Normal Prostate (13) Class 2: Prostate Carcinoma (44) Ttest: 3.136 Pvalue: 0.004
TUT1	1	
		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 2.926 Pvalue: 0.004
SYNJ2	4	
		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 5.758 Pvalue: 9.6E8

		Study: Vanaja_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (8) Class 2: Prostate Adenocarcinoma (27) Ttest: 5.338 Pvalue: 4.3E5
		Study: Welsh_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (9) Class 2: Prostate Carcinoma (25) Ttest: 4.627 Pvalue: 2.4E4
		Study: Nanni_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (1), Benign Prostatic Hyperplasia (2) Class 2: Prostate Carcinoma (22) Ttest: 3.924 Pvalue: 0.002

Name	# of Studies	Study Details
TACC1	9	
		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 8.674 Pvalue: 1.2E13
		Study: Dhanasekaran_Prostate_2 Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Adjacent Prostate (2), Normal Adult Prostate (7), Normal Pubertal Prostate (3) Class 2: Prostate Carcinoma (25) Ttest: 7.941 Pvalue: 5.3E8
		Study: Yu_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (23) Class 2: Prostate Carcinoma (64) Ttest: 4.807 Pvalue: 1.6E5

		Study: Liu_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Adjacent Normal Prostate (13) Class 2: Prostate Carcinoma (44) Ttest: 4.782 Pvalue: 2.9E5
		Study: Vanaja_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (8) Class 2: Prostate Adenocarcinoma (27) Ttest: 4.901 Pvalue: 1.2E4
		Study: Dhanasekaran_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (6), Benign Prostatic Hyperplasia (16) Class 2: Primary Prostate Carcinoma (59) Ttest: 4.236 Pvalue: 1.2E4

		Study: Welsh_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (9) Class 2: Prostate Carcinoma (25) Ttest: 5.818 Pvalue: 1.4E4
		Study: Luo_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Benign Prostatic Hyperplasia (9) Class 2: Prostate Carcinoma (16) Ttest: 3.765 Pvalue: 0.001
		Study: Varambally_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Benign Prostate (6) Class 2: Prostate Carcinoma (7) Ttest: 3.711 Pvalue: 0.004
RBM38	3	
		Study: Dhanasekaran_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (6), Benign Prostatic Hyperplasia (16) Class 2: Primary Prostate Carcinoma (59) Ttest: 3.28 Pvalue: 0.002
		Study: Luo_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Benign Prostatic Hyperplasia (9) Class 2: Prostate Carcinoma (16) Ttest: 3.355 Pvalue: 0.003

		Study: Tomlins_Prostate Experiment Type: mRNA Analysis: Prostate Epithelium Type Class 1: Benign Prostate (22) Class 2: Prostate Carcinoma (30) Ttest: 2.81 Pvalue: 0.007
IGF2BP3	2	
		Study: Varambally_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Benign Prostate (6) Class 2: Prostate Carcinoma (7) Ttest: 4.973 Pvalue: 5.7E4
		Study: Tomlins_Prostate Experiment Type: mRNA Analysis: Prostate Epithelium Type Class 1: Benign Prostate (22) Class 2: Prostate Carcinoma (30) Ttest: 3.144 Pvalue: 0.003
IREB1	4	
		Study: Yu_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (23) Class 2: Prostate Carcinoma (64) Ttest: 4.784 Pvalue: 1.6E5
		Study: Singh_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (50) Class 2: Prostate Carcinoma (52) Ttest: 3.371 Pvalue: 0.001
		Study: Dhanasekaran_Prostate_2 Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Adjacent Prostate (2), Normal Adult Prostate (7), Normal Pubertal Prostate (3) Class 2: Prostate Carcinoma (25) Ttest: 3.479 Pvalue: 0.003

		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 2.87 Pvalue: 0.005
ACF	1	
		Study: Lapointe_Prostate Experiment Type: mRNA Analysis: Prostate Type Class 1: Normal Prostate (41) Class 2: Prostate Carcinoma (62) Ttest: 2.854 Pvalue: 0.006

Name	# of Studies	Study Details
IREB1	1	
		Study: Huang_Thyroid Experiment Type: mRNA Analysis: Thyroid Type Class 1: Normal Thyroid (8) Class 2: Thyroid Carcinoma (8) Ttest: 5.24 Pvalue: 1.3E4

Name	# of Studies	Study Details
RBM28	3	

		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 6.081 Pvalue: 2.6E4
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 5.228 Pvalue: 0.002
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 3.316 Pvalue: 0.004

