

Supplementary Table 1. Immunophenotyping results on several occasions of the index patient bearing the homozygous *RASGRP1* stop gain mutation.

Patient/ Age	7 years old	8 years old	12 years old	12.5 years old
ALC (cells/mm ³) (normal range)	4064 (1100-5900)	3100 (1100-5900)	2700 (1000-5300)	2700 (1000-5300)
Immunoglobulins				
Ig A (g/dl)	660 (70-303)	680 (70-303)	640 (67-433)	766 (67-433)
Ig G (g/dl)	2030 (764-2134)	3400 (764-2134)	1720 (835-2094)	1600 (835-2094)
Ig M (g/dl)	188 (69-387)	346 (69-387)	218 (47-484)	306 (47-484)
Lymphocyte subsets				
CD3 (%)	64 (55-78)	61	75	81
(cells/mm ³)	2600 (700-4200)	1891 (700-4200)	1125 (800-3500)	2187 (800-3500)
CD4 (%)	13 (27-53)	10 (27-53)	13 (25-48)	8 (25-48)
(cells/mm ³)	528 (300-2000)	310 (300-2000)	195 (400-2100)	216 (400-2100)
CD8 (%)	55 (19-34)	52 (19-34)	58 (9-35)	70 (9-35)
(cells/mm ³)	2235 (300-1800)	1612 (300-1800)	870 (200-1200)	1890 (200-1200)
CD19 (%)	10 (10-31)	11 (10-31)	9 (8-24)	3 (8-24)
(cells/mm ³)	406 (200-1600)	341 (200-1600)	135 (200-600)	81 (200-600)
CD16+56 (%)	22 (4-26)	26 (4-26)	11 (6-27)	13 (6-27)
(cells/mm ³)	894 (90-900)	806 (90-900)	165 (70-1200)	351 (70-1200)
CD45RA			55	
CD45RO			51	
TCR αβ			62	
TCR γδ			17	
ALC, absolute lymphocyte count; TCR, T cell receptor.				

Supplementary Table 2. Immunophenotyping of the healthy mother.

Value	Mother
WBC (x10 ³ /μl)	5.9 (4.1-11.2)
ALC (x10 ³ /μl)	2.3 (1.0-2.8)
ANC (x10 ³ /μl)	2.3 (1.8-6.4)
Monocytes (x10 ³ /μl)	0.5 (0.3-0.9)
Eosinophils (x10 ³ /μl)	0.4 (0.1-0.5)
Basophils (x10 ³ /μl)	0 (0-0.2)
Immunoglobulins	
Ig A (g/dl)	5.42 (0.9-4.5)
Ig G (g/dl)	18.5 (8-18)
Ig M (g/dl)	3.29 (0.7-2.5)
Lymphocyte subsets	
CD3 (%)	68 (55-83)
CD3 (1000/μl)	1.5 (1.0-2.8)
CD4 (%)	42 (28-57)
CD4 (1000/μl)	1.0 (0.3-1.4)
CD8 (%)	25 (10-39)
CD8 (1000/μl)	0.7 (0.2-0.9)
CD19 (%)	9 (6-19)
CD19 (1000/μl)	0.2 (0.1-0.5)
CD16+56 (%)	14 (7-31)
CD16+56 (1000/μl)	0.3 (0.09-0.6)

Supplementary Table 3. Relative immunophenotyping values of patient, mother brother and a healthy donor.

Subset (rel. proportions in %)	Healthy Donor	Mother	Brother	Patient
CD3+	47.8	66.7	67.0	80.0
CD3+CD4+	66.1	19.3	24.0	10.8
CD3+CD8+	28.9	53.9	52.9	74.7
CD19+	19.6	21.6	14.3	2.51
Gated on CD3+:				
iNKT cells (TCR α 24+TCR β 11+)	0.25	0.89	0.25	0.03
recent thymic emigrants (CD3+CD4+CD45RA+CD31+)	17.2	17.4	12.2	5.92
TCR α 7.2+CD161+ (MAIT)	3.21	1.53	1.13	0.06
Gated on CD3+CD4+:				
CD45RA+CCR7-	2.2	1.0	1.9	0.33
CD45RA+CCR7+	26.6	45.6	27.3	10.8
CD45RA-CCR7+	68.9	52.7	69.7	51.9
CD45RA-CCR7-	2.2	0.7	1.1	37.0
CD3+CD4+FOXP3+	5.51	n.d.	n.d.	2.38
CD45RA+CD27-	0.064	n.d.	n.d.	0.44
CD45RA+CD27+	70.6	n.d.	n.d.	16.7
CD45RA-CD27+	26.3	n.d.	n.d.	76.8
CD45RA-CD27-	3.09	n.d.	n.d.	6.04
CD45RA+CD28-	0.073	n.d.	n.d.	0.6
CD45RA+CD28+	70.1	n.d.	n.d.	16.2
CD45RA-CD28+	29.8	n.d.	n.d.	81.9
CD45RA-CD28-	13.3	n.d.	n.d.	1.32
Gated on CD3+CD8+:				
CD45RA+CCR7-	18.2	5.6	6.6	17.4
CD45RA+CCR7+	31.1	63.9	66.7	2.45
CD45RA-CCR7+	35.1	27.0	23.1	1.16
CD45RA-CCR7-	12.0	3.5	3.7	79.0
CD45RA+CD27-	0.2	n.d.	n.d.	18.1
CD45RA+CD27+	83.5	n.d.	n.d.	3.08
CD45RA-CD27+	14.5	n.d.	n.d.	16.8
CD45RA-CD27-	1.82	n.d.	n.d.	62
CD45RA+CD28-	2.98	n.d.	n.d.	21.7
CD45RA+CD28+	81.2	n.d.	n.d.	1.3
CD45RA-CD28+	14.2	n.d.	n.d.	2.9
CD45RA-CD28-	1.56	n.d.	n.d.	74.1
Gated on CD19+:				
IgD+CD27- (naïve)	58.9	48.4	44.9	84.0
IgD+CD27+(memory non-switched)	7.0	16.3	22.1	7.09
IgD-CD27+(memory)	30.5	31.1	28.0	3.15
IgD-CD27-	3.6	4.0	5.0	5.75
CD21 ^{low} CD38-	2.6	4.78	8.09	20.6
CD21 ^{low} CD38+	0.61	1.14	1.17	17.3

Supplementary Table 4. Evaluation of the detected variant in *RASGRP1* (Chr15: 38805097: G>A; c. C726T, p. Arg246*) in exome variant databases (last accessed July, 13, 2016).

Source	Found
dbSNP v142 ¹	no
60,706 Exomes from the Exome Aggregation Consortium ²	no
6503 Exomes from NHLBI Exome Sequencing Project ³	no
CBio Portal ⁴	no
108 Exomes Hacettepe Exome Project	no
In-house collection of 500 individuals (219 Exomes including 96 PID patients, 12 Genomes and 269 target enrichment screens on 229 PID patients and 40 SCID patients)	no

¹<http://www.ncbi.nlm.nih.gov/SNP/>; ²<http://exac.broadinstitute.org/>;

³<http://evs.gs.washington.edu/EVS/>; ⁴http://www.cbioportal.org/data_sets.jsp

Supplementary Table 5a. Evaluation of the estimated population frequency severe homozygous *RASGRP1* variants. To date, no severe (stopgain, frameshift, splice-site) homozygous variants have been observed in *RASGRP1*. This table lists rare heterozygous variants in *RASGRP1* which have been identified and indicates an estimated population frequency of homozygous *RASGRP1* loss-of-function individuals based on heterozygous carrier frequency, as obtained from ExAC browser (accessed April 29, 2016).

