

Supplementary Information

Pyruvate Kinase M2 Activates mTORC1 by Phosphorylating AKT1S1

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Supplementary Data

Supplementary Figures S1-S6

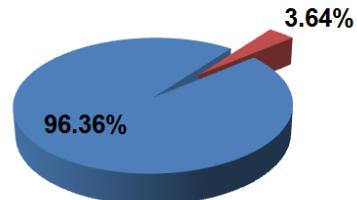
Figure S1

A

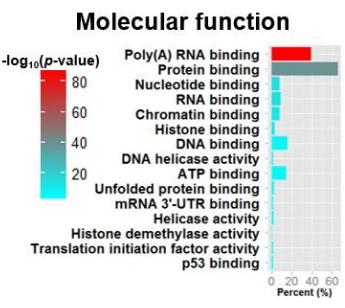
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.....DS...	14.07	46	277	45642	1048208	3.81
.....S.S...	7.91	43	159	96036	901183	2.49
.....SD.D...	32	42	436	2910	1093935	35.93
.....R.R...	12.24	42	231	5843	1002569	3.1
.....DS.E...	43.66	40	851	592	1108364	88
.....S.DE...	32	40	567	3288	1095554	23.59
.....S..DE...	32	35	527	3304	1096264	22.04
.....SEEE...	40.04	32	759	1000	1106980	46.67
.....R.G.S...	47.35	32	699	1162	1107171	50.7
.....ES...	26.59	30	452	4159	1092962	46.02
.....D...S...	8.55	30	189	42929	944112	3.49
.....SD.E.D...	38.66	29	811	381	1107772	103.97
.....R.RG.S...	41.04	28	627	1209	1101926	40.7
.....DS.E...	27.45	26	462	3428	1088603	17.98
.....S...S...	25	25	302	6580	1058500	12.8
.....S...SP...	19.24	24	326	10068	1065126	7.77
.....SD.EK...	41.14	23	782	411	1107391	79.25
.....E..SE...	24.18	23	394	4539	1082485	13.92
.....S...S...E...	22.09	23	349	7317	1072443	9.66
R.....S...	7	23	116	44720	803147	3.56
.....S...S...	20.13	22	371	55206	1079369	11.1
.....SE.E.E...	38.22	21	648	580	1102505	61.6
.....R.RG.S...	55.06	20	668	155	1102660	212.99
.....S.T...	8.74	43	179	63146	705184	2.68
.....T...E...	7.25	28	136	42009	642038	3.15

B

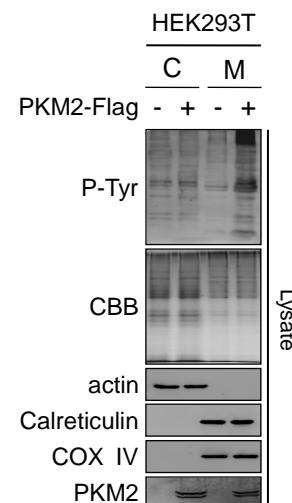
■ Disordered ■ Ordered



C



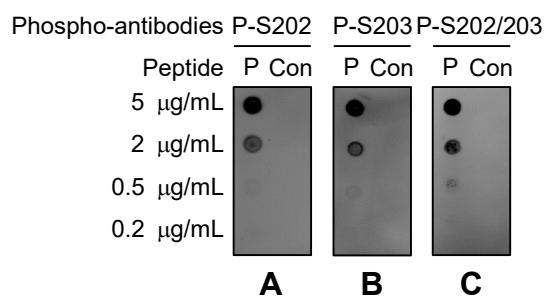
D



Supplementary Figure S1, Bioinformatics Analysis of Substrates of PKM2

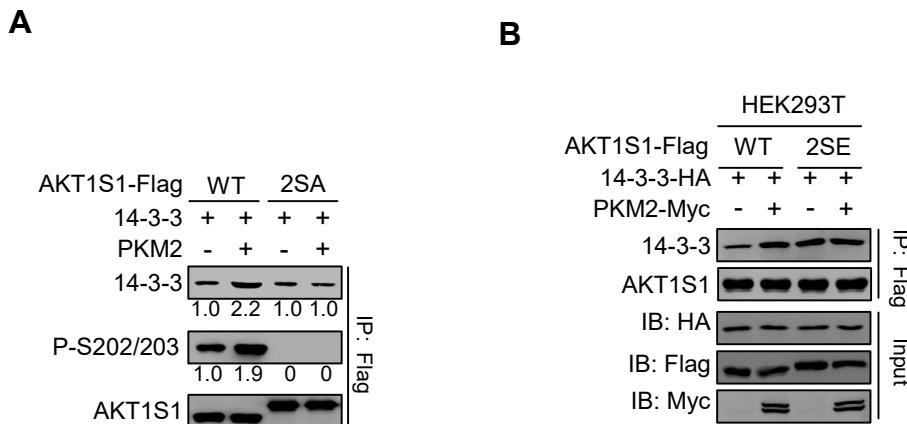
(Related to Figure 2). Bioinformatics analysis results are shown. **A**, Motif-x analysis was employed to identify consensus sequences in substrates (S and T) of PKM2. **B**, The distribution of phosphorylation sites of PKM2 in ordered and disordered regions of proteins. **C**, KEGG molecular functions enrichment of substrates of PKM2. **D**, PKM2 increases membrane-bound P-Tyr. Flag-tagged PKM2 was overexpressed in HEK293T cells. The levels of P-Tyr in cytosolic (C) and membrane (M) fractions were determined.

Figure S2



Supplementary Figure S2, Validation of P-202, P-203 and P-202/203 antibodies (Related to Figure 3). Dot blot assays were carried out to determine the specificities of AKT1S1 P-S202 (**A**), P-S203 (**B**) and P-S202/203 (**C**) antibodies. Anti-sera were used at 1:1000 dilutions. Synthetic phosphor-serine containing peptides (P) that was used in making each antibody and the control peptides (Con) that have the same amino acids sequence but devoid of phosphor-serines were fixed onto nitrocellulose membrane at the concentrations as indicated, the signal of antibodies reactive to each antigen and to each control peptide was detected by chemiluminescence on Typhoon (GE).

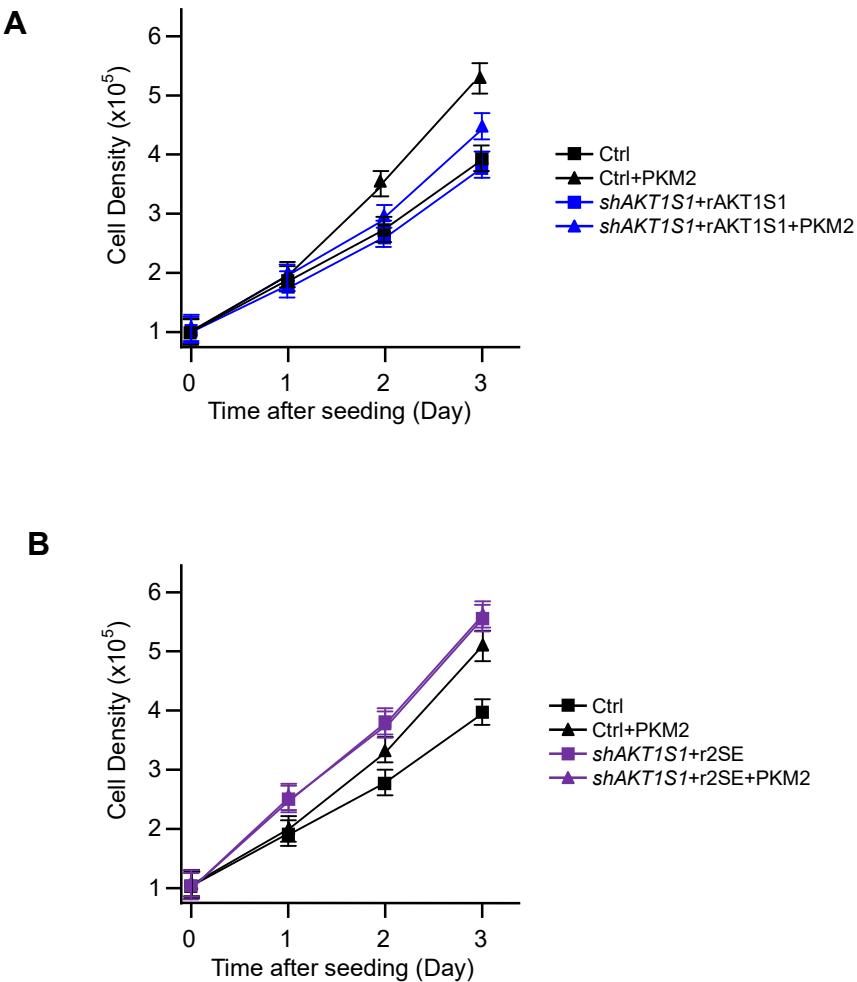
Figure S3



Supplementary Figure S3, S202/203 phosphorylation increases AKT1S1-14-3-3 interaction (Related to Figure 4).

A, Purified 14-3-3 was incubated with AKT1S1 or 2SA with or without PKM2 *in vitro*. After reaction, the levels of P-S202/203 of Flag beads purified AKT1S1 or 2SA and the amount of 14-3-3 pulled down by purified AKT1S1 or 2SA was determined (western blot) and quantified (numeric). **B**, 14-3-3 was co-expressed with either AKT1S1 or 2SE in HEK293T cells, the amount of 14-3-3 co-immunoprecipitated with either AKT1S1 or 2SE with expressing PKM2 or without expressing PKM2 were determined.

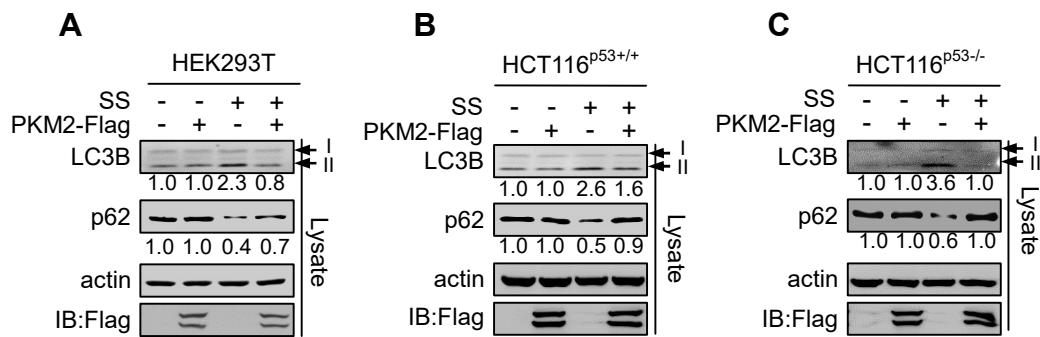
Figure S4



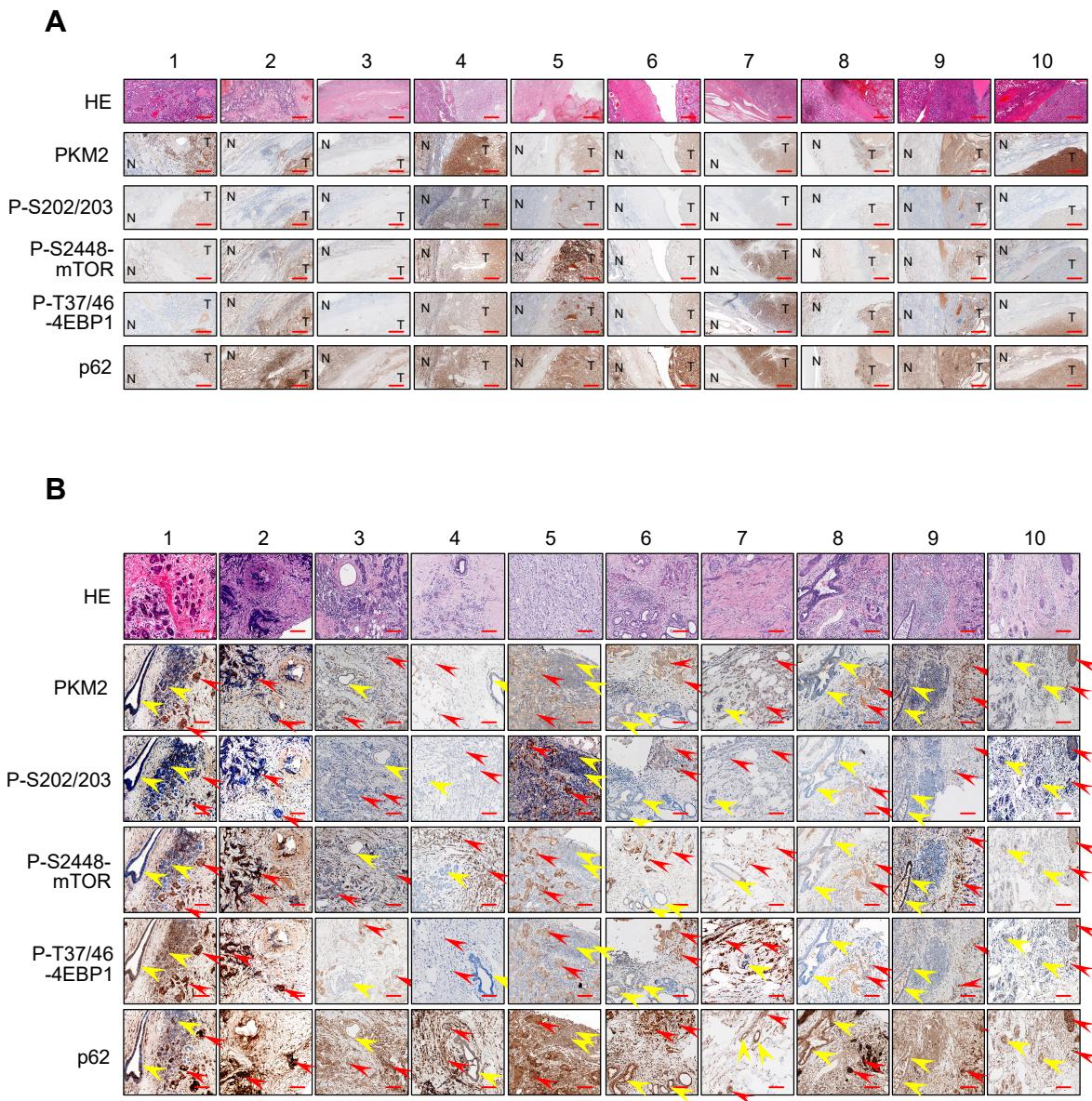
Supplementary Figure S4, PKM2 activates mTORC1 (Related to Figure 5).

Growth curves of AKT1S1 knockdown HEK293T cells that stably overexpressing shRNA resistant AKT1S1 (rAKT1S1) (**A**) or 2SE (r2SE) (**B**) were detected with and without overexpressing PKM2. Growth curves of HEK293T cells with and without overexpressing PKM2 were determined in each experiment as controls.

Figure S5



Supplementary Figure S5, PKM2 overexpression prevented serum starvation induced autophagy in HEK293T, HCT116^{P53+/+} and HCT116^{P53-/-} cells (Related to Figure 6). Cells were cultured under either normal DMEM or serum starvation (SS) conditions. The levels of LC3B and p62 in HEK293T cells and HEK293T cells expressing PKM2 (**A**), in HCT116^{P53+/+} cells and HCT116^{P53+/+} cells expressing PKM2 (**B**), in HCT116^{P53-/-} cells and HCT116^{P53-/-} cells expressing PKM2 (**C**), were determined and quantified.



Supplementary Figure S6, IHC analysis of RCC and breast cancer samples

(Related to Figure 7). PKM2, P-S202/203, P-S2448-mTOR, p-T37/46-4EBP1 and p62 levels of the same patient were detected in RCC (A) and breast cancer (B) tissues and adjacent normal tissues. For RCC samples, normal and tumor tissues are marked by N and T, respectively. For breast cancer, tumor and normal tissues are marked by red and yellow arrows, respectively. Pathologic results were confirmed by experienced pathologists. Bar scales were 100 μ m.

