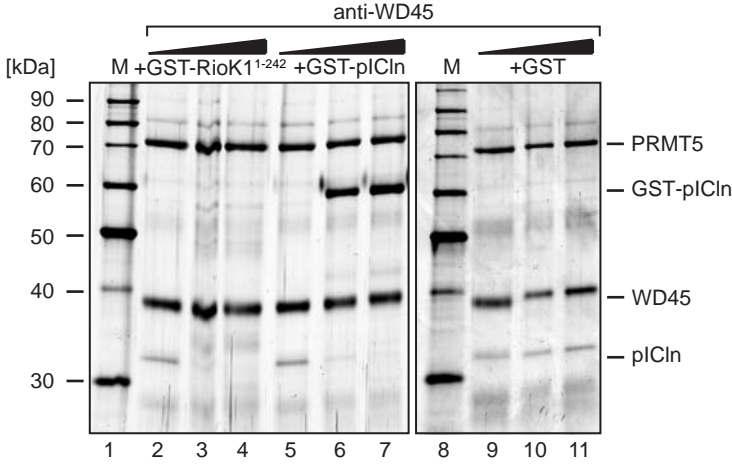


SUPPLEMENTAL FIGURE LEGEND

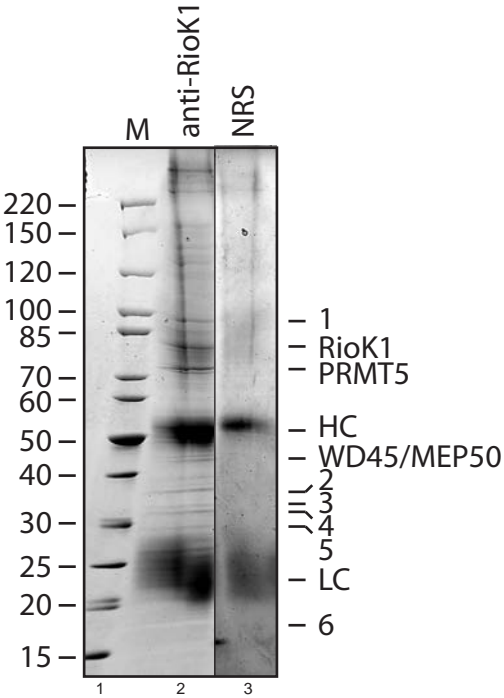
SUPPLEMENTAL FIGURE S1. **Competition assay of the PRMT5 complex with GST-RioK1, GST-pICln and GST.** The PRMT5 complex was immunoprecipitated with anti-WD45/MEP50 antibodies from HeLa cell extract. The purified complex was incubated with increasing amounts (0.1 to 10 μ g) of GST-tagged RioK1 (AA1-242, lanes 2-4), pICln (lanes 5-7) or GST (lanes 9-11). The complex composition was afterwards analyzed by silver staining.

SUPPLEMENTAL FIGURE S2. **Identification of RioK1-interacting proteins.** Immunoprecipitation from HeLa extract with anti-RioK1 antibody (lane 2) or NRS (lane 3). The immunoprecipitates were washed extensively, resolved by SDS-PAGE, Coomassie stained and the marked bands were analyzed by mass spectrometry.

Supplemental Figure S1



Supplemental Figure S2



SUPPLEMENTAL TABLE LEGEND

SUPPLEMENTAL TABLE S1. **List of putative RioK1-interacting proteins.** Proteins identified in Fig. S1 and from a pull-down in HeLa extract with GST-tagged truncated RioK1 (AA 450-568), covalently crosslinked to GSH-sepharose are listed alphabetically. Putative methylated RG-motives are in bold letters. Information about DNA-and RNA-binding capacity of proteins was derived from UniProtKB/Swiss-Prot database. Corresponding UniProtKB/Swiss protein accounts are listed in the last column.

protein	short	sequence-ID	DNA-binding	RNA-binding	putative RG-motive
116 kDa U5 small nuclear ribonucleoprotein component	U5S1	Q15029	-	-	FTEQERGVGIKS, TITEPRGNEEAQ, QEPLHRGGQII, VLARRRGHVTQD
26S protease regulatory subunit 4	PRS4	P62191	-	-	KVDDLRTGTPMSV, DGFDSRGGDVKVI
26S proteasome non-ATPase regulatory subunit 3	PSMD3	Q43242	-	-	GSARRRGADKAK
28S ribosomal protein S7, mitochondrial precursor	RT07	Q9Y2R9	-	+	AVKVARGWISGLA
28S ribosomal protein S9, mitochondrial precursor	RT09	P82933	-	-	LLLWGRGSLARK
3' histone mRNA exonuclease 1	THEX1	Q8IV48	-	+	FKLETRGVKVDVL
39S ribosomal protein S4, mitochondrial precursor	RM54	Q6P161	-	-	-
39S ribosomal protein L1, mitochondrial precursor	RM01	Q9BYD6	-	+	-
39S ribosomal protein L11, mitochondrial precursor	RM11	Q9Y3B7	-	-	LGRAARGLRKPKE, PVLGQRGVSINQ
39S ribosomal protein L19, mitochondrial precursor	RM19	P49406	-	-	EFIPRRGRDTPD, IQRSGRGLGATF
39S ribosomal protein L22, mitochondrial precursor	RM22	Q9NWU5	-	-	HNLRSRGLALG, LAKLIRGMSIDQ, ESTSGRGQCLKR, IRYHGRGRFGIM
39S ribosomal protein L38, mitochondrial precursor	RM38	Q96DV4	-	-	ECRRWRGFSTSA, PFPARGSGIHR
3-ketoacyl-CoA thiolase, mitochondrial	THIM	P42765	-	-	MALLRGVVFVA, HELRRRGKGYAV
40S ribosomal protein S10	RS10	P46783	-	-	QSLKSRGYVKEQ, PARLTRGEADRD, TEFQFRGGFGRGRGQPP
40S ribosomal protein S11	RS11	P62280	-	+	GNVSIRGRILSG
40S ribosomal protein S13	RS13	P62277	-	+	-
40S ribosomal protein S14	RS14	P62263	-	+	RKGGRRGR
40S ribosomal protein S15	RS15	P62841	+	-	RKFTYRGVDLDQ, RRRLNRGLRRKQ
40S ribosomal protein S15a	RS15A	P62244	-	+	NNAEKRGKRQVL
40S ribosomal protein S16	RS16	P62249	-	+	VAHCKRGNGLIK
40S ribosomal protein S17	RS17	P08708	-	-	MKRIQRGPPVRSIKL, NFKTPRGP
40S ribosomal protein S18	RS18	P62269	-	+	KIRAHRLRHFV, WGLRVRGQHTKT, KTTGRRGRTVGV
40S ribosomal protein S19	RS19	P39019	-	+	RHLYLRGGAGVG, PSFHSRGSKVA
40S ribosomal protein S2	RS2	P15880	-	+	PGMGNRGGFRGGFGSG, FGSGIRGRGRGRGRGRGRGARGKAED, VATAIRGAILA, IVPVRRGYWGNK, LIPAPRGTGIVS, CYTSARGCTATL
40S ribosomal protein S20	RS20	P60866	-	+	CADLIRGAKEKN
40S ribosomal protein S23	RS23	P62266	-	-	MGKCRGLRTAR
40S ribosomal protein S25	RS25	P62851	-	+	ERLKIRGLARA
40S ribosomal protein S27a	RS27A	P62979	-	-	-
40S ribosomal protein S3	RS3	P23396	+	+	EKVATRGLCAIA, VSGKLRGQRAKS
40S ribosomal protein S3a	RS3A	P61247	-	+	-