Consequence	Annotation	Homozygous Alleles	Heterozygous Allele count							
			African	East Asian	European (Non-Finnish)	Finnish	Latino	Other	South Asian	Allele count
p.Gln793Ter	stop gained	0	0	0	1	0	0	0	0	1
p.Leu788Phefs	frameshift	0	0	0	1	0	0	0	0	1
p.His786Glnfs	frameshift	0	0	0	1	0	0	0	0	1
c.2260-1G>A	splice acc.	0	0	0	1	0	0	0	0	1
p.Arg98Thrfs	frameshift	0	0	0	1	0	0	0	0	1
p.Gly92Ter*	stop gained	0	0	0	0	0	1	0	0	1
p.Met76Valfs	frameshift	0	0	23	2	0	1	0	8	34
p.Val55Ter	frameshift	0	0	0	0	0	2	0	0	2
p.Gln545Ter	stop gained	0	0	0	1	0	0	0	0	1
p.Thr201Ilefs	frameshift	0	1	0	0	0	0	0	0	1
c.570-1G>A	splice acc.	0	0	0	1	0	0	0	0	1
c.570-1G>T	splice acc.	0	0	0	0	0	0	0	1	1
c.1428+1G>A	splice donor	0	0	1	0	0	0	0	0	1
p.Asp371Ilefs	frameshift	0	0	0	1	0	0	0	0	1
p.Glu367Ter	stop gained	0	0	0	1	0	0	0	0	1
p.Ile354Phefs	frameshift	0	0	0	1	0	0	0	0	1
c.327-1C>G	splice acc.	0	0	0	0	0	0	0	3	3
p.Arg48Ter	stop gained	0	0	1	1	0	0	0	0	2
p.Glu27Argfs	frameshift	0	0	0	0	0	0	0	1	1
p.Phe3_Serfs	frameshift	0	1	0	4	0	0	0	0	5
p.Pro8Hisfs	frameshift	0	0	0	1	0	0	0	0	1
SUM of putative loss-of-function population alleles			2	25	18	0	4	0	13	62
Total population alleles			9802	8622	66716	6614	11574	898	16510	120736
loss-of-function carrier frequency			2.040E-04	2.900E-03	2.698E-04	0.0001511	3.456E-04	0.00111	7.874E-04	5.135E-04
Expected rate of nullizygous per million (1 in X non-consanguineous conceptions)			24	0.12	13.7	< 43.8	8.3	<0.80	1.6	3.8

Supplementary Table 5b. *RASGRP1* conservation evaluation. CADD Score, RVIS Score and expected versus observed number of nonsense variants in *RASGRP1* are summarized below.

Performed Test	Score
Chr15: 38805097: G>A variant CADD (v1.3) score	35
CCDS Size (including canonical splice) ENSG00000172575	5023
Number of unique protein-coding variants (ExAC)	594
RVIS Score Version 2 of RASGRP1	-0.82 (11.77% most intolerant)
CCDS ExAC 'Expected' number of nonsense variants	28.3
CCDS Observed nonsense variants (ExAC)	7
Binomial Exact test p-value	1.29619E-06

Supplementary Table 6. Comparison of human RASGRP1 deficiency with findings from *Rasgrp1*^{-/-} knockout mice^{2,3,4,5}.

	Human RASGRP1 Deficiency	<i>Rasgrp1</i>^{-/-} mouse model
Peripheral T-cells		
CD3+ CD4+	↓	↓ ⁸
CD3+CD8+	↑	↓ ⁸
TCRαβ	↓	n.d.
TCRγδ	↑	↑ ⁸
Proliferation	↓	↓ ⁸
ERK signaling	↓	↓ ⁸
iNKT cells	↓	↓ ¹⁸
CD19+		
NK cells		
CD16+CD56+	normal	normal
Cytotoxicity	↓	n.d.
Peripheral B-cells		
CD19+	↓	normal ⁸
CD19+IgD+	normal	n.d.
CD19+IgM+	↑	↑ ⁸
CD19+CD21 ^{low}	↑	n.d.
CD19+ CD38+	↑	n.d.
Autoimmunity		
Splenomegaly	no	yes, older mice ²⁴
Autoantibodies	no	Yes, ANCA ²⁴
Immunodeficiency		
Viral infections	↑	n.d.
Bacterial Infectios	↑	n.d.

Supplementary Table 7. Absence of all autoantibodies tested in index patient P1.

Nuclear Antibodies	
Anti-nuclear antibody (ANA)	negative
Rheuma Serology	
Rheumatoid-factor IgG	negative
Rheumatoid -factor IgA	negative
Rheumatoid -factor IgM	negative
Rheumatoid factor latex test	negative
Anti-streptolysine O	negative
Collagen type I	negative
Collagen type II	negative
Collagen type III	negative
Collagen type IV	negative
Thyroid gland Antibodies	
Thyreoglobulin antibody	negative
Vessel & kidney Antibodies	
Neutrophil cytoplasmic antibody (ANCA)	negative
Proteinase-3 antibody	negative
Myeloperoxidase	negative
Phospholipid antibodies	
Cardiolipin IgG	negative
Cardiolipin IgA	negative
Cardiolipin IgM	negative
B2-glycoprotein IgG	negative
B2-glycoprotein IgA	negative
b2-glycoprotein IgM	negative
Intestinal Antibodies	
Gliadin IgG	negative
Gliadin IgA	negative
Tissue transglutaminase IgG	negative
Tissue Transglutaminase IgA	negative

Supplementary Table 8a. FCA, FCB and SAINT Score of identified RASGRP1 interactors in HEK293 cells.

BAIT	PREY	FC_A	FC_B	SAINT
RASGRP1	RASGRP1	81.83	68.16	1
RASGRP1	DYNLL1	34.49	34.49	1
RASGRP1	DYNLL2	31.78	31.78	1
RASGRP1	TUBA1A	25.44	25.44	1
RASGRP1	TUBA1C	25.14	25.14	1
RASGRP1	TUBB4B	15.18	15.18	1
RASGRP1	TUBB3	13.37	13.37	1
RASGRP1	TUBB	12.92	9.68	1
RASGRP1	TUBA8	12.16	12.16	1
RASGRP1	TUBB2B	11.56	11.56	1
RASGRP1	TUBA1B	8.6	5.12	1
RASGRP1	SLC25A5	6.43	6.43	1
RASGRP1	EMD	5.22	5.22	1
RASGRP1	HSPA1B	4.35	2.44	1
RASGRP1	RPS3	4.02	4.02	1
RASGRP1	SLC25A6	3.93	3.28	1
RASGRP1	ATP5A1	3.72	3.72	1
RASGRP1	CAD	3.11	3.11	1
RASGRP1	IRS4	3.03	2.73	1
RASGRP1	PPP6R3	2.81	2.81	1
RASGRP1	SLC25A13	2.51	2.51	1
RASGRP1	LGALS3BP	2.51	2.51	1
RASGRP1	RHOT2	1.91	1.91	1
RASGRP1	HSPA1L	3.41	2.22	0.99
RASGRP1	HNRNPH1	2.81	2.81	0.99
RASGRP1	HNRNPF	2.51	2.51	0.99
RASGRP1	RPS27	2.21	2.21	0.99
RASGRP1	ATP1A1	1.91	1.91	0.99
RASGRP1	HSPD1	1.6	1.6	0.99
RASGRP1	TUFM	2.21	2.21	0.98
RASGRP1	ATP2A2	1.91	1.91	0.98
RASGRP1	DCAF7	1.91	1.91	0.98
RASGRP1	PPP6C	2.21	2.21	0.97
RASGRP1	RCN2	2.21	2.21	0.97
RASGRP1	SLC25A1	1.91	1.91	0.97
RASGRP1	SLC25A11	1.6	1.6	0.97
RASGRP1	RMDN3	1.6	1.6	0.97
RASGRP1	ACSL3	1.6	1.6	0.97
RASGRP1	HLA-B	1.91	1.91	0.96
RASGRP1	HSP90AA1	1.6	1.6	0.96
RASGRP1	HSPA8	2.54	1.42	0.91
RASGRP1	HSPA5	1.74	1.14	0.79