Name	# of Studies	Study Details
U2AF2	1	
		Study: Haslinger_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal BCells (11) Class 2: Chronic Lymphocytic Leukemia (100) Ttest: 3.457 Pvalue: 0.005 Reporter: 32557_at
ROD1	3	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 6.888 Pvalue: 6.9E6
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 5.168 Pvalue: 1.8E4
		Study: Haslinger_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal BCells (11) Class 2: Chronic Lymphocytic Leukemia (100) Ttest: 3.222 Pvalue: 0.007
CPEB2	3	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 10.568 Pvalue: 7.4E9

		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 5.906 Pvalue: 3.2E6
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 6.123 Pvalue: 6.5E5
SAFB2	2	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 4.503 Pvalue: 1.3E4
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 3.321 Pvalue: 0.003
SRP14	1	
		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 2.913 Pvalue: 0.008
SR140	2	
		Study: Durig_Leukemia Experiment Type: mRNA Analysis: CD3+ Purified Peripheral Blood Cells Type Class 1: Normal (8) Class 2: inv(14)(q11q32)Positive TCell Prolymphocytic Leukemia (5) Ttest: 4.776 Pvalue: 5.8E4

		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 2.915 Pvalue: 0.009 Reporter: 212058_at
SNRP70	3	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 5.524 Pvalue: 2.4E5
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 6.759 Pvalue: 3.7E5

		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 4.508 Pvalue: 4.4E4
SFRS12	1	
		Study: Haslinger_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal BCells (11) Class 2: Chronic Lymphocytic Leukemia (100) Ttest: 3.517 Pvalue: 0.004
TACC1	4	

		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 9.063 Pvalue: 4E12
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 9.8 Pvalue: 8.4E8
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 7.154 Pvalue: 1.1E7
		Study: Haslinger_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal BCells (11) Class 2: Chronic Lymphocytic Leukemia (100) Ttest: 7.349 Pvalue: 4.3E6
SLTM	2	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 4.222 Pvalue: 3.6E4
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 4.117 Pvalue: 0.002

RBMS1	3	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 11.291 Pvalue: 1.6E10
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 6.017 Pvalue: 3.3E6
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 6.986 Pvalue: 4.7E6
MINT	2	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 5.02 Pvalue: 3.6E6
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 5.013 Pvalue: 3.5E5
LKAP	3	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 5.181 Pvalue: 7.6E4

		Study: Haslinger_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal BCells (11) Class 2: Chronic Lymphocytic Leukemia (100) Ttest: 4.024 Pvalue: 8.1E4
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 4.653 Pvalue: 0.001
NOL8	2	
		Study: Durig_Leukemia Experiment Type: mRNA Analysis: CD3+ Purified Peripheral Blood Cells Type Class 1: Normal (8) Class 2: inv(14) (q11q32)Positive TCell Prolymphocytic Leukemia (5) Ttest: 3.922 Pvalue: 0.003
		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 3.323 Pvalue: 0.004
MTHFSD	1	

		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 3.907 Pvalue: 0.002
FMR1	2	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 6.49 Pvalue: 5.6E5

		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 3.165 Pvalue: 0.007
ERAL1	3	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 10.715 Pvalue: 2.4E9
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 6.964 Pvalue: 8E6
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 4.974 Pvalue: 3.7E5
LARP7	1	
		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 7.237 Pvalue: 1.5E7
KRR1	4	

		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 12.623 Pvalue: 1E13
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 6.694 Pvalue: 2.1E5
		Study: Haslinger_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal BCells (11) Class 2: Chronic Lymphocytic Leukemia (100) Ttest: 5.036 Pvalue: 1.2E4
		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 3.839 Pvalue: 0.002
RBM25	1	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 4.068 Pvalue: 0.002
RBM23	1	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 3.26 Pvalue: 0.005
RBM44	1	

		Study: Maser_Leukemia Experiment Type: DNA Analysis: Leukemia Type Class 0: TCell Acute Lymphoblastic Leukemia (18) One Sample Ttest: 11.511 Pvalue: 1.9E9
RBM26	1	
		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 3.891 Pvalue: 7.1E4
ACIN1	4	
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: BCell Acute Lymphoblastic Leukemia (87) Ttest: 7.86 Pvalue: 8.5E9
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 6.224 Pvalue: 1.2E6
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: TCell Acute Lymphoblastic Leukemia (11) Ttest: 7.818 Pvalue: 1.4E6

		Study: Durig_Leukemia Experiment Type: mRNA Analysis: CD3+ Purified Peripheral Blood Cells Type Class 1: Normal (8) Class 2: inv(14) (q11q32)Positive TCell Prolymphocytic Leukemia (5) Ttest: 4.235 Pvalue: 0.003
DND1	1	