40S ribosomal protein S4, X isoform	RS4X	P62701	-	+	MARGPKKHL, WISLPRGKIGRL
40S ribosomal protein S5	RS5	P46782	-	+	-
40S ribosomal protein S6	RS6	P62753	-	-	KRKSVRGCIVDA
40S ribosomal protein S7	RS7	P62081	-	+	-
40S ribosomal protein S8	RS8	P62241	-	-	HTVRVRRGNKKY
40S ribosomal protein S9	RS9	P46781	-	+	-
40S ribosomal protein SA	RSSA	P08865	-	-	EVLRMRTISRE
5'-3' exonuclease 1	XRN1	Q8IZH2	+	+	KMNQQRGRRFRS, FQSSRGENMML, SIFIGRGSRRNP, VPVGLRGTIIGI, RCSPGRGYRLPT
60 kDa heat shock protein, mitochondrial precursor	CH60	P10809	-	-	PVEIRRGVMLAV; GMKFDRGYISPY
60S acidic ribosomal protein P0	RLA0	P05388	-	+	IRMSLRGKAVVL, MRKAIRGHLENN, LLPHIRGNVGFV, TTKISRGTIEIL
60S acidic ribosomal protein P2	RLA2	P05387	-	+	-
60S ribosomal protein L10	RL10	P27635	-	-	KSRFCRGPVDAK, LQTGMRGAFGKP, KYIPNRGPLDKW
60S ribosomal protein L10a	RL10A	P62906	-	+	-
60S ribosomal protein L10-like	RL10L	Q96L21	-	-	KSRFCRGPVDAK, LQTGMRGAFGKP
60S ribosomal protein L11	RL11	P62913	-	+	VHCTVRGAKAEE
60S ribosomal protein L12	RL12	P30050	-	+	-
60S ribosomal protein L13	RL13	P26373	-	+	KVRAGRGSLEE
60S ribosomal protein L13a	RL13A	P40429	-	-	LVLDGRGHLLGR, NTNPSRGPYHFR, FWRTVRGMLPHK, PHKTKRGQAALD
60S ribosomal protein L14	RL14	P50914	-	+	-
60S ribosomal protein L17	RL17	P18621	-	-	KSCKSRGSNLRV
60S ribosomal protein L18	RL18	Q07020	-	+	KFERARGRRASRGYK
60S ribosomal protein L18a	RL18A	Q02543	-	+	-
60S ribosomal protein L19	RL19	P84098	-	+	-
60S ribosomal protein L21	RL21	P46778	-	+	TKGKRRTRYMF
60S ribosomal protein L22	RL22	P35268	-	+	-
60S ribosomal protein L23	RL23	P62829	-	-	MSKRGRGGSSGA
60S ribosomal protein L23a	RL23A	P62750	-	+	-
60S ribosomal protein L24	RL24	P83731	-	+	-

60S ribosomal protein L24	RS24	P83731	-	+	-
60S ribosomal protein L26	RL26	P61254	-	+	EVQVVRGHYKGG
60S ribosomal protein L26-like 1	RL26L	Q9UNX3	-	-	EVQVVRGHYKGG
60S ribosomal protein L27a	RL27A	P46776	-	+	KTRKLRGHVSHG, KHPGGGRNAGGL
60S ribosomal protein L28	RL28	P46779	-	+	-
60S ribosomal protein L29	RL29	P47914	-	+	-
60S ribosomal protein L3	RL3	P39023	-	+	RSSRHRGKVKSF, YVETPRGLRFTK, PRKTHRGLRKA
60S ribosomal protein L30	RL30	P62888	-	+	-
60S ribosomal protein L31	RL31	P62899	-	+	-
60S ribosomal protein L32	RL32	P62910	-	-	NWRKPRGIDNRV
60S ribosomal protein L35	RL35	P42766	-	+	KARDLRGKKKEE
60S ribosomal protein L36	RL36	Q9Y3U8	-	-	RHSRRRGLRCLKH
60S ribosomal protein L36a	RL36A	P83881	-	-	-
60S ribosomal protein L36a-like	RL36L	Q969Q0	-	-	-
60S ribosomal protein L38	RL38	P63173	-	+	-
60S ribosomal protein L4	RL4	P36578	-	+	RIPRVRRGGGTHR, FGNMCRGGRMFA, RRIQRGPCIY
60S ribosomal protein L5	RL5	P46777	-	-	-
60S ribosomal protein L6	RL6	Q02878	+	+	NPVLRVIGRYS, LTGRHRGKRVVF
60S ribosomal protein L7	RL7	P18124	+	+	FVIRIRGINGVS, ELIYKRGYKIN, KLSSPRGGMKKK
60S ribosomal protein L7a	RL7A	P62424	-	+	-
60S ribosomal protein L8	RL8	P62917	-	+	MGRVIRGQRKGA, IHDPRGAPLAK, EKPGDRGKLARA, CWPRVRGVAMNP, RTGRLRGTKTVQ
6-phosphofructokinase type C	K6PP	Q01813	-	-	CNLLQRGITNLC, LGHVQRGGTPSA, DAVRLRGRSFAG, KTTIQRGLVLRN, KLKEARGRGKFTT
78 kDa glucose-regulated protein precursor	GRP78	P11021	-	-	GKEPSRGINPDE, IPPAPRGVPQIE
Actin, cytoplasmic 1	ACTB	P60709	-	-	EAQSKRGILTLL, KILTERGYSFTT
Actin, cytoplasmic 2	ACTG	P63261	-	-	EAQSKRGILTLL, KILTERGYSFTT
Activator of basal transcription 1	ABT1	Q9ULW3	+	+	LQSVERGQFLA
Alpha-centractin	ACTZ	P61163	-	-	KAEHRGLLSIR
Alpha-enolase	ENOA	P06733	+	-	EIFDSRGNPTVE
AP-1 complex subunit mu-1	AP1M1	Q9BXS5	-	-	CRNYRGDVDMS, FDNTGRGKSKS
AP-2 complex subunit mu-1	AP2M1	Q96CW1	-	-	-
Arginyl-tRNA synthetase, cytoplasmic	SYRC	P54136	-	-	VSLIERGESFYQ, KEFEDRGFVQVD, KMLDDRGNNTAA
Aspartyl-tRNA synthetase, cytoplasmic	SYDC	P14868	+	+	YDMFMRGEEILS
ATP synthase subunit alpha, mitochondrial precursor	ATPA	P25705	-	-	SLVPIGRGQRELI, QQLSRGVRLTE, IYAGVRGYLDKL