RASGRP1	CHUK	1.54	1.11	0.53
RASGRP1	RPL23	0.5	0.32	0.09
RASGRP1	EEF1A1	1.3	1.3	0
RASGRP1	RPLP2	1.3	1.3	0
RASGRP1	FASN	1.3	1.3	0
RASGRP1	ANKRD28	1.3	1.3	0
RASGRP1	BAG2	1.3	1.3	0
RASGRP1	SLC25A10	1.3	1.3	0
RASGRP1	ACTA2	0	0	0
RASGRP1	HMGB1P1	0	0	0
RASGRP1	HSPA9	0	0	0
RASGRP1	MYL6	0	0	0
RASGRP1	KRT1	0	0	0
RASGRP1	TBC1D4	0	0	0
RASGRP1	HNRNPU	0	0	0
RASGRP1	HSP90AB1	0	0	0
RASGRP1	GFP	0	0	0
RASGRP1	FBXO31	0	0	0
RASGRP1	UBB	0	0	0
RASGRP1	ACTB	0	0	0

Supplementary Table 8b. FCA, FCB and SAINT Score of identified RASGRP1 interactors in Jurkat T cells.

BAIT	PREY	FC_A	FC_B	SAINT
RASGRP1	RASGRP1	41.6	41.6	1
RASGRP1	NA	13.57	3.39	0.3
RASGRP1	TUBB4B	9.99	5.92	0.94
RASGRP1	DYNLL1	9.81	9.81	0.47
RASGRP1	TUBB	7.09	4.28	0.5
RASGRP1	EIF3I	4.83	4.83	0.99
RASGRP1	EMD	4.04	4.04	0.87
RASGRP1	TUBA1B	4.01	2.73	0.05
RASGRP1	EEF2	3.86	2.44	0.5
RASGRP1	TUBB3	3.82	3.82	0.5
RASGRP1	HSPA9	3.41	2.18	0.99
RASGRP1	PDIA3	2.93	2.93	0.49
RASGRP1	TKT	2.91	1.23	0.5
RASGRP1	RPS19	2.82	1.28	0.98
RASGRP1	EEF1A1P5	2.73	2.73	0.49
RASGRP1	NME1	2.63	1.65	0.42
RASGRP1	ATP5A1	2.6	1.43	0.94
RASGRP1	SERBP1	2.56	2.01	0.5
RASGRP1	VCP	2.54	2.54	0.46
RASGRP1	HNRNPM	2.43	2.13	0.5

RASGRP1	MSN	2.38	1.43	0.5
RASGRP1	CCT2	2.34	2.34	0.49
RASGRP1	PGK1	2.33	0.98	0.5
RASGRP1	HSPA8	2.3	1.46	0.14
RASGRP1	SSBP1	2.29	2.29	0.49
RASGRP1	HSPD1	2.25	0.7	0.5
RASGRP1	RBBP4	2.24	2.24	0.49
RASGRP1	HMGB2	2.19	2.19	0.49
RASGRP1	CCT5	2.14	2.14	0.49
RASGRP1	TRMT112	2.14	2.14	0.49
RASGRP1	CCT3	2.13	1.74	0.5
RASGRP1	NME2	2.12	1.33	0.5
RASGRP1	CCT7	2.09	2.09	0.49
RASGRP1	SUB1	2.09	2.09	0.5
RASGRP1	PKM	2.09	0.88	0.81
RASGRP1	MDH2	2.08	1.06	0.5
RASGRP1	HINT1	2.04	2.04	0.5
RASGRP1	PAFAH1B3	2.04	2.04	0.5
RASGRP1	PGAM1	2.03	0.95	0.5
RASGRP1	DUT	1.99	1.19	0.5
RASGRP1	SFPQ	1.99	1.45	0.5
RASGRP1	PCNP	1.99	1.99	0.5
RASGRP1	SOD1	1.99	1.99	0.49
RASGRP1	CCT4	1.94	1.94	0.47
RASGRP1	EZR	1.94	1.94	0.46
RASGRP1	PRKCSH	1.89	1.89	0.48
RASGRP1	YWHAG	1.89	1.89	0.49
RASGRP1	RPL5	1.89	1.08	0.5
RASGRP1	LAP3	1.89	1.89	0.48
RASGRP1	ADA	1.89	1.89	0.5
RASGRP1	CCT8	1.85	1.06	0.5
RASGRP1	CCT6A	1.84	1.84	0.48
RASGRP1	RBBP7	1.84	1.84	0.48
RASGRP1	PPM1G	1.84	1.84	0.49
RASGRP1	PSMA5	1.84	1.84	0.49
RASGRP1	RPL8	1.84	1.84	0.49
RASGRP1	ATP5H	1.84	1.84	0.49
RASGRP1	PRDX5	1.81	1.32	0.5
RASGRP1	NPM1	1.8	1.03	0.5
RASGRP1	NUDC	1.79	1.79	0.49
RASGRP1	SF3A1	1.79	1.79	0.49
RASGRP1	STMN1	1.78	0.71	0.5
RASGRP1	PPIA	1.77	0.68	0.48