		Study: Durig_Leukemia Experiment Type: mRNA Analysis: CD3+ Purified Peripheral Blood Cells Type Class 1: Normal (8) Class 2: inv(14) (q11q32)Positive TCell Prolymphocytic Leukemia (5) Ttest: 3.549 Pvalue: 0.005
RBM16	2	
		Study: Haslinger_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal BCells (11) Class 2: Chronic Lymphocytic Leukemia (100) Ttest: 5.036 Pvalue: 2.3E4
		Study: Andersson_Leukemia Experiment Type: mRNA Analysis: Leukemia Type Class 1: Normal Bone Marrow (6) Class 2: Acute Myeloid Leukemia (23) Ttest: 3.357 Pvalue: 0.005
PSPC1	1	
		Study: Maia_Leukemia Experiment Type: mRNA Analysis: BCell Type Class 1: Normal Pre BCell (CD19+, 2lg) (5), Normal Immature BCell (CD19+, slg+) (3) Class 2: Acute Lymphoblastic Leukemia (TEL/AML1 Genotype) (20) Ttest: 3.928 Pvalue: 0.001

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Supplementary file 14

Biological Process	Gene	PubMed ID
Apoptosis	ACO1	11856741 , 15169927
	ADAR	14613934 , 15677478 , 15746098 , 16126896
	AKAP1	11483602 , 12427737
	APOBEC1	16055734
	NKRF	10777530 , 12533512
	NOL8	15132771
	NPM1	7721868 , 9881708 , 10233871 , 10362140 , 10391679 , 10597193 , 11054668 , 11110708 , 11125022 , 11280786 , 11522649 , 11704868 , 11756229 , 12067985 , 12244208 , 12374805 , 12691918 , 12748172 , 12748950 , 12882984 , 14508522 , 14583466 , 14656879 , 14962911 , 15087454 , 15126367 , 15297316 , 15310764 , 15374880 , 15459011 , 15514966 , 15831697 , 15883821 , 15886198 , 15893727 , 15928040 , 15964625 , 16046528 , 16076867 , 16103750 , 16105984 , 16135792 , 16166282 , 16170336 , 16199867 , 16210342 , 16304056 , 16317043 , 16455956 , 16469875 , 16540653 , 16597596 , 16709933 , 16735125 , 16763206 , 16818631 , 16827655 , 16857742 , 16880530 , 16982741 , 17072349 , 17073594 , 17111047 , 17185414 , 17277230 , 17483340 , 17611412 , 17612934 , 18214681 , 18288131 , 18345031 , 18420587 , 18574026 , 18974134
	PABPC4	11352910
	PNPT1	16410805 , 17804700
	PTBP1	16885029
	PUF60	16452196
	RBM10	15514923
	RBM25	18663000
	RPL7	9566973 , 10672495 , 10859237
	RPS5	10821535

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	SSB	9794444 , 12102760 , 16906225
	TACC1	16288026
	TIA1	7964902 , 8759725 , 9486710 , 9664088 , 9973508 , 10404082 , 10613902 , 10672058 , 11514562 , 12359741 , 12533540 , 14736875 , 15316071 , 15572676 , 17869012 , 18756326 , 18959781
	TROVE2	16301677 , 16906225
Cell Cycle	ADAR	17224799
	AKAP1	10995432 , 11994283 , 12697839 , 12925532 , 15470020
	CSTF2	9736695 , 10859161 , 11598190 , 11943782 , 12446762 , 16371507
	CSTF3	16882666
	EIF3B	8150258 , 9362495 , 11042177 , 11134033
	FUS	12040005
	GARS	16763051
	MARS	9843569 , 15073333 , 18761010
	NONO	2336466 , 16148043
	NPM1	1874752 , 2026181 , 3308448 , 6345184 , 8526928 , 8694782 , 9181305 , 9350304 , 9914158 , 10208875 , 10233871 , 10471330 , 10503877 , 10512052 , 10931858 , 11278991 , 11812799 , 11888936 , 12058066 , 12237316 , 12464630 , 12861014 , 14583466 , 14635188 , 14636574 , 14739278 , 15082766 , 15126367 , 15184379 , 15190079 , 15265990 , 15314174 , 15388344 , 15644315 , 15665273 , 15684379 , 15699019 , 15814843 , 15872011 , 16024624 , 16140928 , 16199867 , 16501600 , 16540521 , 16540653 , 16608843 , 16648475 , 16738315 , 16818631 , 16912078 , 17069796 , 17111047 , 17113241 , 17179202 , 17215513 , 18165222 , 18729828 , 18763323 , 18931307 , 18951898
	PNPT1	16055741 , 16410805 , 17804700
	PSMA1	9247291
	PTBP1	15685555