ATP-binding cassette sub-family F member 1	ABCF1	Q8NE71	-	+	PAPKPRGGKTK; TEYLQRGFNLPI
ATP-citrate synthase	ACYL	P53396	-	-	QLIKRRGKLG, TIFVRRGGPNYQ, EYKICRGIKEGR, SICDERGQELIY
ATP-dependent RNA helicase A	DHX9	Q08211	-	+	GPTWDRGANLKD, SVVIIRGATGCG, RVAFERGEEPGK, LEAGIRGISHVI, GSGYRRGGSSYS, VGGYRGVSRGGFRGNSGGD, SGGDYRGPSSGGYRSGGFQRRGGGRGAYGTG, YFGQGRGGGG
ATP-dependent RNA helicase DDX1	DDX1	Q92499	-	+	MNPYDRGSFAFI, TDVAARGIDIHG, HVCSSRGKGCYN
ATP-dependent RNA helicase DDX3X	DDX3X	O00571	-	+	SRSDSRGKSSFF, SFFSDRGSGRGRFDDRGRSDYD, DGISSRGRSGF, FGKFERGGNSRW, IRDLERGCHLLV, VDMMERGKIGLD, TAVAARGLDISN, YKGSSRGRSKSS, GHGSSRFGGGGG
ATP-dependent RNA helicase DDX3Y	DDX3Y	O15523	-	+	GSRDSRGKPGYF, GYFSEGRSGRGRFDDRGRSDYD, IRDLERGCHLLV, VDMMERGKIGLD, TAVAARGLDISN, YKGSSRGRSKSN, GFGASRGSSSRS, GYGNSRFGGGGG
ATP-dependent RNA helicase DDX50	DDX50	Q9BQ39	-	+	KLLKGRGVTYLF, TNVAARGLDIPE, YQPRERGLRYV
ATP-dependent RNA helicase DDX54	DDX54	Q8TDD1	-	+	GLRKRGAASQARSDSED, TDLAARGLDIPL, ASLELRGLARVA, RSGPNRGAKRNR, DFDSEGLSISG, AQLNTRGRQQLK, GASDRRGPERRGGKRDGQASR, LHFLQRGLKQL, QGAFGRGARSKK
ATP-dependent RNA helicase DDX55	DDX55	Q8NHQ9	-	+	TDVMARGIDIPE, FASLARGFALLR, KMPELRGKQFPD
BAG family molecular chaperone regulator 1	BAG1	Q99933	-	-	MAQRGGARRPRGDRERL, EPPAQRGPPPSR, HDRPTRGAAAGA
Beta-centractin	ACTY	P42025	-	-	KAEHRGLLTI
Bifunctional aminoacyl-tRNA synthetase	SYEP	P07814	-	+	RFPTVRGLRRGMTVEG, IQLQRRGFICD
Brix domain-containing protein 2	BXDC2	Q8TDN6	-	-	TKRKRGGFAVQ, LIFSSRGINFRT
Bystin	BYST	Q13895	-	-	KFKAARGVGGQE, VREKRGRGTGEA, VLEVYRGVREVL
CAD protein	PYR1	P27708	-	-	DGSVLRGQPFGA, RCLCQRGAETV, MRYGNRGNQPC, LLEQHRGQPLPP, INLSMRGAGGRR, KAAKARGLPVT, RRVVLRGEVAYI, PERPRRGIPGLP, AFVASRGTKQEE
Carbamoyl-phosphate synthase [ammonia]	CPSM	P31327	-	-	RLLVKRGAEVHL, MSMANRGNQPV, VELFKRGVLKEY
Carbonyl reductase [NADPH] 1	CBR1	P16152	-	-	ARDVTRGQAAVQ
Carbonyl reductase [NADPH] 3	CBR3	O75828	-	-	VTGANRIGLAI, ARDVARGQAAVQ
Cell division cycle 5-like protein	CDC5L	Q99459	+	+	KRKRKRGVDYNA
Cell growth-regulating nucleolar protein	LYAR	Q9NX58	-	-	-
Chromodomain helicase-DNA-binding protein3	CHD3	Q12873	+	-	VKKRKRGPKKQK, GRKKLRGKMAP, TKKLRGRPRGR, EALNSRGGGNQV, VRDLRGKTEKE, PEPGYRGDREKS, STPGERGEKPL
Cisplatin resistance-associated overexpressed protein	CROP	Q86Y74	-	-	ERRIRRGHARLA
Clathrin heavy chain 1	CLH1	Q00610	-	-	FCFAVRGQAGGK, LILVRRGQF, CVAYERGQCDLE, NYYQDRGYFEEL
Cleavage and polyadenylation specificity factor 7	CPSF7	Q8N684	-	+	CVRVPRGGIPPR

Coatamer subunit delta	COPD	P48444	-	-	-
Coiled-coil domain-containing protein 86	CCD86	Q9H6F5	-	-	-
Cold shock domain-containing protein E1	CSDE1	O75534	+	+	FGFIERGDVVE, VLRQPRGPDNSM
Complement component 1 Q subcomponent-binding protein	C1QB	Q07021	-	-	GLLRPRGPCACG, DFLADRGVDTNF
Core histone macro-H2A.1	H2AY	O75367	+	-	MSSRGGKKKS, LLAKKRGSKGL
Crossover junction endonuclease MUS81	MUS81	Q96NY9	+	-	DIGETRGGGHRP, LALLTRGLQRLY, QLMQVRGVSGEK
Cullin-associated NEDD8-dissociated protein 1	CAND1	Q86VP6	-	-	-
D-3-phosphoglycerate dehydrogenase	SERA	O43175	-	-	VVNCARGGIVDE
Developmentally-regulated GTP-binding protein 1	DRG1	Q9Y295	-	-	KDGKGRGRQVIA
DNA mismatch repair protein MSH6	MSH6	P52701	+	-	DDSPTRGWWVSKR, LANYSRGGDGM, VDELGRGTATFD
DNA polymerase delta catalytic subunit	DPOD1	P28340	+	-	PPKRARGGLWDD, GPPPSRGSVPVL, ISRDSRGGRELT, SYLLSRGQQVKV, EAVLLRGDHTRC
DNA replication licensing factor MCM3	MCM3	P25205	+	-	NGSHIRGDINIL, IPTTGRGSSGVG, MVLADRGVVICD
DNA replication licensing factor MCM6	MCM6	Q14566	+	-	QAEPRGSIPRS, ETEGIRGLRALG, NDEVKRGVLLML, EGTSLRGDINVC
DNA-binding protein A	DBPA	P16989	+	+	RRRYRRGYGRRRGPPRNY, PRYSRGGPPRPR, QPSVRRGYRRPY