RASGRP1	PAICS	1.74	1.74	0.49
RASGRP1	HMGB1	1.74	1.1	0.5
RASGRP1	CNPY2	1.74	1.74	0.49
RASGRP1	MTHFD1	1.74	1.74	0.44
RASGRP1	PSME2	1.74	1.74	0.5
RASGRP1	LUC7L2	1.74	1.74	0.49
RASGRP1	P4HB	1.74	1.74	0.47
RASGRP1	PA2G4	1.74	1.74	0.49
RASGRP1	ENO2	1.73	1.27	0.5
RASGRP1	PRSS1	1.73	1.73	0
RASGRP1	TPM3	1.71	1.16	0.5
RASGRP1	SHMT2	1.69	1.69	0.48
RASGRP1	NAP1L4	1.69	1.69	0.47
RASGRP1	SRRM2	1.69	1.69	0.24
RASGRP1	PFDN2	1.69	1.69	0.5
RASGRP1	RANBP1	1.67	1.47	0.5
RASGRP1	TPI1	1.67	0.57	0.5
RASGRP1	CAPZB	1.64	1.64	0.47
RASGRP1	GDI2	1.64	1.64	0.48
RASGRP1	AIP	1.64	1.64	0.49
RASGRP1	IDH2	1.64	1.64	0.46
RASGRP1	TALDO1	1.64	1.64	0.47
RASGRP1	RPS29	1.64	1.37	0.5
RASGRP1	PFDN1	1.64	1.64	0.49
RASGRP1	YWHAB	1.62	1.09	0.49
RASGRP1	HSPE1	1.6	0.55	0.48
RASGRP1	RNASEH2C	1.59	1.59	0.49
RASGRP1	LCP1	1.59	1.1	0.5
RASGRP1	THYN1	1.59	1.59	0.5
RASGRP1	MYL6	1.59	1.59	0.5
RASGRP1	PARK7	1.59	0.75	0.5
RASGRP1	RPS20	1.59	1.59	0.49
RASGRP1	ATP5B	1.59	0.65	0.5
RASGRP1	NONO	1.57	1.11	0.5
RASGRP1	PSMA1	1.55	1.29	0.5
RASGRP1	MANF	1.54	1.54	0.5
RASGRP1	COX4I1	1.54	1.54	0.49
RASGRP1	DPYSL2	1.54	1.54	0.46
RASGRP1	PSME1	1.54	1.54	0.47
RASGRP1	SARNP	1.54	1.54	0.48
RASGRP1	SEP6	1.54	1.54	0.46
RASGRP1	ANXA6	1.54	1.54	0.43
RASGRP1	EEF1D	1.54	0.85	0.5

RASGRP1	DSTN	1.54	1.54	0.49
RASGRP1	EIF4A1	1.53	1.25	0.5
RASGRP1	CALR	1.53	1.34	0.5
RASGRP1	SET	1.53	1.15	0.5
RASGRP1	NEDD8	1.5	1.5	0.49
RASGRP1	LASP1	1.5	1.5	0.49
RASGRP1	TBCA	1.5	1.5	0.5
RASGRP1	EIF3A	1.5	1.5	0.35
RASGRP1	PHGDH	1.5	1.5	0.44
RASGRP1	PAK2	1.5	1.5	0.47
RASGRP1	KHDRBS1	1.5	1.5	0.45
RASGRP1	SF3B2	1.5	1.5	0.41
RASGRP1	YWHAH	1.5	1.5	0.5
RASGRP1	ATIC	1.5	1.5	0.48
RASGRP1	MCM7	1.5	1.5	0.46
RASGRP1	ATP5I	1.5	1.5	0.49
RASGRP1	PDIA4	1.5	1.5	0.47
RASGRP1	G3BP1	1.5	1.5	0.44
RASGRP1	PCNA	1.49	0.71	0.5
RASGRP1	HNRNPK	1.48	0.86	0.47
RASGRP1	PRDX6	1.47	1.09	0.5
RASGRP1	ILF3	1.47	1.16	0.5
RASGRP1	RPS21	1.46	0.69	0.5
RASGRP1	TCEB2	1.45	1.45	0.49
RASGRP1	CCDC58	1.45	1.45	0.49
RASGRP1	FLNA	1.45	1.45	0.23
RASGRP1	ACAT2	1.45	1.45	0.47
RASGRP1	COX5B	1.45	1.45	0.5
RASGRP1	PSMC3	1.45	1.45	0.47
RASGRP1	LRRC47	1.45	1.45	0.44
RASGRP1	TLN1	1.45	1.45	0.19
RASGRP1	MDH1	1.45	1.45	0.49
RASGRP1	KHSRP	1.45	1.45	0.42
RASGRP1	MYH9	1.45	1.45	0.26
RASGRP1	TFG	1.45	1.45	0.47
RASGRP1	NSFL1C	1.45	1.45	0.47
RASGRP1	GPI	1.45	1.45	0.46
RASGRP1	SMAP	1.45	1.45	0.49
RASGRP1	HN1	1.45	1.21	0.5
RASGRP1	FAM136A	1.45	1.45	0.49
RASGRP1	CDV3	1.45	1.45	0.49
RASGRP1	PSMB6	1.45	1.45	0.48
RASGRP1	CACYBP	1.45	1.45	0.49

RASGRP1	YWHAQ	1.42	1.03	0.5
RASGRP1	EIF5A2	1.4	1.4	0.48
RASGRP1	ACP1	1.4	1.4	0.48
RASGRP1	COX5A	1.4	1.4	0.49
RASGRP1	NAP1L1	1.4	1.4	0.45
RASGRP1	PPA1	1.4	1.4	0.49
RASGRP1	EIF5A	1.4	1.4	0.49
RASGRP1	LSM7	1.4	1.4	0.5
RASGRP1	GLO1	1.4	1.4	0.48
RASGRP1	FARSA	1.4	1.4	0.43
RASGRP1	MCM2	1.4	1.4	0.41
RASGRP1	STMN2	1.4	1.4	0.47
RASGRP1	PFDN6	1.4	1.4	0.49
RASGRP1	PSMB7	1.4	1.4	0.48
RASGRP1	HIST1H1A	1.39	0.68	0.26
RASGRP1	RPS12	1.39	1.22	0.49
RASGRP1	GSTP1	1.37	1	0.49
RASGRP1	MZB1	1.37	1	0.49
RASGRP1	RPS3A	1.36	1.12	0.5
RASGRP1	TCOF1	1.35	1.35	0.32
RASGRP1	WDR1	1.35	1.35	0.45
RASGRP1	DBNL	1.35	1.35	0.44
RASGRP1	GART	1.35	1.35	0.41
RASGRP1	PSAP	1.35	1.35	0.43
RASGRP1	HINT2	1.35	1.35	0.49
RASGRP1	PGD	1.35	1.35	0.42
RASGRP1	RUVBL1	1.35	1.35	0.46
RASGRP1	EIF4H	1.35	1.35	0.48
RASGRP1	TCP1	1.35	1.35	0.44
RASGRP1	TCEB1	1.35	1.35	0.47
RASGRP1	HCLS1	1.35	1.35	0.45
RASGRP1	YWHAE	1.35	1.35	0.47
RASGRP1	MAT2A	1.35	1.35	0.46
RASGRP1	H2BFS	1.35	1.35	0.35
RASGRP1	PHB	1.35	1.35	0.48
RASGRP1	PSMA7	1.35	1.35	0.47
RASGRP1	FEN1	1.35	1.35	0.47
RASGRP1	SRP14	1.35	1.35	0.48
RASGRP1	HNRNPUL1	1.35	1.35	0.34
RASGRP1	SNRPG	1.35	1.35	0.49
RASGRP1	EIF5	1.35	1.35	0.46
RASGRP1	EEF1G	1.34	1.18	0.49
RASGRP1	ENO3	1.32	0.83	0.49