Supplemental Table S1

Table S1: Identified PKM2 Substrates

UniProt Acces	Gene name	Position	Code	Peptide Sequence	Fold Change
1 O75643	SNRNP200	225	S	EEGDEDVYGEVREEAS DDDMEGDEAVVRCTL	3166.18
2 Q96EV2	RBM33	205	S	GGMETLELQKDIKEES DEEEEDEESGRRLF	2576.20
3 O95400	CD2BP2	49	S	SGGPGSRFKGKHSLDS DEEEDDDGGSSKYD	1828.91
4 Q8WVC0	LEO1	197	S	DEKMQNTDDEERPQLS SDDERQQLSEEKANS	1769.91
5 Q96ST2	IWS1	196	S	QISDSEEEP PRHQAS DSEN EPPKPRMSDS	1769.83
6 P25205	MCM3	711	S	QPDAKGDSYDPYDF S DTEEE MPQVHTPKTA	1415.93
7 Q96ST2	IWS1	235	S	QVSDSEEEP PRHQAS DSEN EELPKPRISDS	1356.93
8 Q96ST2	IWS1	198	S	SDSEEEP PRHQAS DSEN EELPKPRISDS	1317.60
9 Q96ST2	IWS1	237	S	SDSEEEP PRHQAS DSEN EELPKPRISDS	1254.30
10 Q12872	SFSWAP	870	S	RSRSRTKSKARSQSV S PSKQAAPRPAAPAAH	1163.40
11 Q13435	SF3B2	436	S	KKGFE EEHKDSDDD S DDEQEKKPEAPKLSK	1144.61
12 O60841	EIF5B	214	S	GQKKNQKNKPGPNIE S GNE DDASF KIKTV A	1140.61
13 Q13435	SF3B2	435	S	KKKGFEE EHKDSDDD S DDEQEKKPEAPKLS	1061.95
14 Q96ST2	IWS1	248	S	QASDSENEELPKPRI S DSE SEDPPRHQASDS	983.28
15 Q4G0J3	LARP7	261	S	TSISKMKRSRPTSEG S DIESTEPQKQCSKKK	964.25
16 P52701	MSH6	227	S	TTYVTDKSEEDNEIE S EEEVQPKTQGSRRSS	943.95
17 Q9UBB9	TFIP11	98	S	AGLKKGA AAEAELED S DDEEKPVQDDFPKD	865.29
18 Q5C9Z4	NOM1	320	S	GKRVRAFAEDEEKSEN S EDGDITDKSLCGSG	814.29
19 Q9UQ35	SRRM2	1925	S	RSRSRASPVSRRRSR S RTPPVTRRRSRSRTP	806.29
20 P42166	TMPO	66	S	PPLPAGTNSKGPPDF S DEEREPTPVLGSGA	786.95
21 Q9H0D6	XRN2	499	S	GSPSP LGGIKRKAED S DSEPEPEDNVRLWEA	786.63
22 O95218	ZRANB2	188	S	EDEDDADLSKYNLDA S EEEDSNKKSNRRSR	781.63
23 Q13428	TCOF1	277	S	RAKKPEEESES SEEG S ESEE EAPAGTRSQVK	766.96
24 P05455	SSB	366	S	GKGKVQFQGKTKFAS S DDEHDEHDENGATGP	747.30
25 P48634	PRRC2A	350	S	YTEKLKF S DEEDGRDS S DEEGAEGHRDSQSAS	727.63
26 Q5C9Z4	NOM1	317	S	KRRGKRVRAFAEDEEKSEN S SEDGDITDKSLC	707.96
27 Q13247	SRSF6	303	S	DIKS KRSRSRSQSRNS S PLPVPPSKARSVSPP	688.51
28 Q9H0G5	NSRP1	248	S	ILQTDV KVEENPDADS S DFDAKSSADDEIEET	688.30
29 Q9HCN4	GPN1	338	S	ILTRGT LDEEDEAADS S TDDIDHRVTEESHE	684.34
30 O43719	HTATSF1	600	S	ENVLDKELEENDSEN S E FEDDGSEKVLDEEG	648.97
31 O43719	HTATSF1	624	S	KVLDEEGSEREF D ED S DEKE EEE DTYEKVFD	629.30
32 O43719	HTATSF1	597	S	ELHENVLDKELEENDSEN S E FEDDGSEKVL	625.58
33 P35659	DEK	32	S	PASEKEPEMPGPREE S EEE EE DE DEEEEEE	609.64
34 Q5VT52	RPRD2	374	S	PEPVTDNRDV EDMEL S D V EDDGSKII IVEDRK	609.27
35 P25205	MCM3	672	S	VLEKEKKRKKRSEDE S ETEDEE EKSQEDQEQ	571.30
36 P49756	RBM25	703	S	KLPVD S VFNKF EDED S D DV PRKRKL VPLDY G	570.83
37 P51116	FXR2	453	S	TGGPAYGPSSDV STA S E TESEKREEPN RAGP	570.30
38 O60231	DHX16	103	S	RALLEKNRSYRLLED S E S ESSEETV RAGSSL	570.16
39 P52701	MSH6	261	S	KKRRVIS D SES D IGGS S D V E F K P D T KEEGSSD	550.71
40 P14625	HSP90B1	306	S	EPM EEE AAKEEKEE S D E AA V EEEEKKP	550.67
41 Q9H3N1	TMX1	247	S	LKKV EEE QE A DEED V S E EEAES K EGTNKDFP	550.64
42 Q8WVC0	LEO1	300	S	RMKRKN A IAS D SE A DS D T E V P K D NS G T M LF	550.28
43 O43719	HTATSF1	642	S	KEEE E DTYEKVFD D DE K E EE YADEKGLE	530.97
44 P51532	SMARCA4	695	S	PTLPVE KKK IP D PD D SD V SE V D A R H I E NA	512.30
45 Q13435	SF3B2	431	S	SAAPKKGF EE H K D S DD S D E QEKKPEA	511.75
46 O00505	KPNA3	60	S	LLKKRNVPQ EES LED S D V D A F K AQN V T LE A	511.31
47 P24534	EEF1B2	106	S	GATDSK DDDD IDLFG S D EE ESE A KRLREE	493.24
48 P52701	MSH6	274	S	GGSD V E F K P D T KEEG S D E ISS G VG D SE E SEG	491.92
49 Q8WVC0	LEO1	294	S	SEDEV L R M K R KN A IAS D SE A DS D T E V P K D NS	491.64
50 Q8WVC0	LEO1	296	S	DEVL R M K KN A IAS D SE A DS D T E V P K D NS G T	491.28
51 Q9H0G5	NSRP1	255	S	VEENPD A DS D FA K S S AD D IE E TR V N C R RE	472.21
52 O95218-2	ZRANB2	305	S	SRSSSSGDRKKR R TRS R S P E S Q V I G E N T K QP	471.98
53 Q6QNY0	BLOC1S3	65	S	RPTGLRVAGEAAET D S E PE E PE P TAAPRDL	471.69
54 Q99856	ARID3A	88	S	GHPASPGG SE D G PPG S EE D A R E G TPGSPG	471.42
55 O95671	ASMTL	239	S	VKHDSI P A D T F E D L S D V EG G SE T Q R D A G	462.51

56 P49756	RBM25	583	S	EAERRRQPQIKQEPE\$EEEEEEKQEKEEKRE	453.25
57 P55081	MFAP1	116	S	LARHRKIVEPEVVGE\$DSEVEGDAWRMERED	452.72
58 Q86UE4	MTDH	426	S	SLLKSQEPIPDDQKV\$DDDKEKGEGALPTGK	452.57
59 Q9NTJ3	SMC4	28	S	RREEGPPPPSPDGAS\$DAEPEPPSGRTESPA	452.47
60 Q9UEG4	ZNF629	28	S	QGPEQSPNDAHRAAES\$ENEESPRQESSGEE	452.31
61 P67809	YBX1	313	S	QDGKETKAADPPAENS\$APEAEQGGAE*****	432.83
62 Q8NHM5	KDM2B	975	S	RTENSLANENQQPIK\$EPESEGEEPKRPPGI	432.79
63 Q96EV2	RBM33	41	S	AERSWRRAADEWD\$ELEDDLGEDLLSGK	432.65
64 Q9NTJ3	SMC4	27	S	RRREEGPPPPSPDGAS\$DAEPEPPSGRTESP	432.57
65 P13861	PRKAR2A	78	S	PPEPGPDRVADAKGD\$ESEEEDEDLEVVPVPSR	432.14
66 P35269	GTF2F1	307	S	PQQEEGPKGVDQS\$SEESEEEKPPEEDKE	431.73
67 P52701	MSH6	252	S	GSRRSSRQIKKRRVI\$DSES DIGGSDVEFKP	413.31
68 P52701	MSH6	256	S	SSRQIKKRRVISDSES\$DIGGSDVEFKPDTKE	413.14
69 P55209	NAP1L1	10	S	*****MADIDNKEQS\$ELDQDLDDVEEVEEE	412.98
70 Q9BW71	HIRIP3	223	S	AKKVEGNKGTKSLKE\$EQESEEEILAQKKEQ	412.87
71 O43719	HTATSF1	616	S	Efeddgsekvldeeg\$erefdeddsdekeeee	393.82
72 O60293	ZFC3H1	28	S	GLSPKEEGELEDGEIS\$DDDNNNSQIRSRSSSS	393.74
73 Q6P6C2	ALKBH5	64	S	EPYPVSGAKRKYQED\$DPERSDYEEQQQLQKE	393.62
74 Q8IWX8	CHERP	817	S	SKSYSPGRRRRSRSRSPTPPSSAGLGSNSAP	393.58
75 Q8N1G4	LRRC47	520	S	EMKKYTLENKEEGSL\$DTEADAVSGQLPDPT	393.49
76 Q9H0G5	NSRP1	254	S	KVEENPDADSDFDAKS\$ADDEIEETRVNCRR	393.31
77 Q7KZ85	SUPT6H	75	S	DDDDDEGEDEGE\$SGDSEDDVGHKRKRT	393.28
78 P55081	MFAP1	118	S	RHRKIVEPEVVGES\$EVEGDAWRMEREDSS	373.81
79 O43719	HTATSF1	714	S	DDDSNEKLFDEEEDS\$EKLFD\$DERGT\$LG	373.72
80 O76031	CLPX	617	S	YIRAPTKESSEEEYD\$GVEEEGWPRQADAAN	373.67
81 P05388	RPLP0	307	S	AAAPAKVEAKEESE\$EDMGFLFD*****	373.65
82 Q13428	TCOF1	87	S	QAKKTRVSDPIST\$E\$SEEEEEEAAETAKAT	373.58
83 Q13442	PDAP1	60	S	DGAAGDPKEKKSLDS\$DESEDEDDYQQKRK	373.47
84 Q14527	HLTF	397	S	SSRPKRRKTAQYIES\$DSEEIETSEL\$PQKM	373.39
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91 Q9UNS1	TIMELESS	1173	S	KEPLKAAPKKRQLLD\$DEEQEEDEGRNRAPE	353.98
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96 O60841	EIF5B	182	S	DEDNSKKIKERSRIN\$SGESGDESDEFQSR	353.74
97 P13861	PRKAR2A	80	S	E\$PGPDRVADAKGD\$ESEEEDEDLEV\$PVPSRFN	353.69
98 Q7KZ85	SUPT6H	73	S	INDDDEGEDEGE\$SGD\$EDDVGHK\$KRT	353.57
99 Q8TF01	PNISR	290	S	EGGDGPRLPQR\$KFD\$DEEEEDTENVAASS	353.48
100 Q9P2I0	CPSF2	419	S	EAAKKLEQSKEADID\$SDES D\$DIEIDQPSA	351.21
101 Q9UHB7	AFF4	222	S	HQRSKSPRD\$P\$SRVPFSSQHQSTQS	347.61
102 Q9UJV9	DDX41	23	S	RKRARTDEV\$AGGS\$RSEAEDDED\$DYV\$PY	335.60
103 Q9UQ35	SRRM2	497	S	GR\$RSRS PATAKRGRSRSRTPTKRGHSRSRSPQ	334.93
104 Q9Y2W1	THRAP3	379	S	QTNTDKEKIKE\$GFS\$DTGLGDGKMKSDSFA	334.84
105 O14654	IRS4	775	S	KAPDTNKEDDSKD\$ND\$ESDYMFM\$APGAGAIP	334.78
106 O43719	HTATSF1	676	S	KKAEEGDADEKL\$EE\$DDKEDEDADGKEVED	334.67
107 O60524	NEMF	747	S	QSGRDELNEELIQEE\$SEDEGEYE\$EV\$RKDQD	334.51
108 Q8WXI9	GATAD2B	129	S	PERGR\$LPSPDIIVL\$DNEASSPRSSSRMEE	334.47
109 Q9BWU0	SLC4A1AP	312	S	KQQQILLEKKMLGED\$DEEEEMDTSERKINA	334.38
110 Q92733	PRCC	267	S	AALQVT\$KQITQEE\$DDEEVAPENFFSLPEK	334.37
111 Q9UQ35	SRRM2	1729	S	PETRSRTPPR\$HRRSP\$VSSPEPAEKSRSR\$R	334.36
112 O60841	EIF5B	186	S	SKKIKERSRIN\$SGE\$GDESDEFQSRKGQK	334.32