DNA-dependent protein kinase catalytic subunit	PRKDC	P78527	+	-	GHQLIRGLGQEC, LSAIRGYGLFA, KRRLPRGFPPSA, TLLYLGRPFSLQ, RRIARRGSCVTO, PAKRVRGKARLP, EYDVLRGIFTSE, GOYDGRGKPLPE, KRIIRRHDERE, YVAVARGSKDHN
DNA-directed RNA polymerase III subunit RPC2	RPC2	Q9NW08	+	-	DIEYTRGSQRII, KASLDRGFGRCL, MHARARGPRAVL
DNA-directed RNA polymerases I and III subunit RPAC1	RPAC1	O15160	+	-	-
DnaJ homolog subfamily A member 1	DNJA1	P31689	-	-	MQRERRGKNVVH, DKCEGRGKKGA, CCPNCRGTGMQI, AVFTRRGEDLFM, DEHHPRGGVQCC
DnaJ homolog subfamily C member 9	DNJC9	Q8WXX5	-	-	DGEVRRGYHKVS
Double-stranded RNA-binding protein Staufen homolog 2	STAU2	Q9NUL3	-	+	KLNERGPAHSK, GLAMKRGEPAIY, ANYNFRGMYNQR, VLLSERGMPPRR
Dynein heavy chain, cytosolic	DYHC	Q14204	-	-	KQCYERGEKPKV, VRPHIRGAIREY, ESTRVGRGTGNV, QIRKARFTDNA, KCRLIRGWDDL, REKEERGEAVDE, GVQYHRGEMTAL, IIDSVRGELQKR, MTRVWRGIFEAL, ARLAKRGGRTMA, FNRVARGMLHQD, FQHFLRGNEIVL, RSFYERGVAVLC
Elongation factor 1-alpha 1	EF1A1	P68104	-	-	KAERERGITIDI, VKDVRRGNAVGD
Elongation factor Tu, mitochondrial precursor	EFTU	P49411	-	-	LPLLCRGLAVEA, PEERARGITINA, YSVPRGTVVTG, TGTLERGILKKG, LGALVRGLKRED, REDLRRGLVMVK
ESF1 homolog	ESF1	Q9H501	-	-	YAVDKRGRPISH, GPDLARGKNIE
Eukaryotic initiation factor 4A-I	IF4A1	P60842	-	+	SELLRGIYAYG, DEMLSRGFKDOI, TDLARGIDVQQ, IHRIGRGGRRFR
Eukaryotic translation initiation factor 2 subunit 1	IF2A	P05198	-	+	KIEEKRGVFNVQ
Eukaryotic translation initiation factor 2 subunit 2	IF2B	P20042	-	+	-
Eukaryotic translation initiation factor 2 subunit 3	IF2G	P41091	-	-	WGQIRRGVTIKP
Eukaryotic translation initiation factor 2C 2	I2C2	Q9UKV8	-	-	-
Eukaryotic translation initiation factor 3 subunit 10	IF3A	Q14152	-	-	DSEWRRGPPEKE, EKEWRRGEGRDE, DRVPRGMDDDRGPRRGPEEDR, DRFSRRGADDDR, IADEDNWNRHA, DRPPRRGLDEDRGSRWTA, TADEDRGRPRGMDDDRGPRRGADDE, NADDDRGRPRGLDDDRGPRGMDDDRGPRRGDDDRGPRRGDDDRGPRRGDDDRGPRRWNA, DRIPRRAEDDRGPWRNM, DRFPRRGDDSRP, EGGWRRGPAEES, DDRDRRGPLRS
Eukaryotic translation initiation factor 3 subunit 6	IF36	P60228	-	-	-
Eukaryotic translation initiation factor 3 subunit 6-interacting protein	IF3I	Q9Y262	-	-	-
Exosome component 10	EXOSX	Q01780	-	+	LEMWERGNGQPV, TGKSDRGFRYNW
Exportin-2	XPO2	P55060	-	-	ACDLVRGLCKFF, RLFTMRGPNAT, QAFLEGSNTIA
Exportin-T	XPOT	O43592	-	+	VDLNPRGVLDYL, AFLDHRGLRHSS, TIHLKRGPECVQ
FACT complex subunit SPT16	SPT16	Q9Y5B9	-	-	EDLLGRGSRAAL, RFTSVRGDKVDI, VHSSGRGNSRGRSHSS
Fatty acid synthase	FAS	P49327	-	-	NPDSLRTHTGV, FFFDFRGPISAL, QPLPVRGGNVI, TAMPPFRGYAVLG, VLGGERRGPEVQ, MGTQWRGMGLSL, LAAYWRGQCIKE, QAVLKRGLKPSK, EFPAPRGTP LIS, KELRLRGYDYG, LHTLLRGHPLGD, VSTLTRGDLSSI, YALVVRGRVPRG, QWLIQRGVQKLV, SLEGARGLIAEA, SVSCGRGNAGQS
FilaminA	FLNA	P21333	-	-	PQLEARGDSTYR, CRAVGRGLQPKG, DRVKARGPGLK, VKYTPRGAGSYT, TYVQDRGDGMYK, FTVETRGAGTGG
G patch domain-containing protein 4	GPTC4	Q5T310	+	+	PEVKSRRGMKFAE, LNLEDRGEETVL, TGSSSRGKRKRQ
GCN1-like protein 1	GCN1L	Q92616	-	-	MLASYRGDTLLQ, RETVLRGLMELH, LQDSNRGVRLKA, AVSGVRGMGFLM
GC-rich sequence DNA-binding factor homolog	GCFC	Q9Y5B6	+	-	FLQEMRGYVQDL
General transcription factor II-I	GTF2I	P78347	+	-	VVGTERGRAFVN, LERIVRGSNKIK, VPFKPRGREFSF
Glutamate-rich WD repeat-containing protein 1	GRWD1	Q9BQ67	-	-	VYLPGRGPPLRE
Glutamyl-tRNA synthetase	SYQ	P47897	-	-	LMEQLRGEALFK, VELIRRGGLAYVC, YVCHQRGEELKG, TALRRRGFPPEA
Glutathione S-transferase Mu 3	GSTM3	P21266	-	-	GYWDIRGLAHA
Glyceroldehyde-3-phosphate dehydrogenase	G3P	P04406	-	-	LWRDGRGALQNI
G-rich sequence factor 1	GRSF1	Q12849	-	+	LGALLRGCNC, AASQTRGLQTP, RDGKRRGDALIE, GVVRLRGLPYSC, FMVDYRGRRTKG, HFVHMRGLPFQA
Guanine nucleotide-binding protein subunit beta 2-like 1	GBLP	P63244	-	-	EQMTLRGTLKGH, PQRALRGHSHFV
Guanine nucleotide-binding protein-like 3	GNL3	Q9BVP2	-	-	KEAKKRGHKKPR, VLAQRGMHQKG
H/ACA ribonucleoprotein complex subunit 4	DKC1	O60832	-	+	-
Heat shock 70 kDa protein 1	HSP71	P08107	-	-	IPPAPRGVPQIE