RASGRP1	ANXA1	1.32	0.89	0.5
RASGRP1	ALDOA	1.32	0.64	0.49
RASGRP1	RPS16	1.32	0.95	0.48
RASGRP1	PSMA6	1.31	1.1	0.5
RASGRP1	SUMF2	1.3	1.3	0.47
RASGRP1	TUBB4A	1.3	0.68	0.47
RASGRP1	STIP1	1.3	1.3	0.45
RASGRP1	GDI1	1.3	1.3	0.43
RASGRP1	ACAT1	1.3	1.3	0.44
RASGRP1	EIF3C	1.3	1.3	0.3
RASGRP1	BTF3L4	1.3	1.3	0.48
RASGRP1	CCDC124	1.3	1.3	0.47
RASGRP1	RANBP3	1.3	1.3	0.42
RASGRP1	PFKP	1.3	1.3	0.34
RASGRP1	EDF1	1.3	1.3	0.48
RASGRP1	SRSF5	1.3	1.3	0.48
RASGRP1	HSPH1	1.3	1.3	0.33
RASGRP1	ARPP19	1.3	1.3	0.48
RASGRP1	C9orf78	1.3	1.3	0.37
RASGRP1	MAP4	1.3	1.3	0.44
RASGRP1	CHORDC1	1.3	1.3	0.47
RASGRP1	CRIP1	1.3	1.3	0.49
RASGRP1	CANX	1.3	1.3	0.39
RASGRP1	ACTR3	1.3	1.3	0.43
RASGRP1	VBP1	1.3	1.3	0.48
RASGRP1	USP14	1.3	1.3	0.44
RASGRP1	RPSA	1.29	1.13	0.49
RASGRP1	RPS5	1.28	1.01	0.5
RASGRP1	DDX41	1.27	1.06	0.49
RASGRP1	YWHAZ	1.27	0.63	0.47
RASGRP1	RPS28	1.27	0.66	0.49
RASGRP1	SNRPGP15	1.27	1.06	0.49
RASGRP1	LDHB	1.27	0.8	0.44
RASGRP1	ANP32E	1.25	1.25	0.44
RASGRP1	ANP32B	1.25	1.25	0.45
RASGRP1	APRT	1.25	1.25	0.47
RASGRP1	ANP32A	1.25	1.25	0.46
RASGRP1	GANAB	1.25	1.25	0.29
RASGRP1	AK2	1.25	1.25	0.47
RASGRP1	PSMC1	1.25	1.25	0.44
RASGRP1	COA6	1.25	1.25	0.48
RASGRP1	REXO2	1.25	1.25	0.48
RASGRP1	CBFB	1.25	1.25	0.48

RASGRP1	EIF4B	1.25	1.25	0.36
RASGRP1	MTHFD2	1.25	1.25	0.45
RASGRP1	ADSS	1.25	1.25	0.4
RASGRP1	IFI16	1.25	1.25	0.33
RASGRP1	RTCB	1.25	1.25	0.44
RASGRP1	ST13P5	1.25	1.25	0.33
RASGRP1	MAT2B	1.25	1.25	0.45
RASGRP1	HN1L	1.25	1.25	0.47
RASGRP1	TPD52	1.25	1.25	0.47
RASGRP1	SF1	1.25	1.25	0.36
RASGRP1	SRSF6	1.25	1.25	0.44
RASGRP1	SEP6	1.25	1.25	0.32
RASGRP1	PSMA2	1.25	1.25	0.47
RASGRP1	TIMM9	1.25	1.25	0.48
RASGRP1	DPP3	1.25	1.25	0.36
RASGRP1	ANXA5	1.25	1.25	0.46
RASGRP1	MCM3	1.25	1.25	0.32
RASGRP1	EIF1AX	1.25	1.25	0.48
RASGRP1	EIF1AY	1.25	1.25	0.48
RASGRP1	PFAS	1.25	1.25	0.23
RASGRP1	CIAPIN1	1.25	1.25	0.46
RASGRP1	PPP2CA	1.25	1.25	0.45
RASGRP1	PPP2CB	1.25	1.25	0.45
RASGRP1	CHTF8	1.25	1.25	0.48
RASGRP1	ST13	1.25	1.25	0.4
RASGRP1	CAP1	1.25	1.25	0.42
RASGRP1	POLDIP3	1.25	1.25	0.39
RASGRP1	HIST1H1E	1.25	0.65	0.22
RASGRP1	THRAP3	1.25	1.25	0.3
RASGRP1	HCFC1	1.25	1.25	0.16
RASGRP1	PRPF19	1.25	1.25	0.38
RASGRP1	DNASE2	1.25	1.25	0.45
RASGRP1	LRRC59	1.25	1.25	0.45
RASGRP1	G3BP2	1.25	1.25	0.42
RASGRP1	MIF	1.24	1.09	0.48
RASGRP1	RAN	1.24	1.09	0.49
RASGRP1	HAGH	1.24	0.68	0.49
RASGRP1	RPS10	1.24	0.84	0.48
RASGRP1	RPL26	1.23	0.67	0.48
RASGRP1	TAGLN2	1.21	0.98	0.47
RASGRP1	CFL1	1.21	0.46	0.48
RASGRP1	HSPA4	1.2	1.2	0.29
RASGRP1	ZC3H15	1.2	1.2	0.38

RASGRP1	LARP1	1.2	1.2	0.25
RASGRP1	PCBP1	1.2	1.05	0.47
RASGRP1	RPS17L	1.2	1.2	0.46
RASGRP1	TIMM10	1.2	1.2	0.47
RASGRP1	PDLIM1	1.2	1.2	0.44
RASGRP1	FLNB	1.2	1.2	0.13
RASGRP1	SMS	1.2	1.2	0.41
RASGRP1	RAP1B	1.2	1.2	0.45
RASGRP1	RAP1A	1.2	1.2	0.45
RASGRP1	DCTN6	1.2	1.2	0.46
RASGRP1	CDC37	1.2	1.2	0.4
RASGRP1	PBK	1.2	1.2	0.43
RASGRP1	RBM3	1.2	1.2	0.48
RASGRP1	PSMD7	1.2	1.2	0.43
RASGRP1	IMPDH2	1.2	1.2	0.36
RASGRP1	RPL22	1.2	1.2	0.47
RASGRP1	PSMC2	1.2	1.2	0.45
RASGRP1	CLTA	1.2	1.2	0.45
RASGRP1	PSMD4	1.2	1.2	0.42
RASGRP1	EIF3G	1.2	1.2	0.43
RASGRP1	THOC3	1.2	1.2	0.4
RASGRP1	GTF2F1	1.2	1.2	0.35
RASGRP1	COPS3	1.2	1.2	0.37
RASGRP1	RNASEH2A	1.2	1.2	0.43
RASGRP1	RPS26P11	1.2	1.2	0.28
RASGRP1	RCSD1	1.2	1.2	0.41
RASGRP1	ARHGDI1	1.2	0.87	0.48
RASGRP1	LSM4	1.2	1.2	0.47
RASGRP1	TXNDC5	1.2	1.2	0.42
RASGRP1	PPP1R2	1.2	1.2	0.47
RASGRP1	EPB41	1.2	1.2	0.28
RASGRP1	MRPS28	1.2	1.2	0.45
RASGRP1	BTF3	1.2	1.2	0.47
RASGRP1	RPS7	1.2	0.73	0.46
RASGRP1	RBM8A	1.2	1.2	0.47
RASGRP1	RPS26	1.2	1.2	0.48
RASGRP1	PPIH	1.2	1.2	0.46
RASGRP1	PPIF	1.2	1.2	0.46
RASGRP1	PNPT1	1.2	1.2	0.31
RASGRP1	SUCLG1	1.2	1.2	0.42
RASGRP1	NDUFS8	1.2	1.2	0.46
RASGRP1	DHX15	1.2	1.2	0.29
RASGRP1	APEH	1.2	1.2	0.34