113 Q8NHM5	KDM2B	979	S	SLANENQQPIKSEPE\$EGEEPKRPPGICERP	334.28
114 P46821	MAP1B	1016	S	EEDMDEAIEKGEAEQ\$EEEADEEDKAEDARE	334.15
115 Q13428	TCOF1	769	S	VKAEKQEDSESSEEE\$DSEEAAASPAQVKTS	314.93
116 Q5T8D3	ACBD5	196	S	TPNAKTVNGKAESS\$GAESEEEEAQEEVKG	314.88
117 Q8NAV1	PRPF38A	193	S	EPRVSALEEDMDDVES\$EEEEEEDEKLERVP	314.82
118 Q9Y2W2	WBP11	353	S	AGQEIPEEGREVEEF\$EDDDDEDSDDSEAEK	314.73
119 Q9P2I0	CPSF2	423	S	KLEQSKEADIDSS\$DES DIEEDIDQPSAHKT	314.67
120 Q13308	PTK7	1064	S	KDRPSFSEIASALGD\$TVDSKP*****	314.65
121 Q14527	HLTF	400	S	PKRRKTAQYIESS\$SEEIETSELPOQKMK	314.63
122 P21127	CDK11B	47	S	IKRLKNSSDRDRSKRD\$LEEGERDHRMEITI	314.50
123 P22059	OSBP	351	S	VGSGKDQCCSGKGDM\$DEDDENEFFDAPEII	311.40
124 P34741	SDC2	115	S	QTKSPEETDKEKVHLS\$D SERKMDPAEEDTNV	298.34
125 Q14684	RRP1B	452	S	KALKARVAEPGAEAT\$STGEESGEHPPAVP	295.25
126 Q9H307	PNN	347	S	EEAGEEEEKEIAIVHS\$DAEKEQEEEEQKQEM	295.15
127 Q9NWV8	BABAM1	49	S	AEDRAVGAQASVGSR\$E\$EGE\$EAASADDGSLN	294.99
128 Q9UQ35	SRRM2	1541	S	KTVARTPLGQRSRSG\$SQELDVKPSASPQER	294.84
129 Q9P2I0	CPSF2	420	S	AAKKLEQSKEADIDSS\$DES DIEEDIDQPSAH	294.73
130 P35269	GTF2F1	308	S	QQEEGPKGVD EQSDS\$EESEEEKPPEEDKEE	294.35
131 Q8IY81	FTSJ3	599	S	SPLYQDEAPKGTEASS\$GTEATGLEGEEKDG	287.34
132 O95218-2	ZRANB2	307	S	SSSSGDRKKRTRRS\$PESQVIGENTKQP**	284.50
133 Q14527	HLTF	398	S	SRPKRRKTAQYIESS\$DSEEIETSELPOQKMK	275.72
134 O60231	DHX16	106	S	LEKNRSYRLLEDSEES\$SEETVS RAGSSLQKK	275.68
135 Q13308	PTK7	1055	S	MQRCWALSPKDRPSF\$EIASALGD\$TVDSKP	275.53
136 O15541	RNF113A	84	S	KTRD\$GKQKAAYGDL\$SEEEENEPESLGVV	275.47
137 P18858	LIG1	66	S	SPVKRPGRKAARVLG\$E\$GGEE\$EADELPAKGQ	275.32
138 P51532	SMARCA4	1627	S	GRRRPSRGSR A\$RAKPVV\$DDDSEEEQEDRSGS	275.29
139 P55081	MFAP1	132	S	DSEVEGDAWRMEREDS\$SEEEEEEIDDEEIER	275.24
140 Q13427	PPIG	257	S	KKEKKRKKS\$KKS\$AS\$SESEAENLEAQ\$PQ\$STV	268.34
141 Q99613	EIF3C	11	S	*****MSRFFTTGSD\$ESESS\$LSGEELVT\$K	267.30
142 Q9UQ35	SRRM2	625	S	ARRGRSRSRTPARR\$RTRSPVRRSRSRSP	264.83
143 Q9Y2W2	WBP11	361	S	GREVEEF\$E\$DDDE\$DD\$SEAEKQSQKQHKE	264.37
144 P55081	MFAP1	133	S	SEVEGDAWRMEREDS\$SEEEEEEIDDEEIER	264.31
145 P49736	MCM2	139	S	AGRGLGRMRGRLLYD\$DEEDEER PARKRRQV	261.90
146 Q13895	BYSL	98	S	RERTTRLGPMPQDG\$D DEEDEEWPTLEKAAT	256.47
147 Q6KC79	NIPBL	1154	S	RSSGGGRYRNRS\$PSD\$DMEDYSPPP\$LS\$EV	255.79
148 Q8TEU7	RAPGEF6	230	S	VGDVDLTRLPEGPV\$D EDEEDEEIDRTDP	255.73
149 Q96S55	WRNIP1	92	S	SSALKQPATPTAAE\$SE\$EGE\$EE\$GGDDGETES	255.67
150 Q99613	EIF3C	9	S	*****MSRFFTTGSD\$ESESS\$LSGEELVT	255.65
151 Q9BW71	HIRIP3	370	S	LGSTS\$GEESDLEREV\$D SEAGGGPQGERKNR	255.61
152 Q9Y467	SALL2	802	S	GSESGGEKAISVRGD\$EEASGAEEVGTVAA	255.59
153 O14617	AP3D1	632	S	VPEGLDLD\$A WINEPL\$DSESED\$ERPRAVFHE	255.53
154 O15541	RNF113A	253	S	EGRYGVYEDENYEVGS\$DDEEIPFKFC\$ICRQS	255.40
155 O43823	AKAP8	323	S	FQLYEEPDTKLARV\$D EGDFSENDDAAGDFR	255.35
156 O60341	KDM1A	131	S	AKVEYREMDES LANL\$E DEYYSEERNAKAE	241.36
157 P35659	DEK	301	S	DSSTTKNQN\$SS\$K\$E\$ESEDSS\$DEPLIKKLKKPPT	236.27
158 P35659	DEK	306	S	KKNQN\$SS\$KE\$ESEDSS\$DEPLIKKLKKPPT	236.15
159 P52756	RBM5	624	S	ENPLKRLVAAYSGD\$D NEEELVERLESEE	235.99
160 Q13428	TCOF1	764	S	PTVTQVKAEKQED\$E\$EE\$DS\$EEAAASPA	235.92
161 Q13442	PDAP1	63	S	AGDPKKEKKSLD\$DE\$EDEDDYQQKRKGVE	235.84
162 Q14498	RBM39	129	S	LPHSIKLSRRRSRSK\$PFRKDK\$PVREPIDN	235.83
163 Q8IWX8	CHERP	813	S	SRSRSKSYSPGRRRS\$RSRSPTPPSSAGLGS	235.80
164 Q8N1G4	LRRC47	518	S	MAEMKKYTLLENKEEG\$LS DTEADAVSGQLPD	235.78
165 Q8NHQ9	DDX55	544	S	EKKKKMNEKRKREEG\$DIEDEDMEELLNDTR	235.76
166 Q8WVC0	LEO1	171	S	DEKIQNSDDEERAQGS\$DEDKLQNS\$DD\$DEKMQ	235.72
167 Q99613	EIF3C	178	S	PEQSADEDAEKN\$E\$GSS\$DEDEDEDGVSA	218.27
168 Q9H501	ESF1	663	S	KQKALAAEASEEELP\$DV DLNDPYFAEEVKQ	218.12
169 Q8NAV1	PRPF38A	194	S	PRVSALEEDMDDVES\$EEEEEEDEKLERVPS	217.62

170 Q5T8D3	ACBD5	200	S	KTVNGKAESSDSGAES EEEEAQEEVKGAEQS	217.58
171 O43524	FOXO3	280	S	GRAAKKAAALQTAPES ADDSPSQLSKWPGSP	217.53
172 O43823	AKAP8	328	S	EPDTKLARVDSEGDF SENDDAAGDFRSGDEE	217.32
173 O60841	EIF5B	190	S	KERSRINSSGESGDES DEFIQLQSRKGQKKNQK	217.19
174 Q00341	HDLBP	31	S	RSGLVPQQIKVATLN SEEESDPPTYKDAPP	216.93
175 Q12872	SFSWAP	283	S	LAENKSDEKKKSGVSS DNEDEDDDEEDGNYLH	216.78
176 Q5VZL5	ZMYM4	122	S	TDDSLEVERRVTQHES DNENEIQIQNKLKD	216.64
177 Q7Z417	NUFIP2	112	S	LNGNAGEREISLKNL SDEATNPISRVLNGN	216.62
178 Q7Z4V5	HDGFRP2	240	S	KKKAPSASDSDSKADS DGAKPEPVAMARSAS	216.59
179 Q86VM9	ZC3H18	83	S	DEEDRASEPKSQDQDS EVNELSRGPTSSPCE	216.49
180 Q8N7H5	PAF1	456	S	EDEARAARDKEEIFGSDADSEDDADSDDEDR	216.47
181 Q93009	USP7	18	S	HQQQQQQQQKAGEQQL SEPEDMEMEAGDTDDP	216.36
182 Q99543	DNAJC2	63	S	ASFQELEDKKELSEES SEDEELQLEEFPMPLKT	216.32
183 Q9H501	ESF1	657	S	KKRLKRKQKALAAEAS SEEELPSVDLNDPYF	199.25
184 Q9UKJ3	GPATCH8	890	S	DASSDQSCYSRQRSY SDDSYSYSDSRSRRHS	198.53
185 Q9UKY1	ZHX1	47	S	PPVLTVENTRAESIS SDEEVHESVSDDNQQ	198.34
186 Q9UPT8	ZC3H4	808	S	QDRENEEGDTGNWYSS DEDEGGSSVTSILKT	197.66
187 Q9UQ35	SRRM2	416	S	RDRSPPKSPEKLPQSS SSSESSPPSPQPTKVS	197.38
188 Q7Z4V5	HDGFRP2	370	S	ERADRGEAERGSGGS GDELREDDEPVKKRG	197.35
189 P51532	SMARCA4	1631	S	PSRGSRACKVVS DDDSSEEQEDRSGSGSEE	197.29
190 O60841	EIF5B	183	S	EDNSKKIKERSRINS SGESGDESDEFIQLSRK	197.15
191 Q99613	EIF3C	182	S	ADEDAEKNEEDSEGS SDEDEDEDGVSAATFL	196.83
192 O00267	SUPT5H	19	S	SEDSNFSEEDSERSS SDGEEAEVDEERRSAA	196.81
193 O75475	PSIP1	106	S	KFSSQQAATKQSNASS SDVEEVEEKETSVSKED	196.76
194 O95831	AIFM1	118	S	GLTPEQKQKKAALSA SEGEEVPQDKAPSHVP	196.72
195 P52701	MSH6	275	S	GSDVEFKPDTKEEGS SDEISSVGVDSESEGL	196.67
196 Q13428	TCOF1	272	S	LPPAKRAKKPEEESES SSEEGSESEEEEAPAGT	196.66
197 Q4G0J3	LARP7	258	S	TSNTSISKMKRSPRT SEGSDIESSTEPQKQCS	196.59
198 Q5C9Z4	NOM1	280	S	GDVEKEKKAQEAEAQ SSEDDDEDTEEEQGEEK	196.53
199 Q5VTR2	RNF20	136	S	LTERKALVVPEPEPD SDSNQERKDDRERGE	196.48
200 Q8N5F7	NKAP	157	S	APEVWGLSPKNPEPD SDEHTPVEDEEPKKST	196.38
201 Q96B36	AKT1S1	202	S	SVPVWGFKEKRTEARS SDEENGPPSSPDLLR	196.27
202 Q96DR7	ARHGEF26	392	S	AVLYQNYKEKALDIDS SDEESEPKEQKSDEKI	195.75
203 Q9BW71	HIRIP3	196	S	APGKASVSRKQAREE SSEESEAEPVQRTAKV	194.25
204 Q9NYF8	BCLAF1	385	S	RAEGEWEDQEALDYF SSDKESGKQKFNDSEGD	187.34
205 Q9P1Y6	PHRF1	915	S	SAMSKLRGAVAAEGA SSDTEREEPTESQGLAA	186.24
206 Q9ULX3	NOB1	201	S	EEEEEEENG FEDRKDDSSDDGGGWITPSNIKQ	179.34
207 Q9UQ35	SRRM2	1102	S	QSKSQTSPKGGRSRSS SSPVTELASRSPIRQD	177.36
208 Q9BQG0	MYBBP1A	738	S	SEEGEDNRSS SEEESEGESEEEERDGDVD	177.26
209 Q7Z4V5	HDGFRP2	396	S	VKKRGRKGRGRGPP SSDSEPEAELEREAKK	177.12
210 Q7Z4V5	HDGFRP2	236	S	GGRK KKKAPSASDSDSKADSDGAKPEPVAMA	176.99
211 Q9UQ35	SRRM2	1923	S	RRRSRSRASPVSRRRS SRSRTPPVTRRSRSR	176.97
212 Q8WVC0	LEO1	279	S	EEEQDHKSESARG SSEDEVLRMKRNIAAS	176.96
213 Q9UKY1	ZHX1	48	S	PVLTPVENTRAESIS SSDEEVHESVSDDNQN	176.93
214 Q92733	PRCC	159	S	PVKIAAPELHKGD SSEEDEPTEKKTILQGS	176.86
215 Q9UQ35	SRRM2	1382	S	PSLDMKEQSTRSSGH SSSSELSPDAVEKAGMS	176.81
216 Q9UPT8	ZC3H4	807	S	KQDRENEEGDTGNWY SSSDEDEGGSSVTSILK	176.79
217 Q8N7H5	PAF1	460	S	RAARDKEEIFGSDAD SSEDDADSDDEDRGQAQ	176.73
218 Q9BQG0	MYBBP1A	734	S	AEDKSEE GEDNRSSSEEESEGESEEEERD	176.56
219 Q7Z4V5	HDGFRP2	395	S	PVKK RGRKGRGRGPPSSDSEPEAELEREAK	176.55
220 Q8WVC0	LEO1	220	S	SEE EKANSSDDERPVASSDNDEKQNSDDEQP	176.24
221 Q13428	TCOF1	762	S	TGPTVTQVKAEKQED SSESSSEEESDSEEEAAS	176.17
222 Q99613	EIF3C	181	S	SADEDAEKNEEDSEGS SSEDEDEDEDGVSAATF	175.68
223 Q13428	TCOF1	765	S	TVTQVKAEKQED SSEEESDSEEEAASPAQ	175.49
224 Q9H501	ESF1	313	S	SGPDLARGKGNIET SSEDEDDTADLFPEESG	174.91
225 Q8WVC0	LEO1	179	S	DEERAQGSDEDKLQ NSDDEKMQNTDDEERP	174.85
226 Q7Z4V5	HDGFRP2	399	S	RGRKGRGRGPPSS SSDSEPEAELEREAKKSAK	172.54

227 Q8N7H5	PAF1	466	S	EEIFGSDADSEDDADS DDEDRGQAQGGSDND	171.30
228 O15541	RNF113A	85	S	TRDSGKQKAAYGDL S EEEEENEPESLGVVY	168.63
229 O60841	EIF5B	137	S	KSKKTAKPKVEMYSG S DDDDFNKL P KKAKG	168.34
230 Q13610	PWP1	50	S	LIAEAKELQEEGGGS S EEETGSP S EDGMQS	167.39
231 Q7Z4V5	HDGFRP2	232	S	RGPLGGRKKKKAPSAS D SDSKADSDGAKPEP	167.35
232 Q9UQ35	SRRM2	1499	S	PEPKALPQT P RPRSRS P SSPELNNKCLTPQR	166.53
233 Q8WVC0	LEO1	277	S	DDEEEQDHKS E SARG S DSEDEVLRMKRKN A I	165.39
234 Q9BQG0	MYBBP1A	732	S	KGAEDKSEE G DNRS S EESEE G EESE EEE	165.37
235 Q9BVG4	PBDC1	197	S	GEEENTKNGGEKGAD S GEEKEGINREDKTD	165.34
236 Q9NYF8	BCLAF1	397	S	DYFS D KESGKQKFND S EGDDTEETEDYRQFR	164.69
237 Q9UKV3	ACIN1	838	S	KISVVSATKGPAG N S T EGGQ P GRKRRWGA	164.54
238 Q6P6C2	ALKBH5	69	S	SGAKRKYQEDSDPERS D YEEQQLQKEEEARK	164.35
239 O14617	AP3D1	634	S	EGLLDL A WINEPL S SE E DERPRAVFHEEE	164.30
240 Q7Z4V5	HDGFRP2	234	S	PLGGRKKKKAPSAS D SKADSDGAKPEPVA	164.20
241 Q7Z4V5	HDGFRP2	397	S	KKRGRKGRGRGP P SS S DSEPEAELEREAKKS	163.42
242 P35659	DEK	307	S	KNQN S KK E SE S ED S SD E PLIKLK K PPTD	162.48
243 Q13428	TCOF1	279	S	KKP E ESES S EE G SE S EE E APAGTRSQVKAS	159.42
244 Q7Z4V5	HDGFRP2	366	S	ERRRERADRGEAERG S GGSSGDELREDDEPV	159.37
245 Q8IWA0	WDR75	782	S	IPEDVD M EEKE S ED S DEENDFTEVQDTSN	158.97
246 Q99613	EIF3C	166	S	DFESHITSYKQNPEQ S ADEDAE K NEEDSEGS	158.75
247 Q9Y5B6	PAXBP1	154	S	YKEDLEKSKIKTELNS S AESEQPLDKTGHVK	158.64
248 Q8WVC0	LEO1	212	S	SDDERQQL S EE E KAN S DDERPVASDND D E K Q	157.88
249 Q96B36	AKT1S1	203	S	VPVWGFKEKRTEARS S DEENGPPSP D L D R I	157.82
250 Q9BTC0	DIDO1	809	S	PRQEAI P LED S PPV S D S EEQQESARAVPEK	157.73
251 Q9H307	PNN	417	S	DQEV M ETNRVESVEP S NEASKELEPEMEFE	157.69
252 Q9Y2U8	LEMD3	259	S	VNGSRLVPYS C RENY S D S EE E DDD V ASSRQ	157.66
253 Q9Y6E2	BZW2	414	S	KKF V EWLQNAEE E SE G E N *****	157.52
254 O60524	NEMF	748	S	SGRDELNEELIQEES S DEGEYE E VRKDQDS	157.47
255 Q7Z4V5	HDGFRP2	369	S	RERADRGEAERG S GGG S GD E L R DEPVKKR	157.34
256 Q9BW71	HIRIP3	199	S	KASVSRKQAREE S EE A EPVQRTAKKVEGN	157.33
257 O14617	AP3D1	636	S	LDLD A WINEPL S DE E DERPRAVFHEEEQR	156.47
258 Q9UQ35	SRRM2	2702	S	DSRSLSYSPVERRRP S PQPSPRDQQ S SS S ER	154.31
259 O15357	INPPL1	132	S	VEGEREPDP P DDR D AS D GEDEK P PL P RS G S	148.32
260 Q13428	TCOF1	270	S	GALPPAKR A K P EEE S ES S EE E APA	147.65
261 O75683	SURF6	206	S	E P REP P GLIFNKVEV S DE E PASKAQRRKEKR	145.13
262 P52701	MSH6	279	S	EFKPDTKEEGSS E IS S VGVD S E S GLNSPV	139.79
263 Q12872	SFSWAP	604	S	DLLPLEKRN V KL D DD S DD D EESKE G QESSSS	139.66
264 O95400	CD2BP2	194	S	GARGGGKGRKGPGQ P SPQR L DR L SG I ADQM	138.89
265 O95232	LUC7L3	425	S	SDTKNEVNGT S EDI K SEGDTQSN*****	137.91
266 P45973	CBX5	13	S	***MGKKTKRTAD S SS E EEYYV V EVVLDR	137.83
267 P78316	NOP14	146	S	GQSLADIEKHNDIV D SD S DAEDRG T L S AELT	137.77
268 Q03468	ERCC6	1142	S	GECSNSSGT K T S MP S G D E S IDEKL G LSYKR	137.68
269 Q5VTL8	PRPF38B	529	S	DSKDQSDKHD R RR S Q S IEQESQE K H K N K DE	137.63
270 Q8IYB3	SRRM1	551	S	RQKETSPRGRR R SP S PPP T RR R SP S PAPP	137.55
271 Q8NC51	SERBP1	25	S	GCVVTNRFDQLFD D SD P F V L K AENKK E	137.48
272 Q8NFC6	BOD1L1	2986	S	QKSVSD V DE V DK K QE S DEEEEEEEDEPSGA	134.20
273 P35269	GTF2F1	311	S	E G PKG V DEQ S DS S SEE S EE E KP E EDK E EE	133.85
274 Q6PD62	CTR9	970	S	RKKKKR R R R HPKG E EGGS S DDDETENG P PK K RR	132.61
275 Q8WVC0	LEO1	630	S	EEDKAQ R L K AK K L T S D EE E GE P SG K R K A ED	131.25
276 Q99459	CDC5L	303	S	FTKKRSK L VLP A P Q I S D AE L QEV V KVGQASE	129.34
277 Q9UQ35	SRRM2	1501	S	PKALPQT P R R RS R SP S SP S PELNNKCLTPQR R	128.34
278 Q9Y467	SALL2	806	S	GGEKAISVRGD S EE A GA E EV G TVAAAATA	128.26
279 P11717	IGF2R	2409	S	KSVKALSS L HGDDQ D SE D EV L T I PEVKVHSG	127.99
280 P21127	CDK11B	277	S	PAQLKEEK M ERD L LS D Q D IS D S E R K T S A	127.96
281 P35611	ADD1	358	S	LLNPEKYKAKSRSP G SP V GE G T G S P PK W Q I G	126.93
282 P51532	SMARCA4	1570	S	SVFTSVRQKIEKED D SE G EE S EE E EE G EE E GG	126.83
283 Q9BW71	HIRIP3	291	S	CKQKSQAK R L G D S DEEEQ K EAASSGDDSG	126.14