Heat shock 70 kDa protein 1L	HS71L	P34931	-	-	CADLFRGTLEPV, IPPAPRGVPQIE
Heat shock 70 kDa protein 6	HSP76	P17066	-	-	VRVCYRGEDKTF, YGLDRRGAGERN, IPPAPRGVPQIE
Heat shock protein HSP 90-alpha	HS90A	P07900	-	-	GEPMGRGTVIL, YLNFIRGVVDSE
Heat shock protein HSP 90-beta	HS90B	P08238	-	-	GEPIGRGTVIL, YLNFIRGVVDSE, ERVRRKGFVVY
Heterogeneous nuclear ribonucleoprotein A/B	ROAA	Q99729	-	+	VPEASRGRGWTGAA, NTRGSRGFGFIL, KLNKRGRVFVIT, YGSGGRNRRNRGRGSGGGG, GKSQRRGGHQNN

Heterogeneous nuclear ribonucleoprotein A0	ROA0	Q13151	-	+	SESLRGLGHFEAF, QSGKKRGGFVFY, GSRSSRGGRRGGRRGGRRDQ, NSGPYRGGYGGG
Heterogeneous nuclear ribonucleoprotein A1	ROA1	P09651	+	+	NTKRSRGGFVFT, EIMDRGSGKKRGFVFT, ASSSQRRSGSG, NFGGRRGGGFGG, NDNFRGGNFSGRGGFGGSRGGGGYG, YSGGSRGGYSGG
Heterogeneous nuclear ribonucleoprotein A3	ROA3	P51991	-	+	RRRRRRGEEGHD, QTKRSRGGFVFT, QSGKKRGFVFT, SAGSQRRGGGSGN, GNFMGRGGNFGG, GGNFGRGGNFGGRRGGYGGG, GGGGSRGSGYGGG, PGYSSRGGYGGG
Heterogeneous nuclear ribonucleoprotein D0	HNRPD	Q14103	+	+	ITGRSRGGFVFL, KTNKRGGFCFIT, QQWGSRGGFAGR, FAGRARGGGGPSQ, GKVSRRGGHQNS
Heterogeneous nuclear ribonucleoprotein D-like	HNRDL	O14979	+	+	VTGRSRGGFVFL, KTNERGGFCFIT, QQKGGRGAAGGRGGTRGGRRGGQGNW, YGKASRGGGNHQ
Heterogeneous nuclear ribonucleoprotein F	HNRPF	P52597	-	+	FVVKLRGLPWSC, GFVLRGLPFGC, HCVHMRGLPYKA
Heterogeneous nuclear ribonucleoprotein G	HNRPG	P38159	-	+	ETNKSRRGFVFT, FESGRGPPPPP, PPRSRRGPPRGLRGGRRGGSGGTRGPPSRGGHMDD, NMSSSRGPLPVKRGPPPRS, RAPVSRGRDSYG, DDYPSRGGYSDRD, SAPPTRGGPPSY, VGRQERGLPPSM, PPSMERGGYPPPR, YSSSSRGAAPGGGGGSRSDRGGGSR
Heterogeneous nuclear ribonucleoprotein H	HNRH1	P31943	-	+	FVVKVRGLPWSC, GFVLRGLPFGC, RFGAGRGGYNSIGRGGAGFER, FERMRGGAYGGG, HCVHMRGLPYRA
Heterogeneous nuclear ribonucleoprotein H'	HNRH2	P55795	-	+	FVVKVRGLPWSC, GFVLRGLPFGC, RFGAGRGGYNSIGRGGAGFER, FERMRGGAYGGG, HCVHMRGLPYRA
Heterogeneous nuclear ribonucleoprotein H3	HNRH3	P31942	-	+	GTVLRGLPFGC, RPIGGRRGGYGA, YGAGRGGSMYDR, YDRMRGGDGYD, RMRDGRMGGGH, HFVHMRGLPFRA, LGGYRGGGGSG, SGGGWRGM
Heterogeneous nuclear ribonucleoprotein K	HNRPK	P61978	+	+	MFDDRRGRPVGF, VGFPMRGGGFDRM, RMPGRGGRRPMP, DMSPRGGPPPPP, PPPPGRGGRRSRAR, PPPPRGGDLMA, MAYDRRGRPGDR, SYAGGRGGYGD
Heterogeneous nuclear ribonucleoprotein M	HNRPM	P52272	-	+	EKNIKRGGNRF, AEGKSRGCAVVE, KDGKSRGIGTVT, SNALKRGEIIAK, AIEMERGGFGGS
Heterogeneous nuclear ribonucleoprotein Q	HNRPQ	O60506	-	+	LTGLNRGYAFVT, DKKKNRGGFCLE, MPPPTRGGRRGGGGYGY, DYHNYRGGYEDP, FQVGARGGRRGARGAAPSRRGAAPPRGRAGYS, AGYSQRGGPGSARGVRRGARGGAQQRRGGVRRGARGRRGGNVGG
Heterogeneous nuclear ribonucleoprotein R	HNRPR	O43390	-	+	LSGQNRGYAFIT, DKKKNRGGFCLE, VHFEDRGAAVKA, MPPPIRGGRRGGRRGGYGY, DYHNYRGGYEDP, DGYAVRGGGGRRGGRRGAPPPRRGGRRGAPPRGRAGYS, AGYSQRGAPLPG, PLGPPRRGGRRGGGPAQ, PAQQRRGGRRGGRRGGNVGG
Heterogeneous nuclear ribonucleoprotein U	HNRPU	Q00839	+	+	QPQQRGAKEA, GGDKKRGGVRRPR, REDHRRGGYFEYI, LEDRVRRGPKGPE, KNQFNRRGGHRRGGFNMRGGNFRGGAPGNRRGGYNNRGNMPPRRGGGGGG, PVFPGRGYSNRGGYNNRGGMPNRRGGYNNQ, YNQFNRRGGNRRGGYNNQ
Heterogeneous nuclear ribonucleoprotein U-like protein 1	HNRL1	Q9BUJ2	-	+	EELQRRGLDRGLKAE, PYEENRRGGYFEHR, HREDRRGRSPQ, SYGVRGRVCFE, LSERIRGTGPK, KRFDNRGGGGRRGGGGGGF, QRYENRRGGNRRGGFQNRGGGGSG, GGGNYRGGFNRS, NNSNRRGYSNRA
Heterogeneous nuclear ribonucleoprotein U-like protein 2	HNRL2	Q1KMD3	-	+	SELQRRGLDSRGLKVDL, SDGERRGGVRRQR, YGFDGRGLKAEN, RQNRSRGGQYVG, VGGQRRGYDNRA, GQPNRRGGYRNF, FYDRYRGGYDRF
Heterogeneous nuclear ribonucleoproteins A2/B1	ROA2	P22626	+	+	ASKRSRGGFVFT, QSGKKRGGFVFT, SSRSGRRGNFGF, GFGDSRGGGGNF, PGSNFRGGSDGY, GYGSRRGGFDGY, GYGGRRGGYGGG
Histone chaperone ASF1B	ASF1B	Q9NVP2	-	-	-
Histone H1.0	H10	P07305	+	+	-
Histone H1.2	H1.2	P16403	+	-	-
Histone H1.3	H1.3	P16402	+	-	-
Histone H1.4	H1.4	P10412	+	-	-
Histone H1.5	H1.5	P16401	+	-	-
Histone H1x	H1X	Q92522	+	-	GGGERRGAPAAA, DKKPARGGQKPEQ