RASGRP1	HIST1H1B	1.2	0.7	0.5
RASGRP1	RBM26	1.2	1.2	0.27
RASGRP1	RBM14	1.2	1.2	0.41
RASGRP1	RCC2	1.2	1.2	0.37
RASGRP1	PSMB1	1.2	1.2	0.43
RASGRP1	PPP1R2P3	1.2	1.2	0.35
RASGRP1	SLC9A3R1	1.2	1.2	0.44
RASGRP1	WBP11	1.2	1.2	0.3
RASGRP1	LDHA	1.19	0.62	0.45
RASGRP1	XRCC5	1.18	0.78	0.46
RASGRP1	FAU	1.18	0.77	0.45
RASGRP1	DDX39B	1.17	0.88	0.46
RASGRP1	DDX39A	1.17	0.88	0.45
RASGRP1	XRCC6	1.17	0.72	0.46
RASGRP1	ATP5O	1.17	0.98	0.47
RASGRP1	MFAP1	1.15	1.15	0.33
RASGRP1	CNDP2	1.15	1.15	0.31
RASGRP1	SNAP29	1.15	1.15	0.4
RASGRP1	CSNK2B	1.15	1.15	0.45
RASGRP1	PSMD9	1.15	1.15	0.43
RASGRP1	DDB1	1.15	1.15	0.22
RASGRP1	ENSA	1.15	1.15	0.46
RASGRP1	CPOX	1.15	1.15	0.32
RASGRP1	EIF6	1.15	1.15	0.42
RASGRP1	AK6	1.15	1.15	0.28
RASGRP1	EWSR1	1.15	1.15	0.28
RASGRP1	NUDT21	1.15	1.15	0.43
RASGRP1	TOR1AIP1	1.15	1.15	0.31
RASGRP1	FKBP3	1.15	1.15	0.43
RASGRP1	FUBP1	1.15	1.15	0.26
RASGRP1	COX17	1.15	1.15	0.48
RASGRP1	CNBP	1.15	1.15	0.44
RASGRP1	HYPK	1.15	1.15	0.48
RASGRP1	ETFB	1.15	1.15	0.42
RASGRP1	CDK1	1.15	1.15	0.44
RASGRP1	SLIRP	1.15	1.15	0.47
RASGRP1	CSTF2	1.15	1.15	0.31
RASGRP1	TARS	1.15	1.15	0.27
RASGRP1	CAPRIN1	1.15	1.15	0.28
RASGRP1	MAPRE1	1.15	1.15	0.43
RASGRP1	MARCKSL1	1.15	1.15	0.43
RASGRP1	ANXA11	1.15	1.15	0.31
RASGRP1	HSPA4L	1.15	1.15	0.26

RASGRP1	ACTN1	1.15	1.15	0.23
RASGRP1	RPA3	1.15	1.15	0.46
RASGRP1	PSAT1	1.15	1.15	0.42
RASGRP1	MGMT	1.15	1.15	0.44
RASGRP1	EIF4G1	1.15	1.15	0.15
RASGRP1	CTPS1	1.15	1.15	0.28
RASGRP1	TXNL1	1.15	1.15	0.41
RASGRP1	NPM3	1.15	1.15	0.45
RASGRP1	SNRPE	1.15	1.15	0.47
RASGRP1	RPL17	1.13	0.71	0.43
RASGRP1	ENO1	1.11	0.42	0.48
RASGRP1	EEF1B2	1.11	0.75	0.45
RASGRP1	NELFE	1.1	1.1	0.31
RASGRP1	GTF2F2	1.1	1.1	0.37
RASGRP1	DDX46	1.1	1.1	0.14
RASGRP1	SRSF11	1.1	1.1	0.27
RASGRP1	FDX1	1.1	1.1	0.42
RASGRP1	HMG5	1.1	1.1	0.34
RASGRP1	FKBP2	1.1	1.1	0.44
RASGRP1	LSM14A	1.1	1.1	0.29
RASGRP1	TBCB	1.1	1.1	0.35
RASGRP1	PSMG4	1.1	1.1	0.44
RASGRP1	TIMM13	1.1	1.1	0.45
RASGRP1	PDAP1	1.1	1.1	0.41
RASGRP1	DNAJC8	1.1	1.1	0.37
RASGRP1	DCTN2	1.1	1.1	0.28
RASGRP1	PHAX	1.1	1.1	0.28
RASGRP1	PSMD2	1.1	1.1	0.19
RASGRP1	EIF2S3	1.1	1.1	0.26
RASGRP1	DYNLRB2	1.1	1.1	0.45
RASGRP1	NOLC1	1.1	1.1	0.2
RASGRP1	EIF1AD	1.1	1.1	0.43
RASGRP1	PNP	1.1	1.1	0.33
RASGRP1	HDAC1	1.1	1.1	0.27
RASGRP1	COTL1	1.1	1.1	0.43
RASGRP1	POLR2C	1.1	1.1	0.34
RASGRP1	FIP1L1	1.1	1.1	0.23
RASGRP1	NTPCR	1.1	1.1	0.41
RASGRP1	VAT1	1.1	1.1	0.27
RASGRP1	DYNLRB1	1.1	1.1	0.45
RASGRP1	SNX3	1.1	1.1	0.44
RASGRP1	NASP	1.1	1.1	0.23
RASGRP1	PPP5C	1.1	1.1	0.24

RASGRP1	NUDCD2	1.1	1.1	0.42
RASGRP1	APIP	1.1	1.1	0.37
RASGRP1	CIRBP	1.1	1.1	0.42
RASGRP1	ATP6V1F	1.1	1.1	0.44
RASGRP1	RAB10	1.1	1.1	0.42
RASGRP1	CLIC1	1.1	1.1	0.36
RASGRP1	CDK6	1.1	1.1	0.37
RASGRP1	SF3B5	1.1	1.1	0.46
RASGRP1	PRDX3	1.1	1.1	0.39
RASGRP1	MRPS14	1.1	1.1	0.44
RASGRP1	BID	1.1	1.1	0.43
RASGRP1	PSME3	1.1	1.1	0.36
RASGRP1	TUFM	1.1	1.1	0.26
RASGRP1	MLEC	1.1	1.1	0.35
RASGRP1	PPP4R2	1.1	1.1	0.27
RASGRP1	UQCRH	1.1	1.1	0.46
RASGRP1	TOMM6	1.1	1.1	0.45
RASGRP1	RHOA	1.1	1.1	0.4
RASGRP1	PPP1R8	1.1	1.1	0.38
RASGRP1	HDAC2	1.1	1.1	0.29
RASGRP1	ZNF207	1.1	1.1	0.26
RASGRP1	ACOT7	1.1	1.1	0.3
RASGRP1	ARCN1	1.1	1.1	0.27
RASGRP1	PPM1A	1.1	1.1	0.28
RASGRP1	PARP1	1.1	1.1	0.13
RASGRP1	PHF5A	1.1	1.1	0.43
RASGRP1	UBE2S	1.1	1.1	0.39
RASGRP1	BAG2	1.1	1.1	0.4
RASGRP1	UBA1	1.1	1.1	0.15
RASGRP1	HP1BP3	1.1	1.1	0.25
RASGRP1	ADI1	1.1	1.1	0.42
RASGRP1	SEP7	1.1	1.1	0.24
RASGRP1	BCL7A	1.1	1.1	0.39
RASGRP1	PPIE	1.1	1.1	0.34
RASGRP1	PPID	1.1	1.1	0.31
RASGRP1	AIMP1	1.1	1.1	0.33
RASGRP1	ELF1	1.1	1.1	0.24
RASGRP1	SOD2	1.1	1.1	0.4
RASGRP1	APEX1	1.1	1.1	0.29
RASGRP1	ALDOC	1.1	1.1	0.29
RASGRP1	ZC3HAV1	1.1	1.1	0.2
RASGRP1	BPNT1	1.1	1.1	0.34
RASGRP1	MEA1	1.1	1.1	0.4