284 Q9BXJ9	NAA15	855	S	PPGYEEDMKITVNGD\$SAEAEELANEI*****	119.67
285 Q9H6S0	YTHDC2	1090	S	QEPSSFRVDGIPNDS\$DSEMEDKTTANLAAL	119.47
286 Q9H1E3	NUCKS1	132	S	QEEEDEA\$PFQEKGDS\$DED\$FLMEDDDSDY\$G	118.59
287 Q9H501	ESF1	312	S	D\$GPDLARGKGNIET\$SEDED\$TADLFPEES	118.51
288 Q9NXG2	THUMPD1	88	S	YGPEKFTDKDQQPSGS\$EGEDDDAEALKKEV	118.41
289 Q9UQE7	SMC3	1067	S	KATLVMKKGDVEGSQS\$QDEGE\$GESERGSG	118.27
290 P78316	NOP14	148	S	SLADIEKHNDIVDSD\$DAEDRGTLSAELTAA	118.15
291 P17812	CTPS1	574	S	CRLSPRDTYSDRSGSS\$SPDSEITELKFPSIN	117.98
292 P35251	RFC1	69	S	FKQKQPSKKR\$IIYD\$DSESEETLQVKNAKK	117.95
293 P51116	FXR2	410	S	PGSGRGGGSDKAGYS\$TDESSSS\$HATRTY	117.88
294 Q03701	CEBPZ	629	S	KAKPGLRSQLDDHPE\$DDEENFIANDDEDM	117.77
295 Q08AD1	CAMSAP2	1148	S	PPEKADVPVEKYDGE\$DKEQFDDDQKVCCGF	116.92
296 Q13428	TCOF1	273	S	PPAKRAKKP\$EESES\$EEGSESEEAPAGR	116.87
297 Q13428	TCOF1	347	S	AGKPEED\$ESSSEES\$DSEEETPAAKALLQA	116.63
298 Q3YEC7	RABL6	471	S	PLPAGPVPSQDITLS\$SEEEAEVAAPTKGPAP	116.59
299 Q4G0J3	LARP7	337	S	KEASEASKENRDIEIS\$TEEEKDGTGDLKDSSL	116.38
300 Q8N1G2	CMTR1	55	S	SHGAKASTTSLSGSD\$ETEGKQHSSDSFDDA	116.17
301 Q96G74	OTUD5	177	S	GGGS\$PEREEVGAGYN\$EDEYEAAAARIEAMD	115.99
302 Q96ST2	IWS1	157	S	HASDSENEDEVGKHPAS\$DSEIEELQKSPASDS	115.93
303 Q9BW71	HIRIP3	330	S	RKSEDRTQLKGGKRLSGS\$SEDEEDSGKGEPT	115.88
304 Q9NW75	GPATCH2	115	S	SKDYRENHNNNKKDH\$SD\$DDQMLVAKRRPSS	115.83
305 Q9NZ63	C9orf78	15	S	*MPVVRKIFRRRRGD\$ESEED\$E\$QDSEEVRLK	115.79
306 Q9UQ35	SRRM2	510	S	SRSRPTKRGHSRSRS\$PQWRRSRSAQRWGRS	115.71
307 Q9Y2U8	LEMD3	140	S	ASAAPAAGSKVLLGFS\$DES\$DVEASPRDQAG	115.62
308 P51532	SMARCA4	1586	S	E\$GEESEEEEEGEEEG\$ESESRSVVKV\$IKLGR	109.22
309 Q13442	PDAP1	57	S	EGGDGAAGDPKKEK\$LD\$DESEDEEDDYQQ	106.27
310 P51532	SMARCA4	1575	S	VRQKIEKEDD\$E\$EGEE\$EEEEEGEE\$ESES	106.14
311 O75475	PSIP1	273	S	ESKRKNLAKTGVTS\$DSEE\$GDDQEGEKKR	105.88
312 Q9BW71	HIRIP3	332	S	SEDRTQLKGGKRLSGS\$SEDEEDSGKGEPTAK	105.73
313 Q13427	PIPIG	744	S	ENDHVHEKNKKFDHE\$SPGT\$DEDKSG*****	105.32
314 O43847	NRD1	86	S	DLGENSRVARLGADE\$EE\$EGRG\$LSNAGDP	104.58
315 P18858	LIG1	201	S	KPLKTSKAETPTESV\$EPEVATKQELQEEE	104.54
316 Q03468	ERCC6	429	S	QKKVPQEI\$DDFFPS\$GEEAEAASVGEGGG	104.34
317 P35269	GTF2F1	305	S	KAPQQEEGPKGVD\$EQ\$SD\$SSEESEEKPPEED	103.38
318 P51116	FXR2	533	S	PYSL\$LTSEPEPPVDS\$EPGE\$PP\$PASARRRS	103.36
319 Q8IY81	FTSJ3	458	S	ADTFLSDL\$PRDDIYV\$DVEDDGDDTSLSDL	102.85
320 Q8WVC0	LEO1	608	S	RYKGGIREERARIYSS\$D\$DEGSEEDKAQRL\$	101.68
321 P35659	DEK	303	S	STTKKNQN\$SSKKE\$E\$ED\$SD\$DEPLIKLKK	101.37
322 Q6P158	DHX57	127	S	KALLRDLQEQDADAG\$ERGLSG\$EE\$DDEPDC	101.19
323 Q03468	ERCC6	430	S	KKVPQEI\$DDFFPS\$GEEAEAASVGEGGG	99.18
324 P52701	MSH6	285	S	KEEGSSDEI\$SSGVGD\$E\$EGLNSPVKVAR\$R	98.95
325 Q9Y2U8	LEMD3	261	S	GSRLVPYSCRENYSD\$EE\$DDVVASSRQVL	98.83
326 P16383	GCFC2	19	S	RPKRTFQR\$AA\$DSS\$DGAEE\$PAE\$PGAPRE	98.63
327 P05388	RPLP0	304	S	PA\$AAAAPAKVEAKEE\$EES\$EDDMG\$FGLFD**	98.56
328 P27824	CANX	564	S	EEKQKSDAEE\$GGTV\$QEE\$EDRKPKAEEDEI	98.39
329 P55884	EIF3B	125	S	EQARDERS\$DRAQAV\$EDAGGNEGRAAE\$P	98.35
330 Q08945	SSRP1	444	S	GLKEGMNP\$SYDEYAD\$D\$E\$QHDAYLERMKEE	98.33
331 Q13435	SF3B2	309	S	EEMETDARSS\$LGQSA\$E\$EE\$DTVS\$VSKKE\$N	98.31
332 Q14839	CHD4	103	S	EGPEFVEEEEVALRS\$DSEGSDYTPG\$KKKK	98.25
333 Q7Z4V5	HDGFRP2	664	S	RPGSDRQERERARGD\$EALDEES*****	98.14
334 Q86VM9	ZC3H18	34	S	QPQGLS\$DDILRD\$G\$DQ\$LDGAG\$VRA\$DLE	97.57
335 Q86X53	ERICH1	254	S	EEDL\$TRARQEGADA\$E\$ED\$PTPAGE\$DV\$KDA	94.60
336 Q8NAF0	ZNF579	194	S	PST\$LAAPTSAAE\$PRE\$E\$EE\$AAEAGAAELRAE	94.35
337 Q8NE71	ABCF1	105	S	DDGEEKELMERLKKL\$SVPT\$DEE\$DEVPA\$K\$P	94.34
338 Q8WVC0	LEO1	229	S	DERPVASDND\$DEKQNS\$D\$DEEQPQLS\$DEEKMQ	93.54
339 Q92538	GBF1	1298	S	LQATARADAPDAGAQ\$SD\$SELPSYHQNDVSLD	86.34
340 Q92733	PRCC	157	S	KEPVKIAAPELHKGD\$D\$SEED\$EPTKKTILQ	84.39

341 Q92917	GPKOW	35	S	SFGFTRTSARRRLADS GDGAGPSPEEKDFLK	83.54
342 Q99613	EIF3C	18	S	RFFTTGSDSESESSL SGEELVTKPVGGNYGK	82.39
343 Q9UQ35	SRRM2	1652	S	RSGSSSKGRGSPSPEG SSSTESSPEHPPKSRT	81.77
344 Q96ST2	IWS1	159	S	SDSENEVGKHPASD SEIEELQKSPASDSET	81.74
345 Q9BW71	HIRIP3	333	S	EDRTQLKGKRLSGS SEDEEDSGKGEPTAKG	81.71
346 Q9UQ35	SRRM2	1653	S	SGSSSKGRGSPSPEG SSSTESSPEHPPKSRTA	81.68
347 P35251	RFC1	71	S	QKQPSKKR IIYDSDS ESEETLQVKNAKKPP	80.66
348 Q9NW75	GPATCH2	117	S	DYRENHN NNKKDHSDS D DQMLVAKRRPSSNL	80.64
349 Q9UQ35	SRRM2	418	S	RSPPKSP EKLPOQSSS ESSPPSPQPTKVS RH	80.62
350 Q08945	SSRP1	667	S	SSRQLSESFKS KEFV S DESSSGENKSKKR	80.59
351 Q9UQ35	SRRM2	1762	S	SASSPRTK TTSRRGRS PSPKPRGLQR SRSRS	80.57
352 Q9UQ35	SRRM2	1542	S	TVARTPLGQRSRSG SQELDVKPSASPQRS	80.49
353 Q9BW71	HIRIP3	555	S	PDWSHMRGI ISSDGE N*****	79.42
354 Q9GZR7	DDX24	82	S	SLFSKEAPKRKAQAV S EEEEEEGKSSSPKK	79.37
355 O94979	SEC31A	527	S	DGANVALKDSDQVAQS D GEESPAEEQLLGE	79.37
356 Q9H0D6	XRN2	501	S	PSPLGGIKRKAED DS EPEPEDNVRLWEAGW	79.29
357 P19338	NCL	34	S	PPPKEVEEDSEDEEM S EDEEDDSSGEEVVIP	79.29
358 Q9HAW4	CLSPN	65	S	IFVSKKLKNRKVLQ D SDSETEDTNASPEKTT	79.25
359 Q9NQ55	PPAN	359	S	RKKSLEGMKKARV GG S DEEASGIPSRTASLE	79.15
360 Q9P289	MST4	304	S	DRFKRWKAEGHS D DESGDSESTSRENN	78.93
361 Q9UQ35	SRRM2	415	S	TRDRSPPKSPEKLPQ S SSSESSPPSPQPTKV	78.91
362 Q9Y5T5	USP16	415	S	NDKNLKKT V EDQDSEEKDNDSYIKERSD	78.90
363 Q9Y6R4	MAP3K4	66	S	RQEGLGDSACKSP E SDLED F DETNTENLY	78.89
364 Q9UQ35	SRRM2	1764	S	SSPRTK TTSRRGRS SPKPRGLQR SRSRR	78.85
365 Q9Y2U8	LEMD3	144	S	PAAGSKVLLGF S DES D VEASPRDQAGGGGR	78.84
366 Q9UQ35	SRRM2	1379	S	ETDPSLDMKEQSTRS S GHSSSELSPDAVEKA	78.82
367 Q8WVC0	LEO1	238	S	DDEKQNSD EEQPQL S D EEMQNSDDERPQA	78.73
368 Q9UQ35	SRRM2	1657	S	SKGRGPSP E GSSSTE S PEHPPKSRTARRGS	78.69
369 Q8NE71	ABCF1	109	S	EKELMERLKKL S V P TS S DEEDEV P A P KPRGGK	78.67
370 Q9HAW4	CLSPN	67	S	VSKKLKNRKVLQ D SD E TEDTNASPEKTYD	78.65
371 Q9UQ35	SRRM2	1378	S	LETDP S LDMKEQSTRS S GHSSSELSPDAVEK	78.63
372 Q9Y2U8	LEMD3	141	S	SAAPAAGSKVLLGF S DES D VEASPRDQAGG	78.61
373 O60841	EIF5B	595	S	DKKPSKEMSSDSEY D DDDRTKERAYDKAK	78.60
374 Q3YEC7	RABL6	470	S	PPLPAGPVPSQDITL S SEEAEVAAP T KGPA	78.58
375 Q8WVC0	LEO1	273	S	HRHSDEEEQDH K SE S ARGSDSEDEVLRMKR	78.53
376 Q1KMD3	HNRPUL2	161	S	EQGLGKREEDEPEERS G DET P GSEVPGDKAA	78.44
377 O75400	PRPF40A	938	S	TGKDSGNWD T SG S EL S E G ELEKRRRTL LE QL	78.38
378 P16383	GCFC2	17	S	AHRPKRTFRQRAAD S SD D GAEE S PAEP G AP	78.35
379 P17812	CTPS1	575	S	RLSPRD T YSDRG S SS P D S E I TELKFPSINH	78.33
380 P25205	MCM3	668	S	YFKKVLEKEKKRKKR S DE E SETEDEE E KSQE	78.25
381 P52701	MSH6	254	S	RRSSRQIKKRRVISD S ES D IGGS D V E FKPDT	78.18
382 Q07666	KHDRBS1	20	S	DDPAARMSRSSGRSG S MDPSGA H PSVRQTPS	78.14
383 Q5C9Z4	NOM1	321	S	KRVRFAEDEEKSEN S EDG D ITDKSLCGS G E	77.72
384 Q6WKZ4	RAB11FIP1	202	S	DTASAI I IPSTTPSVD S D D ESVVKDKKK SKI	77.66
385 Q86VM9	ZC3H18	46	S	DSGSDQ L DGAGV R AS D LEDEES A ARGPSQE	77.65
386 Q8N1G2	CMTR1	53	S	SVSHGAKASTT S LSG S D S E T EGKQHSSDSFD	77.50
387 Q8NAF0	ZNF579	196	S	TLAAP T SAAE P RE S SEEAEAGAAELRAELA	76.59
388 Q8NFC6	BOD1L1	266	S	RTSED M AD K STAD S GGEGLE T APK S EEFS	76.35
389 Q8WVC0	LEO1	607	S	NRYKGGIREERARIY S SDS D E G SEEDKAQRL	76.34
390 Q96E09	FAM122A	37	S	EGGGSGGGGLRR S NS A PLI H GLSD T SPVFQ	74.36
391 Q96NB3	ZNF830	351	S	LTI K ELQK EE ENAD S D D E G ELQ D LLSQDWR	73.59
392 Q99613	EIF3C	13	S	***MSRFFT G SD S ESS L S G E L V T K P VG	72.31
393 Q9BRS2	RIOK1	21	S	LLMSRV V PGQF D DAD S SD S ENRDLK T V K ED	72.11
394 Q9BTC0	DIDO1	811	S	QEAI P LED S PPV S D S EEQQ E SAR A RAPEK S T	67.34
395 Q9BXJ9	NAA15	856	S	PGYEEDM K ITVNGD S S A EAEELANEI*****	65.47
396 Q9H6Z4	RANBP3	355	S	VSSDANRENAAAES G S ESS SQEA T PEKESLA	65.38
397 Q9NXG2	THUMPD1	86	S	DMYGPEKFTDKDQ Q PS G SEG E DDA E ALKK	64.55