Histone H2A.Z	H2AZ	P0C0S5	+	-	LQLAIRGDEELD
Histone H2AV	H2AV	Q71U9	+	-	LQLAIRGDEELD
Histone H2B type 1-B	H2B1B	P33778	+	-	-
Histone H2B type 1-C/E/F/G/I	H2B1C	P62807	+	-	-
Histone H2B type 1-D	H2B1D	P58876	+	-	-
Histone H2B type 1-H	H2B1H	Q93079	+	-	-
Histone H2B type 1-J	H2B1J	P06899	+	-	-
Histone H2B type 1-K	H2B1K	O60814	+	-	-
Histone H2B type 1-L	H2B1L	Q99880	+	-	-
Histone H2B type 1-M	H2B1M	Q99879	+	-	-
Histone H2B type 1-N	H2B1N	Q99877	+	-	-
Histone H2B type 1-O	H2B1O	P23527	+	-	-
Histone H2B type 2-E	H2B1E	Q16778	+	-	-
Histone H2B type 2-F	H2B2F	Q5QNW6	+	-	-
Histone H2B type F-S	H2B2FS	P57053	+	-	-
Histone H3.1	H3.1	P68431	+	-	LARRIRGER
Histone H3.1t	H31T	Q16695	+	-	LARRIRGER
Histone H3.2	H3.2	Q71D13	+	-	LARRIRGER
Histone H3.3	H3.3	P84243	+	-	LARRIRGER
Histone H4	H4	P62805	+	-	MSGRRGGKGG, RRLARRGGVKRI, IYEETRGVLKVF
Histone-binding protein RBBP7	RBBP7	Q16576	-	-	PDLRLRGGHQKEG
Importin subunit alpha-2	IMA2	P52292	-	-	-
Importin subunit beta-1	IMB1	Q14974	-	-	-
Importin-7	IPO7	Q95373	-	-	IIEALRGTMDPA, LQCKRRGGIDQCI
Insulin-like growth factor 2 mRNA-binding protein 1	IF2B1	Q9NZI8	-	+	PENRRGGFGSRGGPQQG

Protein AATF	AATF	Q9NY61	-	-	-
Protein C14orf166	CN166	Q549M8	-	-	YKIEDRGNLRNI
Protein FAM102B	F102B	Q5T813	-	-	-
Protein FAM111B	F111B	Q6SJ93	-	-	GLFYQRGFNVHA
Protein FAM98B	FA98B	Q52LJ0	-	-	MRGPEPGP
Protein KIAA0082	K0082	Q8N1G2	-	-	SQKGRRLGLTLRQFDQEL, PYEMIRGVFFLN
Protein RCC2	RCC2	Q9P258	-	-	AGPRKRGGPAGR, FDFPGRGASQIY
Protein SDA1 homolog	SDA1	Q9NVU7	-	-	LQKKFRGKPTFA, SDEEPRGELLSL
Pumilio homolog 1	PUM1	Q14671	-	+	QQQVLRRGGASQR, LAERIRGHVLSL, IVAEIRGNVLSL
Putative ATP-dependent RNA helicase DHX30	DHX30	Q7L2E3	-	+	PLRDSRGSSEFEM, ELWRRRGPVWQEQ, YVTEGARGARCNV, SKPPSRGGALLF, LHIDARGEPGGI, NVIQRGRGRAGRC, LAELLRGGPCGSF
Putative ATP-dependent RNA helicase DHX57	DHX57	Q6P158	-	+	GKGSRRGGRRGSHA, DAGSERGLSGEE, NVQLQRGEFVVS, LCTCPRGSRIS
Putative pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15	DHX15	O43143	-	+	DRERDRGDRERE, LPGPKRGVACTQ
Putative RNA-binding protein Luc7-like 1	LUC7L	Q9NQ29	-	-	SRSHSRGHRRAS, SGRSERGPPDWR
Putative RNA-binding protein Luc7-like 2	LC7L2	Q9Y383	-	-	-
Putative rRNA methyltransferase 3	RRMJ3	Q8IY81	+	+	CDFLARGGSFIT, SLCTIRGHQLE, ELAGVVRGHQGLR, SWEPLRGGKRSRGPKSDD, RPAGVRGHFKVV
Pyruvate dehydrogenase E1 component subunit beta	ODPB	P08559	-	-	KQKIIRGFCHLC, GFTFTRGLSVRE, TDYKRGDFIPG, PPFVVRGANQWI
Pyruvate kinase isozymes M1/M2	KPYM	P14618	-	-	GIMVARGDLGIE, QAHLYRGIFPVL, NVGKARGFFKKG
Ras GTPase-activating protein-binding protein 1	G3BP1	Q13283	+	+	NIPPQRGPRPIR, RPIMFRGEVRLN, RDNRLRGGPGRGLGGG, LGGGMGRPPRGGMVQK, GFGVGRGLAPR
Ras GTPase-activating protein-binding protein 2	G3BP2	Q9UN86	-	+	PGFPPRGRPRGDMEQN, KPIMFRGEVRLN, RERETRGGGDDR, IRRNDRGPGGPRGIVGGG, RDRDGRGPPRGGMAQK, KLGSRGRTGQME
Ras GTPase-activating-like protein IQGAP1	IQA1	P46940	-	-	PALGLRGLQQQN, LQARCRGYLVRQ, IQSQWRGYKQKK, VVSFNARGRGNALR
Ras-related protein Rab-35	RAB35	P15286	-	-	TSTYYRGTHGVI
Regulator of nonsense transcripts 1	RENT1	Q92900	+	+	WFCNRRGNTSGS, GPAARGTPKPK, KGKTRGGGRQKN
Replication initiator 1	REPI1	Q9BWE0	+	-	LERRCRGPLAMG, LGKESRGLRQQQ, CGQSFRGWVALV, LGPRPRGRPAVT
Rho GTPase-activating protein 1	RHG01	P20936	-	-	TLSNERGAQQHV
Ribonuclease P protein subunit p30	RPP30	P78346	-	-	DLKALRGLVETA, NVAIDRGLAFEL, RPLEIRGPDVA
Ribonuclease P protein subunit p40	RPP40	O75818	-	-	-
Ribonucleases P/MRP protein subunit POP1	POP1	Q99575	-	-	GFVADRGVKHS, IPFIYRQVVRGG, SSEDSSRGGRRAP, RRAPGRQQGLT, QPAAQRGLVLLR
Ribosomal L1 domain-containing protein 1	RL1D1	O76021	-	+	HGKKKRRGKAQVK, NPSTPRGKRRKA
Ribosomal protein L7-like 1	RL7L	Q6DK11	-	-	ELILKRQAKVK, GTPGYRGERINQ
RNA 3'-terminal phosphate cyclase	RTC1	O00442	-	+	CDIKTRGYYPKG, INLTERGCVTKI, SSLKRGVNRADK