RASGRP1	CFDP1	1.1	1.1	0.37
RASGRP1	AHCY	1.1	1.1	0.29
RASGRP1	NMT1	1.1	1.1	0.24
RASGRP1	LYRM7	1.1	1.1	0.43
RASGRP1	PITRM1	1.1	1.1	0.15
RASGRP1	CSNK2A3	1.1	1.1	0.22
RASGRP1	CSNK2A1	1.1	1.1	0.32
RASGRP1	FDPS	1.1	1.1	0.3
RASGRP1	CD2AP	1.1	1.1	0.22
RASGRP1	NIT2	1.1	1.1	0.35
RASGRP1	PCBP2	1.1	0.97	0.45
RASGRP1	NACA	1.1	0.97	0.42
RASGRP1	RAB5C	1.1	1.1	0.4
RASGRP1	HADH	1.1	1.1	0.35
RASGRP1	CD2BP2	1.1	1.1	0.31
RASGRP1	RPL35A	1.1	1.1	0.44
RASGRP1	SNRPB	1.1	1.1	0.38
RASGRP1	TAL1	1.1	1.1	0.32
RASGRP1	SNRPA	1.1	1.1	0.35
RASGRP1	MRPS36	1.1	1.1	0.45
RASGRP1	SNRNP70	1.1	1.1	0.27
RASGRP1	POMP	1.1	1.1	0.42
RASGRP1	GLUD1	1.1	1.1	0.25
RASGRP1	BCL7B	1.1	1.1	0.4
RASGRP1	EPB41L2	1.1	1.1	0.16
RASGRP1	PRPS2	1.1	1.1	0.33
RASGRP1	CMPK1	1.1	1.1	0.39
RASGRP1	GCLC	1.1	1.1	0.24
RASGRP1	ESD	1.1	1.1	0.35
RASGRP1	SNRPN	1.1	1.1	0.36
RASGRP1	CAT	1.09	0.84	0.45
RASGRP1	DDX17	1.08	0.71	0.44
RASGRP1	CORO1A	1.07	0.48	0.42
RASGRP1	TUBB2A	1.06	0.68	0.37
RASGRP1	RPS4X	1.06	0.65	0.39
RASGRP1	ATP6V1B1	1.05	1.05	0
RASGRP1	SMARCC1	1.05	1.05	0
RASGRP1	C21orf33	1.05	1.05	0
RASGRP1	B2M	1.05	1.05	0
RASGRP1	MZT2A	1.05	1.05	0
RASGRP1	DBN1	1.05	1.05	0
RASGRP1	ZNF706	1.05	1.05	0
RASGRP1	DRAP1	1.05	1.05	0

RASGRP1	OSGEP	1.05	1.05	0
RASGRP1	DIAPH1	1.05	1.05	0
RASGRP1	FKBP4	1.05	1.05	0
RASGRP1	GRAP2	1.05	1.05	0
RASGRP1	PDRG1	1.05	1.05	0
RASGRP1	EPDR1	1.05	1.05	0
RASGRP1	MYH10	1.05	1.05	0
RASGRP1	ACO2	1.05	1.05	0
RASGRP1	MRPL53	1.05	1.05	0
RASGRP1	NDE1	1.05	1.05	0
RASGRP1	SNW1	1.05	1.05	0
RASGRP1	USP39	1.05	1.05	0
RASGRP1	CYB5B	1.05	1.05	0
RASGRP1	ACTR1B	1.05	1.05	0
RASGRP1	ACTR1A	1.05	1.05	0
RASGRP1	DHRS4	1.05	1.05	0
RASGRP1	DPY30	1.05	1.05	0
RASGRP1	ATP6V1B2	1.05	1.05	0
RASGRP1	H1FX	1.05	1.05	0
RASGRP1	UQCRC2	1.05	1.05	0
RASGRP1	XPNPEP1	1.05	1.05	0
RASGRP1	FTSJ3	1.05	1.05	0
RASGRP1	SDF2L1	1.05	1.05	0
RASGRP1	TCF7	1.05	1.05	0
RASGRP1	TPT1	1.05	1.05	0
RASGRP1	MLLT11	1.05	1.05	0
RASGRP1	C1orf52	1.05	1.05	0
RASGRP1	LARS	1.05	1.05	0
RASGRP1	MCM4	1.05	1.05	0
RASGRP1	PROSC	1.05	1.05	0
RASGRP1	TPM4	1.05	0.71	0.37
RASGRP1	RPLP1	1.05	0.58	0.45
RASGRP1	ADSL	1.05	1.05	0
RASGRP1	WDR82	1.05	1.05	0
RASGRP1	RAB14	1.05	1.05	0
RASGRP1	RAB1C	1.05	1.05	0
RASGRP1	UNG	1.05	1.05	0
RASGRP1	BOLA2B	1.05	1.05	0
RASGRP1	OGFR	1.05	1.05	0
RASGRP1	RAB1B	1.05	1.05	0
RASGRP1	BDH2	1.05	1.05	0
RASGRP1	UBAP2L	1.05	1.05	0
RASGRP1	WIBG	1.05	1.05	0

RASGRP1	MYEOV2	1.05	1.05	0
RASGRP1	NT5C	1.05	1.05	0
RASGRP1	ADD1	1.05	1.05	0
RASGRP1	DNAJA1	1.05	1.05	0
RASGRP1	API5	1.05	1.05	0
RASGRP1	TMEM109	1.05	1.05	0
RASGRP1	LSM14B	1.05	1.05	0
RASGRP1	ARMC1	1.05	1.05	0
RASGRP1	SAE1	1.05	1.05	0
RASGRP1	SLC3A2	1.05	1.05	0
RASGRP1	ZPR1	1.05	1.05	0
RASGRP1	POLE3	1.05	1.05	0
RASGRP1	ATP6V1G1	1.05	1.05	0
RASGRP1	TP53RK	1.05	1.05	0
RASGRP1	DUS3L	1.05	1.05	0
RASGRP1	SRI	1.05	1.05	0
RASGRP1	RRM2	1.05	1.05	0
RASGRP1	EFHD2	1.05	1.05	0
RASGRP1	GNS	1.05	1.05	0
RASGRP1	MAGOHB	1.05	1.05	0
RASGRP1	PIN1	1.05	1.05	0
RASGRP1	PRRC2A	1.05	1.05	0
RASGRP1	LAMTOR5	1.05	1.05	0
RASGRP1	CPNE1	1.05	1.05	0
RASGRP1	ZRANB2	1.05	1.05	0
RASGRP1	GOT1	1.05	1.05	0
RASGRP1	MAGOH	1.05	1.05	0
RASGRP1	UPF1	1.05	1.05	0
RASGRP1	GLUL	1.05	1.05	0
RASGRP1	COPA	1.05	1.05	0
RASGRP1	MZT2B	1.05	1.05	0
RASGRP1	SSNA1	1.05	1.05	0
RASGRP1	NPC2	1.05	1.05	0
RASGRP1	HMBS	1.05	1.05	0
RASGRP1	RAB7A	1.05	1.05	0
RASGRP1	ARPC5	1.05	1.05	0
RASGRP1	C1orf122	1.05	1.05	0
RASGRP1	DNAJB1	1.05	1.05	0
RASGRP1	HNRNPF	1.05	0.92	0.41
RASGRP1	CLPP	1.05	1.05	0
RASGRP1	IGBP1	1.05	1.05	0
RASGRP1	HADHA	1.05	1.05	0
RASGRP1	MAP7D1	1.05	1.05	0