398 Q9UKV3	ACIN1	898	S	GQEAVVDLHADDSRI S EDETERNGDDGTHDK	64.49
399 O60841	EIF5B	589	S	DSGKTLKKPSKEMSS S DSEYDSDDRTKEER	64.38
400 Q5VVJ2	MYSM1	218	S	GRADPNILNAVKIEKL S DDEEVDTDEVDELS	64.35
401 P19338	NCL	41	S	EDSEDEEMSEDEEDDS S GEEVVI P QKKGKKA	64.30
402 Q9UQ35	SRRM2	1497	S	SSPEPKALPQTPRPRS R SPSSPELNKCLTP	64.25
403 Q8IWX8	CHERP	815	S	SRSKSYPGRRRSRS R SPTPSSAGLGSNS	64.18
404 Q9BW71	HIRIP3	372	S	STSGEESDLEREVSD S EAGGGPQGERKNRSS	63.41
405 Q9UQ35	SRRM2	1539	S	DQKTVARTPLGQRSSRGSSQELDVKPSASPQ	63.34
406 Q8WVC0	LEO1	610	S	KGGIREERARIYSSDS D E G SEEDKAQRLLKA	62.38
407 Q13428	TCOF1	771	S	AEKQEDSESSEEESD S EEAAASPAQVKTSVK	62.31
408 Q9UQ35	SRRM2	2046	S	RSRTPLLPRKRSSRS R PLAIRRSRSRTPRT	61.97
409 Q9UQ35	SRRM2	1654	S	GSSSKGRGSPSPEGSS T ESSPEHPPKSRTAR	61.88
410 Q14839	CHD4	105	S	PEFVEEEEVALRSD S EGSDYTPGKKKKKL	61.82
411 Q9UQ35	SRRM2	1552	S	SRSGSSQELDVKPSAS P QERSESDSSPDASKA	61.34
412 Q8WVX3-2	C4orf3	7	S	*****MTSLINS P INRRPLQNVEGNNR	60.81
413 O14974	PPP1R12A	509	S	LAYVAPTI P RRLAST S DIEEKENRDSSLRT	60.67
414 P16383	GCFC2	16	S	MAHRPKRTFRQRAADS S SDSDGAESPAEPGA	60.54
415 P18858	LIG1	141	S	KQLPKRTI Q EVLEE Q S EDEDREAKRKKEEEE	60.37
416 P45973	CBX5	14	S	**MGKKTKRTADSSS E DEE E YVVEKVLDRR	60.28
417 O75822	EIF3J	127	S	EQLADKLRLLQEE S DLELAKE T FGVNNAV	59.57
418 P27815-7	PDE4A	18	S	RSRSALS V AGTGDERS R ETPESDRANMLGAD	59.36
419 P48634	PRRC2A	342	S	AGAHEEV D YTEKLKF S DEEDGRDSDEEGAEG	59.18
420 P51116	FXR2	447	S	RGRGRRTGGPAYGPS S DVSTASETESEKREE	59.15
421 Q13185	CBX3	95	S	QKAGKEKDGTKRKSL S DSESDDSKSKKRDA	59.14
422 Q13427	PPIG	254	S	RKHKEKKRKKS K KS S ASSESEAENLEAQPO	59.09
423 Q13428	TCOF1	85	S	ALQAKKTRVSDPIST S ESSEEEEEAEAE A TAK	59.05
424 Q13435	SF3B2	307	S	EEEEMETDARSSLGQS A SETEEDTVSVSKKE	59.04
425 Q14137	BOP1	127	S	EMASARIGDEYAED S DEEDIRNTVGNVPLE	58.97
426 Q15054	POLD3	307	S	KVLQKEKKRGKRAL S DDET K ETENMRKKRR	58.96
427 Q3YEC7	RABL6	464	S	DQPRGS P PLPAGPVPS Q DITLS S EEEAEVAA	58.94
428 Q5JRA6	MIA3	288	S	LDLKTKFGSTADALV S DDETTRLVTSLEDDF	58.93
429 Q5T200	ZC3H13	1465	S	AHSLGSGAGEGYEPI S DEDELDEILAGDAEKR	58.88
430 Q6IN85	SMEK1	117	S	GKDP S VDTQDLVDE S EEERFDDMSSPGLEL	58.67
431 Q6NSI8-2	KIAA1841	697	S	PRDGT V SKSNRKSG S SVLLMTVLLC*****	58.64
432 Q6WKZ4	RAB11FIP1	206	S	AIIPSTTPSVDSD S VVVKDKKKSKIKTLL	58.61
433 Q86V59	PNMAL1	335	S	QDARAEAE S PGGASE S DQDGGHESPPKKKAV	58.59
434 Q8IY81	FTSJ3	471	S	YVSDVEDDGDDTLS D LDPEELAGVRGHQG	58.46
435 Q8TEA8	DTD1	197	S	SKERNTPRKEDRSASS G AEGDVSSEREP***	58.34
436 Q8WVC0	LEO1	271	S	EEHRHS D EEEQDHK S ESARGSDSEDEVLRM	57.94
437 Q92538	GBF1	1300	S	ATARADAPDAGAQSD S ELPSYHQNDVSLRG	57.83
438 Q96MU7	YTHDC1	308	S	GSDEKKKERKRARG I S PIVFDRSGSSASESY	57.75
439 Q96ST2	IWS1	183	S	PASDSETEDALKPQ I S DSESEEP R HQASDS	57.68
440 Q99613	EIF3C	16	S	MSRFFTTGSDSE S LSGEELVTKPVGGNY	57.64
441 Q9BQ52	ELAC2	212	S	WQSPERPLSRLSPERS S DSESNENE PH PHG	57.63
442 Q9H7L9	SUDS3	45	S	LESAEDDERSCRGRES D EDTEDASETDLAKH	57.39
443 Q9NR30	DDX21	13	S	***MPGKLRSDAGLE S DTAMKKGETLRKQTE	57.34
444 Q9NXX6	NSMCE4A	30	S	RGRDPHRDRTRSRS S RSPLSPRSRRGSARE	56.91
445 Q9NYF8	BCLAF1	389	S	EWEDQEALDYFSD K SGQKFNDSEGDDTEE	56.65
446 Q9UBB9	TFIP11	210	S	SERTTQSMQDFPVVD S EEE A EEFQKELS Q	56.41
447 Q9UH62	ARMCX3	61	S	VDDAGDCSGARYND W S DDDDDSNESKSIVWY	56.34
448 Q9UKD2	MRT04	235	S	DDLPESASESTEESD S EDDD*****	54.12
449 Q9ULX3	NOB1	184	S	HELQELLIDRGEDVP S EEEEEEENG F EDRKD	52.38
450 Q9UQ35	SRRM2	1103	S	SKSQTSPKGGRSSRSS S PVTEL A RSP I RQDR	51.42
451 Q9Y5B6	PAXBP1	155	S	KEDLEKSKIKTELNS S AESEQPLDKTGHVKD	51.29
452 Q9UQ35	SRRM2	1550	S	QRSRSGSSQELDVKPS A SPQERSESDSSPDS	49.56
453 Q9UQ35	SRRM2	2044	S	RSRSRTPLLPRKRSSRS R SPLAIRRSRSRTP	46.29
454 Q5JSH3	WDR44	403	S	IMRRTKEYVSNDAAQS D DEEKLQSQPTDTG	46.25

455 Q6NSI8-2	KIAA1841	698	S	RDGTVSKSNRKSGLSSVLLMTVLLC*****	42.77
456 Q9NXX6	NSMCE4A	32	S	RDPHRDRTRSRSRSPLSPRSRRGSARERR	40.99
457 Q9UQ35	SRRM2	2042	S	RRRSRSRTPLLPRKRSRSRSPLAIRRSRSR	40.94
458 O94806	PRKD3	30	S	LPTAI PAVLPAASPCSSPKTGLSARLSNGSF	40.93
459 Q5T0N5	FNBPI1L	295	S	DFPFEDYSQHIYRTISDGTISASKQESGKMD	40.91
460 P12694	BCKDHA	347	S	RIGHHSTSDDSSAYRSVDEVNYWDKQDHPI	40.88
461 O15027	SEC16A	1172	S	HRDPYGEVDRRSVHSEHSARSLHSAHLAS	40.84
462 P05386	RPLP1	101	S	AAPAAEKVEAKKEESEESDDDMGFGLFD***	39.85
463 Q9UQ35	SRRM2	2032	S	RSRSRTPPAIRRSRSRTPLLPRKRSRSRSP	39.83
464 O15027	SEC16A	1169	S	PDPHRDPYGEVDRRSVHSEHSARSLHSAHS	39.76
465 O43719	HTATSF1	702	S	KEVEDADEKLFFEDDSNEKLFDEEDSSEKL	39.74
466 P18583	SON	1780	S	RSAASPVSSMPERA SESSSEEKKDYEIFVK	39.72
467 P51532	SMARCA4	655	S	EAWLEMNPGYEVAPRS DSEESGSEEEEEEE	39.71
468 Q15019	SEPT2	218	S	IEEHNIKIYHLPDAES DEDEDDEFKEQTRLLKA	39.69
469 Q14498	RBM39	136	S	SRRRSRSKSPFRKDKSPVREPIDNLTPERD	39.67
470 Q7Z4V5	HDGFRP2	264	S	AMARSASSSSSSSSSDSDVSVKPPRGRKP	39.65
471 P08238	HSP90AB1	226	S	YPITLYLEKEREKEIS DDEAEEEKGEKEEED	39.51
472 Q9Y4P1	ATG4B	383	S	SLDSSDVERLERFFDSEDEDFEILS*****	39.48
473 O60284	ST18	690	S	THGKTEEEKEKDPVSS LENLEEKKFPGAEASI	39.44
474 P07900	HSP90AA1	263	S	EEKESEDKPEIEDVGS DEEEEKKDGDKKKKK	39.40
475 Q9H1E3	NUCKS1	144	S	DGSDEDFLMEDDDDS DYGSSKKKNKKMVKK	39.39
476 Q9UQ35	SRRM2	2020	S	RSRSRTSPVTRRSRSRTPPAIRRSRSRTP	39.35
477 O43719	HTATSF1	713	S	EDDDSNELFDEEEDS SEKLFDDSDERGTLG	39.34
478 Q9Y2W1	THRAP3	53	S	SRSRKRLSSRSRSRSY SPAHNERNHPRVY	39.33
479 P07900	HSP90AA1	231	S	YPITLFVEKERDKEVS DDEAEEKEDKEEEKE	39.27
480 Q08945	SSRP1	668	S	SRQLSESFKSKEFVSS DESSSGENKSKKRR	39.22
481 P35269	GTF2F1	217	S	ASELRIHDLEDDLEM SSADASGEEGGRVP	39.18
482 Q9UJX6	ANAPC2	218	S	LDSRYARRYYRLLQSPLCAGCSSDKQQCWC	39.17
483 Q7Z5K2	WAPAL	77	S	PKVEEESTGDPFGFD SDDESLPVSSKNLAQV	39.15
484 Q8N5F7	NKAP	27	S	REASGSRRRSSSKSPKPSKSARS PRGRRS	39.13
485 Q16629	SRSF7	202	S	SRYFQSPRSRSRSRSIS RPRSSRSKRSRSPS	38.64
486 O75400	PRPF40A	933	S	SPKKKTGKDSGNWDT SGSELSEGELEKRRRT	38.63
487 O75494	SRSF10	133	S	RSRSRSYERRRSRSRSFD DNYRYSRSPRNSR	38.58
488 O60284	ST18	689	S	KTHGKTEEEKEKDPVSS LENLEEKKFPGAEAS	38.57
489 O95218	ZRANB2	120	S	GFNERENVEYIEREES DGEYDEFGRKKKKYR	38.53
490 P51532	SMARCA4	657	S	WLEMNPGYEVAPRS DSEESGSEEEEEEE	38.52
491 P24534	EEF1B2	140	S	QYESKKAKKPALVAKS SILLDVKPWDETDM	37.56
492 Q5H9R7	PPP6R3	617	S	EACCKERIQQFDDGGS DEEDIWEEKHIAFTP	37.38
493 Q5F1R6	DNAJC21	423	S	EGVKVDPEDTNLNQDSAKELEDSPQENVVT	37.37
494 Q9H7J1	PPP1R3E	81	S	SAPAGGGARAPRSRSPDTRKRVRFADALGL	37.36
495 Q8YB3	SRRM1	389	S	PPKRTSSPPRKTRRLSPSASPPRRRHPSPPP	37.33
496 P51531	SMARCA2	1572	S	RPNRGKAKPVVSDFDS DEEQDEREQSEGSGT	36.38
497 Q9BYW2	SETD2	1988	S	PINEETPSQDEEEGVSD VESERSQE QPD KTV	36.35
498 Q9Y2W1	THRAP3	51	S	SLSRSRKRLSSRSRSRSY SPAHNERNHPR	36.34
499 Q9UN86	G3BP2	141	S	VHNDMFRYEDEVFGDSEPELDEESEDEVEEE	36.33
500 Q8NDI1	EHBP1	171	S	CIFLREGKATDEDMQS LASIMSMKQADIGNI	35.69
501 O96007	MOCS2	3	S	*****MS SLEI SSSCF SLET KLP	35.33
502 P19338	NCL	28	S	DPKKMAPPPKEVEEDS EDEEMSEDEEDDSSG	34.52
503 O75475	PSIP1	275	S	KRKNLAKTGVTS TSDSEEEGDDQEGEKKRKG	31.30
504 P49321	NASP	503	S	DKTEEMPND SVLENKS LQENE EEEIGNLELA	31.29
505 Q8TDD1	DDX54	75	S	GPGRPLPTFP TSECTS DVEPD TREVRAQNK	31.28
506 O75400	PRPF40A	935	S	KKKTGKDSGNWDT SGSELSEGELEKRRRTLL	30.33
507 P08238	HSP90AB1	255	S	EDKDDEEKPKIEDVGS DEEDDS GKD KKKT	30.32
508 P05386	RPLP1	104	S	AEEKKVEAKKEESEES DDDMGFGLFD*****	30.31
509 O96007	MOCS2	2	S	*****MS SLEI SSSCF SLET KLI	30.26
510 Q9NW75	GPATCH2	284	S	GLFTNDEGRQGDDEQS DWFYEKESGGACGIT	30.25
511 O94913	PCF11	509	S	SPKRRDRSPKRRQRSMSPTSTPKAGKIRQS	30.24