RNA exonuclease 4	REXO4	Q9GZR2	-	-	DQEASRGSVPSG, DIVPERGDIHKK
RNA U small nuclear RNA export adapter protein	RNUXA	Q9H814	-	+	-
RNA-binding protein 26	RBM26	Q5T8P6	-	+	RYNRRRGRSRSY, KGFCMRGDMCPF, LEAAKRGILSSGRRGIHSRGRGAVHGRGRGRGVRPGHA
RNA-binding protein 35B	RB35B	Q9H6T0	-	+	TVVRARGLPWQS, GLNVARGGVALC, VILRLRGLPFA, DCVRLRGLPYTA
RNA-binding protein 39	RBM39	Q14498	-	+	RDRRFGRYRSP, FNSAIRGKIGLP, TEDMLRGIFEPF
RNA-binding protein 4	RBM4	Q9BWF3	-	+	AVEAIRGLDNTA
RNA-binding protein Musashi homolog 2	MSI2H	Q96DH6	-	+	TTKRSRGGFVFT, TTNRRHGFVFT, FPPGTRGRARGLPYTM, VATYGRGYPGA, AVAAARGSGSNP
RRP15-like protein	RRP15	Q9Y3B9	-	-	QRIATRGVVQLF, FISVLRGMDGST
RRP1-like protein	RRP1	P56182	-	-	LNNITRGIFETI, SEGGERGDALSQ, RLQQRGKGEKE, KRSRRRGVADP, LRDQPRGRGQGRGARQRR
RuvB-like 1	RUVB1	Q9Y265	-	-	ITDKLRGEINKV, IFASNRRGNVCIRGTEDIT
RuvB-like 2	RUVB2	Q9Y230	+	-	AHSHIRGLDLD, IMATNRGITRIRGTSYQS
Serine hydroxymethyltransferase, mitochondrial precursor	GLYM	P34897	-	-	TGEANRGTWQGE, KDRQCRGLELIA, THKTLRGRASGL, DALLERGYSLVS
Serine/threonine-protein kinase 6	STK6	O14965	-	-	-
Serine/threonine-protein phosphatase 2A catalytic subunit alpha isoform	PP2AB	P67775	-	-	GDYVDRGYYSVE, RITILRGNHESR, SDPDRGGWGIS, WGISPRGAGYTF, DPAPRRGEPHVT
Serine/threonine-protein phosphatase 2A catalytic subunit beta isoform	PP2AA	P62714	-	-	GDYVDRGYYSVE, RITILRGNHESR, SDPDRGGWGIS, WGISPRGAGYTF, DPAPRRGEPHVT
Serine-threonine kinase receptor-associated protein	STRAP	Q9Y3F4	-	-	-
Signal recognition particle 14 kDa protein	SRP14	P37108	-	+	-
Small nuclear ribonucleoprotein E	RUXE	P62304	-	+	MAYRGGQKQV
Small nuclear ribonucleoprotein Sm D2	SMD2	P62316	-	-	SKMFLRQDSVIV
Small nuclear ribonucleoprotein-associated protein N	RSMN	P63162	-	+	GLVLLRGENLVS, GRAAGRGVPAGV, LAGPVRGVGGPS, MTPQGRGTVA, QYPPRGTPTPPP, GLPPARGTPIGM, PPPGIRGPPPPP
Small nuclear ribonucleoprotein-associated proteins B and B'	RSMB	P14678	-	+	GLVLLRGENLVS, GRAAGRGVPAGV, LAGPVRGVGGPS, MTPQGRGTVA, QYPPRGTPTPPP, PPPMGRGAPPPP, GIPPRGTPTMGM, PPPMGRGPPPPP
Spectrin alpha chain, brain	SPTA2	Q13813	-	-	AASTNRGKDLIG, WINGIRGLVSSD, NADRIRGVIDMG, NSLIERGACAGS, QLAAARGQRLEE
Splicing factor 3B subunit 1	SF3B1	O75533	-	-	AATPRGDTPGH, TPRTRDGGDSIG, GIRQHRGKGLAA, GRIADRGAEYVS
Splicing factor 3B subunit 2	SF3B2	Q13435	+	+	NRPVLRGEGDGD, PVPFRGPPPPP, KPQVRVGVSSS, DSTRSRGSDSPA, YLQKGRGIEKPP, QPQDSRGGSKKY
Splicing factor 3B subunit 3	SF3B3	Q15393	+	+	EIVSVRGKILEL, LDDPERGMIFVC, YVACGRGPRSSL, DELGERGSIGFL, KALWDRGLLNGA
Splicing factor 3B subunit 4	SF3B4	Q15427	-	+	MGMPRGPFFGS, PPHGMRGPPPLM, PYGYQRGLPPP, PPVPPRGLRGLP
Splicing factor U2AF 35 kDa subunit	U2AF1	Q01081	-	+	MGECTRGGFCNF, SRSRDRGGGGGGG
Splicing factor, arginine/serine-rich 1	SFRS1	Q07955	-	+	GGGVIRGPAGNN, DLKNRRGGPPFA, FPRSGRGTGRGGGGG, GGGAPRGRYGP, SPRRSRGSPPRY
Splicing factor, arginine/serine-rich 10	TRA2B	P62995	-	+	QSRRSRGAFAVY, RDIYDRGYDRGYDDRD, YRSYRGGGGGG, SPYYSRGGYRSR

Splicing factor, arginine/serine-rich 2	SFRS2	Q01130	-	+	YTKESRGFAFVR, SHHSRRGPPRRR
Splicing factor, arginine/serine-rich 3	SFRS3	P84103	-	+	KRSRNRGPPPSW
Splicing factor, arginine/serine-rich 4	SFRS4	Q08170	-	+	IVEHARGPRRDG, VEEEKRGVSVRGRSSEQ, ERSRKRGRSKRDS, SQREGRGESENA
Splicing factor, arginine/serine-rich 5	SFRS5	Q13243	-	+	DIDLKRGFGFVE, ARASRGGGRGRYSDR, EKSQKRGSSRS
Splicing factor, arginine/serine-rich 6	SFRS6	Q13247	-	+	IVEHARGPRRDR, KPKSDRGSHTS
Splicing factor, arginine/serine-rich 7	SFRS7	Q16629	-	+	AEDAVRGLDGKV, SHSRSRGRRYSR, SRSRSGRRRSRS
Splicing factor, arginine/serine-rich 9	SFRS9	Q13242	-	+	GWADERGGEGDG, RTYGGRGGWPRGGRNGP, SRSGSRGRDSPY, SPYQSRGSPHYF
Splicing factor, proline- and glutamine-rich	SFPQ	P23246	+	+	DRFRSRGGGGGG, GGFHRRGGGGGRGLHDF, GLNQNRGPMGPG, PKPPHRGGGEPRGGROHH, DDTPMRGRQLRV, VIVDDRGRSTGK, GGGQPRGMGPGT, PAGYGRGREEYE