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	RBM14	18829545
	RPL7	10672495 , 10859237
	RPS5	18288641
	SLBP	10825184 , 11157774 , 11238936 , 11884612 , 12370293 , 12415002 , 12588979 , 14999087 , 15684398 , 15829567 , 15916543 , 16055702 , 16492733 , 16714279 , 16914750 , 17442888
	SSB	1285896 , 2579168
	TACC1	10637228 , 17359303
Cell Proliferation	ACO1	11082186
	ADAR	11099415 , 14615479
	APOBEC1	9633945
	CPSF6	18636107
	CSTF2	9736695
	EIF3B	3308493
	NKRF	16648473
	NPM1	1697568 , 3392030 , 7474921 , 9819383 , 10233871 , 10878901 , 11125022 , 11280786 , 11309377 , 11494142 , 11522649 , 11756229 , 11888936 , 12057857 , 12067985 , 12086925 , 12374805 , 12623937 , 12691918 , 12748172 , 12882984 , 14635188 , 14729947 , 15082766 , 15289458 , 15485902 , 15588951 , 15616222 , 15644315 , 15665273 , 15684379 , 15805275 , 15831697 , 15872011 , 16024624 , 16045751 , 16051734 , 16135794 , 16199867 , 16254137 , 16267006 , 16317043 , 16357222 , 16469875 , 16508011 , 16540521 , 16540653 , 16648475 , 16705183 , 16709933 , 16782897 , 16785248 , 16857742 , 16870143 , 16912078 , 16982741 , 17015463 , 17069796 , 17108330 , 17113241 , 17179202 , 17277230 , 17308101 , 17612934 , 17949752 , 17998938 , 18024471 , 18070884 , 18574026 , 18809582 , 18931307 , 18951898 , 19033198
	PTBP1	12509450
	RBM14	18635665

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	RBMX	16707624
	SLBP	16914750
	SSB	1592421
	TACC1	15705873
Cell Differentiation	ACO1	8636154 , 10978328 , 11082186
	ADAR	9202292 , 11099415
	APOBEC1	160100694
	CSTF2	10781073
	EIF3G	17990289
	FUS	10442642 , 10828883
	GARS	11226317 , 16763051
	KHSRP	14657238 , 16364914 , 16914741
	NONO	9360842 , 10364157 , 10605820
	NPM1	1568462 , 9484783 , 10233871 , 11054668 , 11125022 , 11756229 , 12374805 , 12882984 , 14520467 , 14583466 , 15310764 , 15899793 , 16109776 , 16199867 , 16365184 , 16449640 , 16455956 , 16540685 , 16608843 , 16766651 , 16857742 , 17318229 , 17448503 , 17949752
	PNPT1	15492272 , 15978720 , 16733069
	PTBP1	11724819
	RBM14	17337438
	RPS5	10821535 , 11684664 , 18288641
SNRPA	15226420	
Cell Survival	ACO1	15178542
	ADAR	14613934 , 14615479
	AKAP1	12427737 , 12716901 , 16251349 , 18323779

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	NONO	18219107
	NPM1	10552961 , 12882984 , 14962911 , 15226403 , 15310764 , 15374880 , 15680399 , 15684379 , 15831697 , 15883821 , 15994285 , 16046528 , 16105984 , 16109776 , 16199867 , 16317043 , 16608843 , 16763206 , 16785248 , 16982741 , 17015463 , 17111047 , 18070884 , 18342811 , 18574026 , 18931307
	TACC1	16288026
	TIA1	12533540 , 14769925 , 15572676 , 16236613
	TROVE2	16890888 , 16906225
Cell Growth	ADAR	17156436
	CSTF2	9710585 , 9885564 , 10669729
	FXR1	17382873
	NCBP2	10733586
	NOL8	14660641
	NPM1	1381705 , 1399149 , 1697505 , 1917161 , 8601891 , 10233871 , 10503877 , 10779644 , 11756229 , 12086925 , 12374805 , 15665273 , 15684379 , 15805275 , 15872011 , 16317043 , 16648475 , 16857742 , 16982741 , 17245342 , 17318229 , 17483340 , 18024471 , 18809582
	PNPT1	15734991 , 17804700
	RBM8A	17545126
	RPL18	9891046
	STRBP	11438536