512 O15027	SEC16A	1175	S	PYGEEVDRRSVHSEHSARSLHSAHSLASRRS	29.54
513 P35269	GTF2F1	218	S	SELRIHDLEDDLEMSDASDASGEGGRVPK	28.51
514 P67809	YBX1	176	S	NYQNSESGEKNEGSESAPEGQAQQRPYRRR	27.34
515 Q96HR8	NAF1	315	S	SWKNDQEPPPEALDFS DDEKEKEAKQRKKSQ	26.48
516 Q13428	TCOF1	935	S	GPSAAQAGKQDDSGSS SEESDS DGEAPA A VT	26.47
517 Q96B97	SH3KBP1	230	S	FGDIFKDPIKL RPRS IEVENDFLPVEKTIG	18.00
518 P24534	EEF1B2	141	S	YESKKAKKP ALVAKSS ILLDVKPWDDETDMA	13.88
519 Q96IZ7	RSRC1	121	S	RRSRSPRRLRSHRS S ERSSHRRTRRS RSRDR	13.56
520 Q92934	BAD	118	S	NLWAAQRYGRE LRRMS DEFVDSFKKGLPRPK	13.49
521 Q9NRW4	DUSP22	58	S	PMLEGVKYL CIPAADS PSQNLTRHF KESIKF	12.40
522 Q8N108	MIER1	160	S	QEIIIRPRRCKYFD TN SEVEEESE EDEDYIIPS	11.25
523 Q9H6Z4	RANBP3	96	S	PEAQ LPPF PRELAGRS AGGSS PEGGEDSDRE	10.18
524 Q04656	ATP7A	1473	S	NYSRASINSLLSDKRS LNSVVTSEPDKHSSL	9.63
525 Q12872	SFSWAP	866	S	KKKR RRSRTKSKARS QSVPSKQ AAPRPAA	8.94
526 Q5SW79	CEP170	1112	S	PTRTSLLR RARLGEA SDSELADADKA VASE	8.69
527 Q9BRS2	RIOK1	22	S	LMSRVVPGQFDDADSS DSEN RDLKTVKEKDD	8.65
528 Q8NC51	SERBP1	394	S	GGRP NRG SRTDKSSA SAPDVDDPEAFPALA*	8.63
529 P49792	RANBP2	2900	S	VGTQSAGKV GEDEDGS DEEVVHNEDIHF EPI	8.11
530 P98175	RBM10	60	S	PREYGSQEGKHDYDD SSEE QSAE DSYEAS PG	7.71
531 Q5THJ4	VPS13D	1042	S	SFDIPTGSLRDSRAQS PVSGPNVAH LTDGAT	7.49
532 Q96IZ7	RSRC1	120	S	TRRSRSRPLRSHRS S ERSSHRRTRRS RSD	7.47
533 Q9UKJ3	GPATCH8	352	S	KSSDQGLQKVGDSDG SS NLDGK KEDED P QDG	7.45
534 P67809	YBX1	174	S	QQNYQNSESGEKNEG SESAPEGQAQQR PYR	7.42
535 Q9H7J1	PPP1R3E	79	S	ARSAPAGGGGARAPR S RSPDTRK RVRFAD AL	7.38
536 Q9UQ35	SRRM2	250	S	KSKRKSKDKKR KRSR S TTPAPK S RRA HRSTS	7.37
537 Q9UQ35	SRRM2	957	S	TSRTT PRRSR SVSPCS NVESRLL PRY SHSGS	7.23
538 Q9Y6X9	MORC2	743	S	KLSPATPSRKR SVAV S DEEEVEEEAERRKER	7.19
539 Q92576	PHF3	1133	S	CKICIGRM APPVDDLS PPKV KVVG VARK HS	7.01
540 P06748	NPM1	112	S	PVVLRLKCGSGPVHIS QQLV LVA VEEDA ESED	6.97
541 P51858	HDGF	132	S	EAAEGD GDKKGNAEG S DEEGKLVIDEPAKE	6.90
542 Q8IXT5	RBM12B	280	S	RKRSHSKSPR RTR SRS PLGFYVHLKNL SLSI	6.74
543 P23588	EIF4B	409	S	PKLERRPRERHPSWR S EETQERERS RTGSES	6.67
544 O15541	RNF113A	47	S	KRPACDPEPGESGSS S DEGCTVVRPEKKR VT	6.54
545 O95382	MAP3K6	916	S	LGDPFLQPGKRSRSP S SPRHAPR PSDAPS AS	6.51
546 Q12789	GTF3C1	1068	S	VRCPRV RKN SSTDQGS DEEGSLQKEQESAMD	6.48
547 Q8IYB3	SRRM1	530	S	GEVGR RRRHPSRS A SPS PRKR QKETSPR GR	6.40
548 Q8N5F7	NKAP	24	S	SPDREASGSG RRRSS SKSPKPSKSARS PRG	6.39
549 Q9H1E3	NUCKS1	130	S	EEQEEEDEAPFQEKD SG S DEDFLM EDD DSD	6.36
550 Q96ST2	IWS1	438	S	DAVSDKSGKREKTIAS DSEEEAGKELSDKKN	6.34
551 Q13185	CBX3	176	S	PQIVIAFYEE RL TWH S CPEDEAQ*****	6.32
552 Q8N9T8	KRI1	93	S	PRIYQK DATFYNRTA SSSDSEEDPEALEKQK	6.31
553 Q8N9T8	KRI1	94	S	RIYQK DATFYNRTA SSSDSEEDPEALEKQK	6.24
554 Q9UQ35	SRRM2	741	S	SSSERK NKSRTS QRR SRSN S SPMKKS RISS	6.23
555 Q96ST2	IWS1	400	S	EEEKVA KRKA AVLSD S DEEKA SAKK SRV VS	6.16
556 Q9C073	FAM117A	213	S	PSGSPV LRLSPCLHRS LEGLNQE LEEV FVKE	6.14
557 P45973	CBX5	12	S	*****MGKKTKRTAD S S SDEE EYV VEV KVL D	6.12
558 P19338	NCL	42	S	DSEDEEMSEDEEDDS GEEV V I P QKKG KAA	6.10
559 P45973	CBX5	11	S	*****MGKKTKRTAD S S SDEE EYV VEV KVL	6.03
560 Q96ST2	IWS1	415	S	SEDEEKASAKK SRV VS DADDSD DAVS DKSG	5.79
561 P26373	RPL13	139	S	LILFPRKPSAPKKGD S SAEELK IAT QLT GPV	5.72
562 Q12872	SFSWAP	868	S	KRRSRSP RSKARS QSVSPSKQ AAPRP A PA	5.57
563 Q5BK Y9	FAM133B	196	S	RKMYSED KPLS S S ESEYIEEVRA KKK	5.54
564 Q5BK Y9	FAM133B	194	S	KKRKMYSED KPLS S S ESEYIEEVRA KKK	5.44
565 Q6PD62	CTR9	1037	S	DEDKLKIADEGHPRNS NSNSD S D E D E Q R K K C	5.43
566 Q8N108	MIER1	166	S	RRCKYFD TN SEVEEE S EED E DYI P S E DWK K E	5.34
567 Q92769	HDAC2	422	S	SIRASDKRIACDEEF S DSEDEGE GEG RRN VAD	5.02
568 P98175	RBM10	736	S	RPSPPRGLVAAYS GE S DSEEEQERGGP EREE	4.92

569 Q5SNT6	FAM21B	451	S	NLKPSSETKTQKGLF\$DEEDSEDLFSSQSAS	4.88
570 Q8IYB3	SRRM1	463	S	RESPSPAPKPRKVEL\$ESEEDKGKMAAADS	4.74
571 Q13428	TCOF1	349	S	KPEEDSESSSEESSDSEEETPAAKALLQAKA	4.69
572 P46937	YAP1	61	S	QAPPAGHQIVHVRGDSSETDLEALFNAVMNPK	4.68
573 Q06265	EXOSC9	306	S	QRITAFKMEKAPI\$DVEEKAEI\$IAEAEP	4.67
574 Q8IYB3	SRRM1	532	S	VGRRRRHSPRSRASP\$PRKRQKETS\$PRGR\$R	4.61
575 Q9BW71	HIRIP3	357	S	GEPTAKGSRKMARLGS\$TSGEESDLEREVSDS	4.56
576 Q9NYF8	BCLAF1	198	S	SQEEPKDTEFHDPS\$IDEFNKSATSGDIW	4.46
577 O15027	SEC16A	164	S	HRPASALVNPLARGD\$PENRTHHPLGAGAGS	4.43
578 Q9UQ35	SRRM2	782	S	SKAKSRLSLRRSLSG\$SPCPKQKSQT\$PPRRS	4.42
579 Q15185	PTGES3	148	S	DEDVDLPEVDGADDS\$QDS\$DEKMPDLE***	4.41
580 Q6ZMG9	CERS6	346	S	SRGKVSKDDRS\$DIES\$SDEED\$PPGKNPHT	4.35
581 Q96IZ7	RSRC1	118	S	SRTRRSRSRPRLRSH\$RSSERSSHRRTRSRS	4.29
582 Q9BW71	HIRIP3	87	S	KLDLTKKGKRPPTPCS\$DPERKRFN\$ESES	4.28
583 Q9UKJ3	GPATCH8	349	S	PDEKSSDQGLQKVGD\$DGSSNLDGK\$EDED\$P	4.27
584 Q13428	TCOF1	934	S	TGPSAAQAGKQDDSG\$SSEE\$SDGEAPAAV	4.25
585 Q96ST2	IWS1	440	S	VSDKSGKREKTIASD\$EEEAGKEL\$DK\$KNEE	4.15
586 Q9BWU0	SLC4A1AP	466	S	RKRKAKNWEDEDFYD\$DDDTFLDRTGLIEKK	4.12
587 Q9C0C2	TNKS1BP1	1620	S	HLFQDSTEPRASRVPS\$DEEVVEEPQSRRTR	4.01
588 P43487	RANBP1	60	S	EEELFKMRAKLFRFA\$ENDLPEWKERGTG\$DV	3.92
589 Q8IYB3	SRRM1	528	S	KN\$EVGRRRHSPRS\$ASP\$PRKRQKET\$PR	3.92
590 Q13428	TCOF1	346	S	KAGKPEED\$E\$S\$EE\$SD\$SEEETPAAKALLQ	3.88
591 Q92769	HDAC2	394	S	GVQM\$QAI\$EDAVHED\$G\$DE\$GED\$E\$PD\$KRISIR	3.84
592 O15042	U2SURP	800	S	HEE\$EEE\$ENQNQ\$EEE\$E\$DE\$E\$DTQ\$SS\$KE\$HH	3.83
593 P49959	MRE11A	689	S	IMSQS\$QVSKGVDFESS\$E\$DD\$DD\$PFMNTSSLR	3.82
594 Q99590	SCAF11	830	S	QSPSPRRETGKESRK\$QSPSPK\$NESAR\$GRKK	3.81
595 Q9UQ35	SRRM2	248	S	TPKSKRK\$KDK\$KR\$KRS\$R\$TT\$PAP\$K\$RRA\$HRS	3.80
596 O95382	MAP3K6	912	S	AQTLLGDPFLQPGKRS\$RSPSS\$PRHAPR\$DA	3.78
597 P07910	HNRNPC	260	S	DET\$NVKME\$EGG\$ADD\$AEE\$G\$D\$LL\$DD\$DN\$EDR	3.75
598 Q96ST2	IWS1	398	S	E\$G\$EEE\$KVA\$KRKA\$AVL\$SD\$E\$DE\$E\$KASAK\$SRV	3.74
599 Q9UQL6	HDAC5	611	S	EEE\$E\$DCI\$QVK\$DE\$E\$G\$SGA\$EE\$GP\$D\$LE\$E\$PGAGY	3.73
600 Q9UQ35	SRRM2	745	S	R\$KN\$K\$RT\$S\$Q\$R\$RS\$R\$S\$NS\$SPE\$MK\$K\$RI\$SS\$R\$R	3.71
601 Q9Y5S9	RBM8A	168	S	GPPKGK\$RRGG\$RR\$RS\$SP\$DR\$RR\$*****	3.69
602 P35269	GTF2F1	221	S	RIHDLE\$D\$DLEM\$SS\$DA\$DAS\$GEE\$GG\$RVP\$K\$AK\$K	3.68
603 P78317	RNF4	95	S	ARRLPQDHAD\$CSV\$VS\$D\$DE\$E\$LS\$R\$D\$R\$D\$V\$Y\$V\$T\$T	3.67
604 Q92625	ANKS1A	661	S	ESLSNC\$SIG\$KK\$R\$LE\$K\$PS\$FA\$E\$W\$D\$E\$IE\$K\$IM\$S	3.66
605 Q99590	SCAF11	834	S	PR\$RET\$GKESRK\$SQ\$SP\$PK\$NES\$AR\$GR\$K\$K\$RS\$Q	3.65
606 Q9UKN8	GTF3C4	19	S	ADQAR\$VGP\$ADD\$GP\$AP\$SGEE\$EG\$GG\$EAG\$G\$K\$E	3.64
607 Q9UQ35	SRRM2	715	S	GR\$RS\$RS\$LV\$RR\$GR\$SH\$S RTP\$Q\$RR\$G\$G\$SS\$ER	3.63
608 P35579	MYH9	1943	S	FVV\$P\$R\$M\$RK\$G\$A\$G\$D\$G\$S\$D\$E\$E\$V\$D\$G\$K\$A\$G\$A\$E\$K\$P	3.62
609 P42166	TMPO	67	S	PL\$P\$AG\$T\$N\$K\$G\$P\$P\$DF\$S\$D\$E\$E\$R\$P\$T\$P\$V\$L\$G\$G\$A\$A	3.61
610 Q08945	SSRP1	672	S	SE\$F\$K\$S\$K\$E\$F\$V\$S\$D\$E\$S\$S\$G\$E\$N\$K\$S\$K\$K\$R\$R\$R\$S\$E\$D	3.60
611 Q15459	SF3A1	359	S	SQL\$Q\$DT\$Q\$V\$Q\$DM\$DE\$G\$S\$D\$EE\$E\$G\$Q\$K\$V\$P\$P\$P\$E\$T	3.60
612 Q66PJ3	ARL6IP4	332	S	Q\$VE\$AL\$P\$G\$P\$S\$LD\$Q\$W\$H\$R\$S\$A\$G\$E\$E\$D\$G\$P\$V\$LT\$D\$E\$Q\$K	3.59
613 Q86US8	SMG6	1196	S	D\$V\$V\$IE\$D\$FE\$D\$S\$E\$A\$E\$G\$S\$G\$G\$E\$D\$D\$I\$R\$E\$L\$R\$A\$K\$K\$LA	3.58
614 Q8TDR0-2	TRAF3IP1	359	S	TS\$K\$R\$R\$S\$K\$N\$S\$V\$E\$G\$D\$S\$T\$DA\$E\$G\$D\$A\$G\$P\$A\$G\$Q\$D\$K\$S\$E	3.57
615 Q96JC9	EAF1	165	S	K\$P\$P\$V\$G\$P\$K\$T\$S\$P\$K\$D\$N\$P\$S\$P\$E\$P\$Q\$L\$D\$D\$K\$R\$E\$L\$R\$A\$E	3.56
616 Q9H792	PEAK1	730	S	R\$G\$Q\$S\$S\$P\$Q\$R\$S\$Y\$S\$S\$H\$S\$S\$PA\$K\$I\$Q\$R\$A\$T\$Q\$E\$P\$V\$A\$K\$I	3.55
617 O15021	MAST4	1831	S	P\$Q\$A\$S\$K\$T\$E\$L\$P\$S\$P\$E\$A\$Q\$S\$P\$S\$P\$G\$D\$V\$R\$A\$S\$V\$P\$P\$V\$L	3.54
618 Q9H0E3	SAP130	300	S	I\$P\$P\$A\$V\$V\$V\$A\$T\$V\$A\$T\$R\$A\$Q\$S\$P\$V\$IT\$T\$A\$A\$H\$A\$T\$D\$S\$A\$L	3.54
619 Q9Y2X3	NOP58	502	S	K\$K\$R\$G\$K\$K\$K\$H\$K\$IE\$E\$P\$S\$E\$E\$E\$P\$C\$T\$S\$T\$A\$A\$S\$P\$E\$K	3.53
620 P08621	SNRNP70	320	S	R\$KE\$E\$L\$R\$G\$G\$G\$D\$M\$A\$E\$P\$S\$E\$A\$G\$D\$A\$P\$D\$D\$G\$P\$P\$G\$E\$L	3.53
621 Q9NXX6	NSMCE4A	35	S	H\$R\$D\$R\$T\$R\$S\$R\$S\$R\$S\$P\$S\$P\$R\$S\$R\$G\$A\$R\$E\$R\$R\$E\$A\$P	3.52
622 Q58EX2	SDK2	2026	S	Y\$T\$R\$S\$P\$P\$R\$P\$S\$P\$G\$S\$L\$H\$Y\$S\$E\$D\$E\$V\$T\$K\$Y\$N\$D\$L\$I\$P\$A\$E\$S	3.52
623 P18583	SON	1782	S	A\$A\$P\$V\$V\$S\$S\$M\$P\$E\$R\$A\$S\$E\$S\$S\$E\$E\$K\$D\$D\$Y\$E\$IF\$V\$K\$V\$K	3.52
624 P19338	NCL	145	S	PA\$K\$G\$A\$K\$G\$K\$N\$A\$K\$K\$E\$D\$S\$D\$E\$E\$D\$D\$S\$E\$E\$D\$E\$D\$D	3.52
625 Q9UQ35	SRRM2	454	S	E\$S\$P\$K\$P\$A\$P\$A\$P\$G\$H\$R\$E\$I\$S\$S\$P\$T\$K\$N\$R\$S\$H\$G\$R\$A\$K\$R	3.51