Squamous cell carcinoma antigen recognized by T-cells 3	SART3	Q15020	-	+	DSIMTRGNAKYA, KKKKIRGPEKRGAEDEDD, PIFSNRGGDFRGYCYVE, QTYGARKGKRTQ
Stress-70 protein, mitochondrial precursor	GRP75	P38646	-	-	GAASRSGPTAAR, IPPAPRGVPOIE
Structural maintenance of chromosomes protein 3	SMC3	Q9UQE7	-	-	KEKEERGIARLA, YAKQGRGSQFTS, DQVSHRGALTTG, QEELDRGYKSIM, SGESERGGSGSQS
Structural maintenance of chromosomes protein 4	SMC4	Q9NTJ3	-	-	ILNEHRGEKLNLR, AMNRSRQKVLDA
Succinyl-CoA ligase [GDP-forming] beta-chain	SUCB2	Q96199	-	-	ILAGGRGKGVFN
Sucrose nonfermenting protein 2 homolog	SMCA5	O60264	+	-	RDYQVRGLNWLI, DGILERGAKTA, MAQIERGEARIQ, AEKKRGRPKPST, GAPDGRGRKKKL
Superkiller viralicidal activity 2-like 2	SK2L2	P42285	+	+	RSMLYRGSEVMR, MRDSERGVVWEE, LPLLKRGIGIHH, GRAGRGMDDRGIVILM
Suppressor of SWI4 1 homolog	SSF1	Q9NQ55	-	-	SFVFRGCTGRN, PVGASRGMKLL, QAVAGRGNMRAQ, RWEMDRGRGRLCDQ, GAQARRGPRGASRDG, SRDGGRRGRGRPGKR
Surfeit locus protein 6	SURF6	O75683	+	+	KIQEARGGGSAK, RLDELRGQDEGK
Target of EGR1 protein 1	TOE1	Q96GM8	+	+	-
T-complex protein 1 subunit alpha	TCPA	P17987	-	-	KYTDIRGQPRYP, ASIILRGANDFM
T-complex protein 1 subunit beta	TCPB	P78371	-	-	CTIVLRGATQI
T-complex protein 1 subunit delta	TCPD	P50991	-	-	GAAGGRGKAYQ, VTIVVRGSNKL
T-complex protein 1 subunit eta	TCPH	Q99832	-	-	TTLGPRGMDKLI, LIVDGRGKATIS, CTFILRGGAEQF, PTAAGRGRGRGP
T-complex protein 1 subunit gamma	TCPG	P49368	-	-	DSCVLRGVMINK, CTILLRGASKEI
T-complex protein 1 subunit theta	TCPO	P50990	-	-	STIVLRGSDNL
Thyroid receptor-interacting protein 13	TRP13	Q15645	-	-	VEVHQRGSSTAK
Transcriptional activator protein Pur-alpha	PURA	Q00577	+	-	LKENQRGRFLRI, RQTVNRGPGLS
Transcriptional activator protein Pur-beta	PURB	Q96QR8	+	-	DSGSEGGGGGP, FQPASRGGGEQE, LKENQRGRFLRI, RQTVNRGGGGFG, KLYERRGGGSGG
Translation initiation factor eIF-2B subunit delta	EI2BD	Q9UI10	-	-	DLQCKRGEHVAL
Tripartite motif-containing protein 25	TRI25	Q14258	-	-	KASKLRGISTKP
Tripartite motif-containing protein 56	TRI56	Q9BRZ2	-	-	GPQPHRGGPRNK, GKGASRGLRALV
tRNA (cytosine-5)-methyltransferase NSUN2	NSUN2	Q08J23	-	+	MGRRSRGRRLQ, EGGGKRGEAGWE, LRIATRGAEQLA, VLCGWRGKASIR
Tubulin alpha-1A chain	TBA1A	Q71U36	-	-	ANNYARGHYTIG, CCLLYRGDVVPK
Tubulin alpha-4A chain	TBA4A	P68366	-	-	ANNYARGHYTIG, CCLLYRGDVVPK
Tubulin beta-2C chain	TBB2C	P68371	-	-	APLTSRGSQQYR, VAAVFRGRMSMK, CDIPRGLKMSA
Tuftelin-interacting protein 11	TFP11	Q9UBB9	-	-	GYVPGRGLGKNA, ENMAQRGIGVAA, VIYIDRGVVVQ
U1 small nuclear ribonucleoprotein 70 kDa	RU17	P08621	-	+	RSQKPRGYAFIE, LVDVERGRTVKG, LGTRRRGGADVN, RKEELRGGGGDM, DREHKRGERGSRGRDEARGGGGQ
U1 small nuclear ribonucleoprotein A	SNRPA	P09012	-	+	RSLKMRGQAFVI
U2 small nuclear ribonucleoprotein A'	RU2A	P09661	-	+	RELDLRGYKIPV, MFKGKRGAQLAK
U2 small nuclear ribonucleoprotein B''	RU2B	P08579	-	+	KTMKMRGQAFVI, IISKMRGTFADK
U3 small nucleolar ribonucleoprotein protein IMP4	IMP4	Q96G21	-	-	AQRMRGRHEVGG, VVHEHRGTPVGL
U4/U6 small nuclear ribonucleoprotein Prp3	PRPF3	O43395	-	-	-
U4/U6 small nuclear ribonucleoprotein Prp4	PRP4	O43172	-	-	LLHTLRGHNTNV
U5 small nuclear ribonucleoprotein 200 kDa helicase	U520	O75643	+	+	LLHDDRGPVLEA, EIVLNRGWAQLT, AKTKVRGLIEII
Uncharacterized protein C1orf156	CA156	O95568	-	-	-
UPF0027 protein C22orf28	CV028	Q9Y310	-	-	LRNACRGGGVGG, ARAKRGRLPQLG, IHSGSRGLGHQV
UPF0384 protein CGI-117	U384	Q9Y3C1	-	-	-
Zinc finger CCHC type antiviral protein 1	ZCC2	Q7Z2W4	-	+	CDHFTRGNCRFP, YQSCPGRGVVPFQ, VQMKRGPDPHQ
Zinc finger CCHC-type with G patch domain-containing protein	ZGPAT	Q8N5A5	+	-	APAAAARGSGSET, WEVHTRGIGSRL, AVVLRGKSLDQ, KPPCRGRGARPGG
Zinc finger CCHC domain-containing protein 8	ZCHC8	Q6NZY4	+	+	NISTPRGIPDEW, TPPLPRGTPPPV

total: 389 identified proteins, 216 nucleic acids-binding proteins, 87 identified proteins without RG-box motive