RASGRP1	NDEL1	1.05	1.05	0
RASGRP1	SNRPC	1.05	1.05	0
RASGRP1	RCC1	1.05	1.05	0
RASGRP1	RUVBL2	1.05	1.05	0
RASGRP1	RPL32	1.05	1.05	0
RASGRP1	PDCD6IP	1.05	1.05	0
RASGRP1	ACTA1	1.04	0.48	0.48
RASGRP1	ACTC1	1.04	0.48	0.48
RASGRP1	HSP90B1	1.03	0.65	0.43
RASGRP1	NIPSNAP1	1.03	0.86	0.42
RASGRP1	RPS25	1.03	0.55	0.44
RASGRP1	RPS14	1.03	0.58	0.36
RASGRP1	SNRPD3	1.02	0.82	0.35
RASGRP1	GAPDH	1.02	0.56	0.47
RASGRP1	PRDX2	1.01	0.72	0.33
RASGRP1	RPS24	1.01	0.79	0.36
RASGRP1	HIST1H1D	1	0.58	0.38
RASGRP1	HIST1H1C	1	0.58	0.38
RASGRP1	CAPZA1	0.99	0.83	0
RASGRP1	HNRNPH1	0.99	0.74	0.38
RASGRP1	TRAP1	0.99	0.83	0
RASGRP1	RBMX	0.98	0.61	0.33
RASGRP1	RPL21	0.98	0.67	0.29
RASGRP1	ILF2	0.97	0.76	0
RASGRP1	RPS17	0.96	0.57	0.3
RASGRP1	RPS3	0.94	0.6	0.18
RASGRP1	RPS11	0.94	0.64	0.24
RASGRP1	TUBB6	0.93	0.47	0.33
RASGRP1	HNRNPR	0.9	0.56	0.27
RASGRP1	HSPA5	0.89	0.45	0.46
RASGRP1	NCL	0.89	0.38	0.37
RASGRP1	SPTBN1	0.89	0.51	0.22
RASGRP1	ALYREF	0.88	0.45	0.25
RASGRP1	RPL24	0.88	0.52	0.27
RASGRP1	EEF1A2	0.87	0.51	0.41
RASGRP1	RPL10L	0.87	0.59	0.09
RASGRP1	RPL10	0.87	0.59	0.07
RASGRP1	HNRNPD	0.87	0.58	0.12
RASGRP1	DDX21	0.85	0.55	0.24
RASGRP1	RPL3	0.85	0.43	0.25
RASGRP1	RPL23A	0.84	0.52	0.08
RASGRP1	SNAP23	0.84	0.5	0
RASGRP1	PFN1	0.84	0.5	0.23

RASGRP1	RPL31	0.83	0.45	0.2
RASGRP1	RPL15	0.82	0.39	0.3
RASGRP1	ACACA	0.82	0.24	0.09
RASGRP1	DDX5	0.81	0.53	0.22
RASGRP1	PRDX1	0.8	0.53	0.18
RASGRP1	ARHGDIB	0.79	0.38	0.3
RASGRP1	RPS6	0.79	0.35	0.25
RASGRP1	RPL36A	0.77	0.39	0.13
RASGRP1	ACTG1	0.77	0.41	0.46
RASGRP1	RPL36AL	0.77	0.39	0.18
RASGRP1	YBX3	0.76	0.5	0
RASGRP1	RPL11	0.76	0.45	0.07
RASGRP1	VIM	0.74	0.26	0.32
RASGRP1	RPS18	0.72	0.31	0.14
RASGRP1	RPL7A	0.71	0.32	0.13
RASGRP1	GRPEL1	0.7	0.28	0.35
RASGRP1	SPTAN1	0.69	0.33	0
RASGRP1	HNRNPA1	0.67	0.41	0.18
RASGRP1	EEF1A1	0.66	0.32	0.17
RASGRP1	ACTB	0.63	0.32	0.44
RASGRP1	HSP90AB1	0.63	0.4	0.27
RASGRP1	HNRNPA2B1	0.61	0.36	0.11
RASGRP1	RPL7	0.61	0.34	0.08
RASGRP1	RPL18	0.6	0.31	0.13
RASGRP1	RPL13	0.56	0.28	0.07
RASGRP1	RPL6	0.56	0.23	0.08
RASGRP1	RPL12	0.54	0.24	0.06
RASGRP1	HSP90AA1	0.54	0.35	0.17
RASGRP1	TUBA1C	0.54	0.36	0.06
RASGRP1	YBX1	0.51	0.27	0
RASGRP1	HNRNPU	0.51	0.28	0.02
RASGRP1	RPL4	0.5	0.2	0.08
RASGRP1	LMNB1	0.49	0.18	0.03
RASGRP1	HIST3H2A	0.48	0.34	0
RASGRP1	HSPA1B	0.48	0.29	0.03
RASGRP1	HSPA1A	0.48	0.29	0.04
RASGRP1	HIST1H2AC	0.48	0.34	0
RASGRP1	HIST1H2AB	0.48	0.34	0
RASGRP1	HIST2H2AC	0.46	0.33	0.01
RASGRP1	HIST2H2AA3	0.46	0.33	0.01
RASGRP1	H2AFJ	0.46	0.33	0.01
RASGRP1	HIST1H2AM	0.46	0.33	0
RASGRP1	HIST1H2AH	0.46	0.33	0.01

RASGRP1	HIST1H2AJ	0.46	0.33	0.01
RASGRP1	HIST1H2AD	0.46	0.33	0.01
RASGRP1	HIST1H2BB	0.43	0.29	0.01
RASGRP1	HIST1H2BO	0.43	0.29	0.01
RASGRP1	HIST1H2BJ	0.43	0.29	0.01
RASGRP1	HIST2H2BE	0.43	0.29	0.01
RASGRP1	HIST2H2BF	0.39	0.27	0
RASGRP1	HIST1H2BD	0.39	0.27	0
RASGRP1	HIST1H2BF	0.39	0.27	0.01
RASGRP1	HIST1H2BL	0.39	0.27	0
RASGRP1	HIST1H2BM	0.39	0.27	0
RASGRP1	HIST1H2BN	0.39	0.27	0
RASGRP1	HIST1H2BH	0.39	0.27	0
RASGRP1	HIST1H2BK	0.39	0.27	0
RASGRP1	ATXN2L	0.18	0.08	0
RASGRP1	PCCB	0.17	0.06	0.02
RASGRP1	KRT1	0.12	0.08	0
RASGRP1	KRT9	0.12	0.08	0
RASGRP1	KRT5	0.11	0.07	0
RASGRP1	KRT2	0.09	0.05	0
RASGRP1	KRT10	0.02	0.01	0
RASGRP1	SNRPF	0	0	0
RASGRP1	SH	0	0	0
RASGRP1	Trypsin	0	0	0
RASGRP1	GFP	0	0	0

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