626 P19338	NCL	153	S	KNAKKEDSDEEEEDDSSEEDDEDDDEDED	3.50
627 Q9UKJ3	GPATCH8	491	S	PGSKAEAKKALGGDVS DQSLESHSQKVSETQ	3.49
628 P85037	FOXK1	249	S	LRSMVSPVPSPTGTIS VPNNSCPASPRGAGSS	3.48
629 P98175	RBM10	738	S	SPPRGLVAAYSGESD SEEQERGGPEREEKL	3.47
630 Q9UPP1	PHF8	854	S	DSLGAFCFDAEYIYPS LESDDDDPALSRPK	3.46
631 Q9UQ35	SRRM2	952	S	SPSRVTSRTTPRSRS SVSPCSNVESRLLPY	3.46
632 Q15154	PCM1	65	S	KKFGVESDKRVNTND IPESSPGVGRRRTKTP	3.45
633 Q13428	TCOF1	583	S	QEKSILGNILQAKPTSS PAKGPPQKAGPVAVQ	3.44
634 Q14677	CLINT1	234	S	FRRKDREDS PERCS SDDEEKKARRGRSPKG E	3.44
635 Q96ST2	IWS1	289	S	SDSESEDPPRNQASD SENELPKPRVSDSES	3.43
636 Q9UQ35	SRRM2	746	S	KNKSRTSQRRSRNS NSPEMKRSRISSRRSRS	3.42
637 P51003	PAPOLA	24	S	GSQQTQPPQKHYGIT SPISLAAKETDCVLT	3.42
638 P98174	FGD1	48	S	SDPGASEPGLLARRGS GSAALGGPLDPQFVGP	3.42
639 O43432-3	EIF4G3	232	S	KQEEKPKPD PVLKSPSPVRLVLSGEKKEQE	3.41
640 O15541	RNF113A	46	S	RKRPA CDPEPGESGSSD E DEGCTVVRPEKKRV	3.40
641 Q14684	RRP1B	458	S	VAEPGA EATSTGEE S GSEHPPAVPMHNKRK	3.39
642 P36507	MAP2K2	293	S	IFGRPVVD GEEGEPHSISPRPRPPGRPVSGH	3.37
643 Q9P219	CCDC88C	227	S	DLT QERDYLQAQHPPS PIKSSAD STPSPTS	3.36
644 Q6AWC2	WWC2	1022	S	PGERNQYICRNL RNSDSDS STLAKKSLFVRNS	3.33
645 Q9BVV6	KIAA0586	376	S	RGNVRLLE QILNNNDSL TRKSESSNTTSLTR	3.29
646 P46937	YAP1	164	S	SGPAATPTAQHL RQSSFEIPDDVPLPAGWEM	3.29
647 P85037	FOXK1	243	S	HIPEDPL RSMVSPVPSPTGTISVPNSCPASP	3.28
648 P46100	ATRX	1352	S	RHRLLRHKLTVSD GE EEKKTKPKEHKEVK	3.27
649 Q7L4I2	RSRC2	222	S	IEKPRRF RSRSLSRTPSP PPFRGRNTAMDAQE	3.26
650 O95218	ZRANB2	153	S	AVGPASILKE VEDKES E GE EEEDDEDLSKYK	3.25
651 Q01831	XPC	884	S	SEAAAPHTDAGGGL SDEEEGTSSQAEAARI	3.25
652 Q13428	TCOF1	1153	S	KAPESSD SEDSSDSSGSEEDGE GPQGAKS	3.25
653 Q8IYB3	SRRM1	874	S	QEEPVAAP EPKKETESEAEDNLDDLEKHLRE	3.25
654 Q9UQ35	SRRM2	743	S	SERKNKSR TSQRRSRSNSSPEMKRSRISSRR	3.25
655 P29692	EEF1D	162	S	TPAEDDEDD DIDLFGSDNEEDKEAAQLREE	3.25
656 Q86US8	SMG6	1191	S	EDEEEDV VIEDFEEDSE AE ^{GSGGEDDIR} ELR	3.24
657 Q8N5S9	CAMKK1	458	S	SWTTVILVK MSLRKRSFGNPFEQARREERS	3.24
658 Q92882	OSTF1	213	S	VRTLSNAEDYLD DEDS D*****	3.23
659 Q5BKY9	FAM133B	191	S	GLSKKRK MYSEDKPLSSESLS ESEYIEEVRA	3.22
660 Q6UN15	FIP1L1	89	S	PKPKV TETEDDDSDSDS DDDED DVHVHTIGDIK	3.21
661 Q96ST2	IWS1	315	S	SDSE SEGPQKGPASD E TEDASRHKQKPESD	3.21
662 O00567	NOP56	520	S	KPKKK KSFSKEELMS SD LEETAGSTSIPKRK	3.20
663 O94842	TOX4	178	S	LPPAQ SPEDRLSTTPSP SSLHEDGVDFRR	3.20
664 P53999	SUB1	17	S	PKSKELV SSSSSGSDSD SEVDKKLKRKKQVA	3.19
665 P67809	YBX1	314	S	DGKETKA ADPPAENS SAPEAEQGGAE*****	3.19
666 P78317	RNF4	94	S	NARRLPQDHAD SCVVSSDDEELSRDRDVYVT	3.19
667 Q16629	SRSF7	204	S	YFQSPSR SRSRSRSIS RPRSSRSKSRSPSPK	3.19
668 Q5BKY9	FAM133B	82	S	ENWKKE LEKHREKLLS GESSSSKRQRKKKE	3.19
669 Q9UQ35	SRRM2	566	S	RGRSR SARRGRSHRSRSPATRGRRSRSRTPARR	3.18
670 Q96ST2	IWS1	313	S	RVSD SEGPQKGPASD E SETEDASRHKQKP E	3.18
671 Q9UQ35	SRRM2	573	S	RRGRSHSRSPATRGR SRSRTPARRGRRSRSRT	3.18
672 Q5UIP0	RIF1	1454	S	QKKERRKE EEKPLQKSPLHIKDDVLPKQKLI	3.17
673 Q86X95	CIR1	202	S	GRNL TANDPSQEYVA SE GEEDPEVEFLKSLT	3.16
674 O14654	IRS4	1231	S	PPRSRRV PRPEREDSD ND DDTHVRMDFARR	3.15
675 P51825	AFF1	206	S	DSAPEREL SPLISLPS PVPLSPIHSNQQT	3.15
676 Q13428	TCOF1	967	S	AQVIK PPLIFVDPNRSP AGPAATPAQAAAS	3.15
677 Q9BW71	HIRIP3	159	S	DEERQRDL PAQRGEESS EEEKGYKGKTRKK	3.15
678 Q9UDT6	CLIP2	294	S	FAPIHK VIRIGFPSTS PAKAKTKRMAMGVS	3.15
679 Q9UGP8	SEC63	576	S	AVKEDEEEV SDKGSD SEEETNRDSQSEKDD	3.15
680 P23588	EIF4B	424	S	SEETQERERSRTG SESSQGTSTSSRNARR	3.15
681 Q9UKM9	RALY	135	S	RDDFYDRLFDYRGRL SP VPVPRAVPVKRPRV	3.15
682 P25788	PSMA3	250	S	EAEKYAKS ELKEEDES DDDNM*****	3.14

683 Q01130	SRSF2	189	S	SSVSRSRSRSRSRSRSRSRSPPPVSKRESKSRS	3.14
684 Q08945	SSRP1	671	S	LSESFKSKEFVSSDESSSGENSKKKRRSE	3.14
685 Q9C0C2	TNKS1BP1	1666	S	LSPSALKAKLPRNRSAEEGELAESKSSQKE	3.14
686 Q9UQ35	SRRM2	575	S	GRSHSRSPATRGRRSRSRTPARRGRSRRTPA	3.14
687 O43524	FOXO3	253	S	PDGGKSGKAPRRRAVSMDNSNKYTKSRGRAA	3.14
688 O95831	AIFM1	116	S	GLGLTPEQKQKKAALSASEGEEVQPDKAPSH	3.14
689 P12268	IMPDH2	416	S	FFSDGIRLKKYRGMGSLDAMDKHLSSQNRYF	3.14
690 Q6P6C2	ALKBH5	361	S	ENRRSVLLPTHRRGFSSENWYRKSYESSE	3.14
691 Q8IYB3	SRRM1	616	S	QRRYSPSPPKRRTASPPPPPKRASPSPPP	3.13
692 Q96ST2	IWS1	287	S	RISDSESEDPPRNQASDSENEELPKPRVSDS	3.12
693 O14654	IRS4	1107	S	FFAAARAASAFPTDSLERDLSPPSAPAVAS	3.11
694 P18583	SON	1784	S	SPVVSSMPERASESSSEEKDDYEIFVKVKDT	3.11
695 P21127	CDK11B	283	S	EKMEERDILSLQDISDSERKTSAAESSSAE	3.10
696 P51114	FXR1	413	S	GYGTNSELNPSETESERKDELSDWLAGED	3.10
697 P53999	SUB1	13	S	***MPKSKELVSSSSGSDSDSEVDKLRK	3.07
698 P85037	FOXK1	257	S	PSPTGTISVPNSCPASPRGAGSSSYRFVQNV	3.05
699 P98175	RBM10	61	S	REYGSQEKGHDYDDSSEEQSAEDSYEAPGS	3.05
700 Q13428	TCOF1	548	S	TPSAQVGKWEEDSESSSEESSDSDGEVPTA	3.04
701 Q2NKX8	ERCC6L	1028	S	VVVAKIRSKARRIVSDGEDEDDSFKDTSSI	3.02
702 Q7Z4V5	HDGFRP2	299	S	LPKPRGRKPKPERPPSSSSDSDSEVDRIS	3.02
703 Q96ST2	IWS1	511	S	LEEKGETQVKEAEDSDSDDNIKRGKHMDFL	2.99
704 Q9BW71	HIRIP3	160	S	EERQRDLPAQRGEESSEEEKGYKGKTRKKP	2.98
705 Q9NR09	BIRC6	480	S	DIPKLEGSDDLLLEDSDSEEHRSRSDSVTGHT	2.97
706 Q9UGP8	SEC63	574	S	EAALKEDEEEVSDKGSDSEEEETNRDSQSEK	2.97
707 Q9UPT8	ZC3H4	92	S	GGPERSRKEKGKHHSDSDEEKSHRLKLRK	2.97
708 Q9UQ35	SRRM2	629	S	RSRSRTPARRSRTRSPVRRRSRSRSPARRS	2.96
709 Q9Y4F5	CEP170B	1135	S	PTRASRLRRARLGASDTEAADGERGSLGNP	2.95
710 Q96ST2	IWS1	167	S	GKHPASDSEIEELQKSPASDSETEDALKPQI	2.94
711 Q04727	TLE4	269	S	LVVDVSNEDPSSPRGSPAHSPRENGLDKTRI	2.94
712 Q07157	TJP1	277	S	RDERATLLNVPDLSDSIHSANASERDDISEI	2.94
713 Q13428	TCOF1	941	S	AGKQDDSGSSSEESDSDGEAPAATSAQVIK	2.94
714 Q53EL6	PDCD4	78	S	LRKNSSRDSGRGDSVSDSGDALRSGLTVP	2.93
715 Q6UN15	FIP1L1	87	S	GVPKPKVTETEDDSDSDDDDEDVHVTIGD	2.92
716 Q7L1Q6	BZW1	413	S	KKFVEWLKNAEESESEAEEGD*****	2.92
717 Q7LBC6	KDM3B	822	S	RQDSSTSNTSDLSDLDSEEEQLQAKTGLKGI	2.91
718 Q92974	ARHGEF2	947	S	SPEERLQDSSDPDTGSEEEGSSRLSPPHSPR	2.91
719 Q9HCK8	CHD8	2071	S	RSVPPVKLEDEDDSDSELDLSKLSPSSSS	2.91
720 Q9NVU0	POLR3E	162	S	AKHREREAAANEAGDSSQDEAEDDVKQITVRF	2.90
721 Q9Y6M1	IGF2BP2	161	S	ENYSFKISYIPDEEVSSPSPPQRAQRGDHSS	2.89
722 O00567	NOP56	569	S	SGSKKKRKFSKEEPVSSGPEEAVGKSSKKK	2.89
723 P49959	MRE11A	688	S	KIMSQSQVSKGVDFFESSEDDDDPFMNTSSL	2.88
724 Q15345	LRRC41	145	S	EAFFSHVLRGTTDVSSDRRLCDQRFSPLLHS	2.87
725 O75179	ANKRD17	2047	S	PTAKEHYPSSPSSPSPPAQPGGVRSRNSPLD	2.87
726 P51114	FXR1	409	S	NYTSGYGTNSELNPSETESERKDELSDWSL	2.87
727 Q06587	RING1	229	S	GTGGGGTGGVGGGAGSEDGDRGGTLGGGTL	2.86
728 Q9NVU0	POLR3E	161	S	DAKHREERAANEAGDSSQDEAEEDDVKQITVR	2.85
729 Q13428	TCOF1	342	S	ASQTKAGKPEEDSESSSEESSDSEEETPAAK	2.84
730 Q9UDY2	TJP2	1159	S	KAPSRPYQDTRGSYGSDAEEEEYRQQLSEHS	2.84
731 Q9Y2U5	MAP3K2	163	S	RKKRLSIIGPTSRDRSSPPPGYIPDELHQVA	2.84
732 Q8TF01	PNISR	408	S	SGDSEDERSDRGSESSDTDDEELRHRIRQKQ	2.84
733 O60832	DKC1	453	S	VAEAAKTAKRKRESESSEDETTPAAPQLIKK	2.84
734 O95218-2	ZRANB2	310	S	SGDRKKRTRRSRSPESQVIGENTKQP*****	2.84
735 P46100	ATRX	1442	S	RIKVQEDSSSENKSNSBBBBBEEEEE	2.83
736 Q6PKG0	LARP1	631	S	MDGRKNTFTAWSDEESDYEIDDRDVNKILIV	2.83
737 Q7Z406	MYH14	1969	S	TRTVRQVFRLEEGVASSDEEAEAQPGSGGPSP	2.83
738 Q92974	ARHGEF2	941	S	ERQELGSPEERLQDSSDPDTGSEEGSSRLS	2.82
739 Q96ST2	IWS1	172	S	SDSEIEELQKPASDSETEDALKPQISDSES	2.81

740 Q9C0C2	TNKS1BP1	1621	S	LFQDSTEPRASRVPS\$DEEVVEEPQSRRTRM	2.81
741 Q9HAS0	C17orf75	18	S	PSLQESMDGDEKELESSEEGGSAEERRLEPP	2.81
742 Q9UQ35	SRRM2	713	S	RRGRSRSLVRRGRSHSRTPQRGRSGSSS	2.81
743 Q9UQ35	SRRM2	455	S	SPKPAPAGSHREISSPTSKNRSHGRAKRD	2.80
744 O96007	MOCS2	8	S	*****MSSLEISSCFSLETKLPLSPPL	2.80
745 P18583	SON	1931	S	KTVRARSRTPSRRRS\$HTPSRRRSRVGRR	2.80
746 Q01130	SRSF2	220	S	RSKSPPKSPEEEGAVS\$*****	2.80
747 Q92769	HDAC2	424	S	RASDKRIACDEEFSD\$DEDEGEGGRRNVADHK	2.80
748 Q9UKJ3	GPATCH8	353	S	SSDQGLQKVGDSDGS\$NLDGKKEDDPQDGG	2.79
749 Q9UPT8	ZC3H4	94	S	PERSRKEKGKHHSD\$DEEKSHRRLKRKRKK	2.78
750 Q9UQ35	SRRM2	954	S	SRVTSRTTPRRRSRSVSPCSNVESRLLPRYSH	2.77
751 Q9Y3T9	NOC2L	672	S	KDEDRKQFKDLFDLNS\$SEEDTEGFSERGIL	2.76
752 P53999	SUB1	15	S	*MPKSKELVSSSSSG\$SDSDSEVDDKLKRKKQ	2.76
753 Q9BW61	DDA1	95	S	EGERSSAPPRKVARTD\$PDMHEDT*****	2.75
754 Q9BWU0	SLC4A1AP	82	S	SQDLSGDFKKPALPV\$PAARSKAPASSSSNP	2.75
755 Q9P206	KIAA1522	669	S	TLTPLQESPVLISKDQ\$PPPSPPPSYHPPPPP	2.74
756 O15021	MAST4	1825	S	IALLSGPQASKTELPS\$PESAQSPSPSGDVRA	2.73
757 Q8IYB3	SRRM1	465	S	SPSPAPKPRKVELSE\$SEEDKGGKMAADSVQ	2.72
758 Q9UGP8	SEC63	570	S	VVGNEAAVKEDEEEVS\$DKGSDEEETNRDS	2.71
759 P55081	MFAP1	52	S	KRYVSGKRPDYAPMES\$DEEDEEFQFIKKAK	2.71
760 Q5M9Q1	NKAPL	140	S	RERIGELGAPEVWGPS\$PKFPQLDSDEHTPVE	2.71
761 Q9UPP1	PHF8	857	S	GACFKDAEYIYPSLES\$DDDDPALKSRPKKKK	2.71
762 P46783	RPS10	146	S	AVPPGADKKAEAGAG\$ATEFQFRGGFGRGRG	2.69
763 Q01130	SRSF2	191	S	VSRSRSRSRSRSRS\$PPPVSRESRSRS	2.68
764 Q01831	XPC	883	S	KSEAAAPHTDAGGGL\$SDEEEGTSSQAEAAR	2.67
765 Q15345	LRRC41	144	S	MEAFFSHVLRGTIDV\$SDRRLCDQRFSPLLH	2.67
766 Q99871-2	HAUS7	391	S	RSGMILLQVVMAMAVADT\$AKAVETVKKQQGEQI	2.67
767 Q8IYB3	SRRM1	560	S	RRRRSPSPPPTRRRRS\$PSPAPPRRRTPTP	2.67
768 Q96DR7	ARHGEF26	222	S	QKSSSEQKLPLQRLPS\$QENELLENPSVVLST	2.67
769 Q96ST2	IWS1	27	S	DDGGATPVQDERDSGS\$DGEDDVNEQHGSDT	2.67
770 P35269	GTF2F1	224	S	DLEDDLEMSDDASDAS\$GEEGGRVPKAKKKAP	2.66
771 P18583	SON	2009	S	TPSRRRSRSRSVRRRS\$FSISPVRLLRSRTPL	2.65
772 P55081	MFAP1	53	S	RYVSGKRPDYAPMES\$DEEDEEFQFIKKAKE	2.65
773 Q7L014	DDX46	804	S	NERKKLQKAALGLQDS\$DDEDAAVDIDEQIES	2.65
774 Q9H1B7	IRF2BPL	215	S	PKPTPEEGPPELNQ\$SPNSSAAVASRNG	2.65
775 Q01130	SRSF2	208	S	PPVSKRESKSRSRS\$PPKSPEEEGAVSS**	2.65
776 Q13428	TCOF1	341	S	VASQTAKGKPEEDSES\$SSEESSDSEEETPAA	2.64
777 Q8WUB8	PHF10	331	S	KGTSDSSGNVSEGES\$PPDSQEDSFQGRQKS	2.64
778 Q08J23	NSUN2	456	S	KLQGKSAETRESTQ\$SPADLTEGKPTDPSKL	2.64
779 Q13428	TCOF1	939	S	AQAGKQDDSGSSSEE\$SDS\$DGEAPAATSAQV	2.64
780 Q13523	PRPF4B	580	S	SEPSSPQSSSTRTRSP\$PDDILERVAADVKEY	2.64
781 Q14191	WRN	1133	S	ISSGSNISSKKSIMVQ\$PEKAYSSSQPVISAQ	2.63
782 Q14498	RBM39	341	S	VTERTDASSASSFLDS\$DELERTGIDLGTTGR	2.62
783 Q16629	SRSF7	209	S	SRSRSRSRSISRPRSS\$RSRSRSPSPKRSRSP	2.61
784 Q4VCS5	AMOT	1041	S	EVPASPATGPGPHRLS\$IPS\$LT\$CNPD\$KT\$DPV	2.61
785 Q5BKY9	FAM133B	192	S	LSKKRKMYSED\$KPLS\$SESLS\$ESE\$YIEEVRAK	2.61
786 Q6PD62	CTR9	1041	S	LKIADEGHPRNSNSNS\$SD\$DEDEQ\$KKCasse	2.61
787 Q6UN15	FIP1L1	85	S	ENGVPKPKVTETEDDS\$SD\$SD\$DD\$EDDVHVTI	2.61
788 Q7L2J0	MEPCE	152	S	QPHRPPGGGGKRRNS\$CNVGGGGGFKH\$PAF	2.61
789 Q7Z2Z1	TICRR	1750	S	LEDFELEGVCQLPDQ\$PPRN\$MPKAEEASSW	2.61
790 Q7Z6E9	RBBP6	1644	S	TRETDEAAFE\$PDYNE\$DSE\$NSV\$VKEE\$SSG	2.61
791 Q86VM9	ZC3H18	32	S	EEQPQGLS\$DD\$ILRD\$GSD\$Q\$LDGAG\$VRA\$D	2.60
792 Q96I25	RBM17	155	S	DRHEASGFARRPDPS\$DEDE\$YERERRKRSM	2.59
793 Q96ST2	IWS1	252	S	SENEELPKPRI\$DSE\$EDP\$PRHQASD\$ENE	2.58
794 Q96ST3	SIN3A	832	S	HHFIPDLLFAQRGDLS\$DVEEEEEE\$MDVDEA	2.58
795 Q99590	SCAF11	832	S	PSPRRETGKESRKS\$Q\$PSPKNESARGRK\$SR	2.58
796 Q9Y5S9	RBM8A	166	S	VRGPPKGKRGGR\$RS\$PSPDR\$RR\$*****	2.58

797	Q9BW71	HIRIP3	359	S	PTAKGSRKMARLGSTSGEESDLEREVSDSEA	2.57
798	O60832	DKC1	451	S	QVVAEAAKTAKRKRESESESDETPPAAPQLI	2.57
799	Q8N5F7	NKAP	23	S	RSPDREASGSGGRRRSSSKSPKSKSARSPR	2.57
800	Q01130	SRSF2	206	S	SPPPVSKRESKSRSRSKSPPKSPEEEGAVSS	2.57
801	P55198	MLLT6	258	S	KDKERLKQKHKKRPESPPSILTPPVPTADK	2.57
802	Q12923	PTPN13	1321	S	KPGISDVTDYSDRGDSDMDEATYSSSQDHQT	2.56
803	Q13428	TCOF1	547	S	ATPSAQVGKWEEDSESSEESSSDSSDGEVPT	2.56
804	Q13428	TCOF1	936	S	PSAAQAGKQDDSGSSSEESDSDEGAPAAVTS	2.56
805	Q9BW71	HIRIP3	363	S	GSRKMARLGSTS GEESSLEREVSDSEAGGGP	2.56
806	Q8IXM2	BAP18	35	S	TKLGELETMQLHPVADS SPAGAKWTETEIEMI	2.55
807	Q8TAP9	MPLKIP	133	S	TSTPFSGRVRERKRMSENELENYFKPSMLEDP	2.54
808	Q9NVU7	SDAD1	585	S	AAPGKSQKRKYIEIDS DEEPRGELLSLRDIE	2.54
809	Q8IXT5	RBM12B	278	S	HFRKRSHSKSPRTRRSRPLGFYVHLKNLSL	2.54
810	Q6PD62	CTR9	1102	S	SDQPSRKR RPSGSEQSDNESVQSGRSHSGVS	2.53
811	Q05519	SRSF11	434	S	KQVTRDYDEEEQGYDSEKEKKEEKPIETGS	2.53
812	Q8N9T8	KRI1	97	S	QKDATFYNR TASSSDSEEDPEALEKQKKVRP	2.53
813	Q13428	TCOF1	343	S	SQTKAGKPEEDSESSSEESSDSEEETPAAKA	2.52
814	P52948	NUP98	888	S	WVFKVSHFSKYGLQDSDEEEEHPSKTSTKK	2.51
815	Q96ST2	IWS1	377	S	SSDSEEEHHKKQKMDSD EDEKEGEEEKVAKR	2.51
816	Q9H019	MTFR1L	237	S	CSSSEEDDCVSLSKASS FADMMGILKDFHRM	2.50
817	Q9HCE7	SMURF1	106	S	GAGFLGCVRLLSN A I SRLKDTGYQRLDLCKL	2.50
818	P04637	TP53	33	S	TFSDLWKLLPENNVL SPLPSQAMDDMLMLSPD	2.50
819	P18583	SON	1935	S	ARSRTPSRRRSRSHTPS RRRRSRVGRRRSFS	2.50
820	P27815-7	PDE4A	6	S	*****MKRSRSALS VAGTG DERSRET	2.49
821	Q9UQ35	SRRM2	495	S	RMGRSRSPATAKGRGRSRSRTPTKRGHSRSRS	2.49
822	Q8WUB8	PHF10	327	S	KRKNKGTS DSSSGNVSEGESPPDSQEDSFQG	2.48
823	O00567	NOP56	519	S	SKPKKKKSFSKEELMS S DLEETAGSTSIPKR	2.48
824	Q9HCK8	CHD8	2069	S	VSRSPVVKLEDEDDSDSELDLSKLSPSSSS	2.47
825	Q9Y2X7	GIT1	388	S	SQSDLDDQHDYDSVAS DEDTDQEPLRSTGAT	2.47
826	Q13428	TCOF1	545	S	GPATPSAQVGKWEEDS ESSSEESSDSSDGEV	2.47
827	Q13428	TCOF1	932	S	GKTGPSAAQAGKQDDS GSSSEESDGEAPA	2.46
828	Q9NUQ3	TXLNG	105	S	TENRNLVSPAYCTQE SREEI PGGEAR TDPPD	2.46
829	Q14684	RRP1B	245	S	ETMEEQKTKVGDGLSAEEI PENEVSLRAV	2.45
830	O60832	DKC1	494	S	AKAGLES GAEPGDGDSDTTKKKKKKAKEV	2.45
831	Q01130	SRSF2	204	S	SRSPPVS KRESKSRSRSKSPKSPEEEGAV	2.45
832	Q01804	OTUD4	1024	S	ANSVDSRVQRPK EES SEDENEVSNILRSGRS	2.44
833	P35611-3	ADD1	358	S	LLNPEKYAKSRS PGSPVGE GTGSPPK WQIG	2.44
834	Q01804	OTUD4	1023	S	GANSVDSRVQRPK EES SEDENEVSNILRSGR	2.44
835	P50502	ST13	79	S	KVEEDLKADEPSSEES DLEIDKEGVIEPD TD	2.43
836	Q99759-2	MAP3K3	175	S	PRSRHLSVSSQNPGRSS PPPGYVPERQQHIA	2.42
837	Q16629	SRSF7	208	S	PSRSRSRSRSISRP RSSRSRSPSPKRSRS	2.42
838	Q7Z4V5	HDGFRP2	190	S	SDLDQASVSPSEEENS ESSSESEKTS DQDFT	2.42
839	Q13523	PRPF4B	578	S	VPSEPSS PQSSTRTRSPSPDDI LERVAADV K	2.41
840	Q9Y2W1	THRAP3	55	S	SRKRLLSSRSRSRSY S PAHN RERNH PRV YQN	2.41
841	Q9UQ35	SRRM2	564	S	QRRGRRSRSARRGRSH S RSPATRGRRSRTPA	2.41
842	Q9UQ35	SRRM2	562	S	NTQRRGRRSRSARRGRSH S HSRSPATRGRRSRT	2.40
843	O15541	RNF113A	43	S	AGRRKRPACDPEPGE SGSSSDEGCTVVRPEK	2.40
844	P16989	YBX3	369	S	AGEAPTE NPA PPTQQS AEE *****	2.39
845	Q9UQ35	SRRM2	456	S	PKPAPAPGSHREISS S P TS KRN RSH GRAK RD K	2.38
846	O75494	SRSF10	131	S	YRRSRSRSYERRRSRSR SF D NYR RSYSPRN	2.38
847	P16383	GCFC2	119	S	HHSSESKDDQGLLSSD SSS LGEKEL S STV KI	2.37
848	Q9NYV4	CDK12	332	S	GSYSGRSPSPYGR RRS SSSPFLSKRSLSRSPL	2.37
849	Q9Y2W2	WBP11	237	S	ALDLPPRRD EMLY S PELA QRGH D DD VS ST	2.37
850	Q9NYV4	CDK12	333	S	SYSGRS PSPYGR RRS S PFLSKRSLSRSPL	2.36
851	Q8IZ21	PHACTR4	270	S	AKQPPI PPKPAH RNS NPVIAEL SQAINSGT	2.35
852	Q9UPU7	TBC1D2B	957	S	F LRERDTSPDKGELV S DEEE DT *****	2.35
853	Q14247	CTTN	438	S	DAASFKAEL SYRG PV S GTEPEPV YSMEA ADY	2.34

854 Q8WX93	PALLD	1121	S	SGHPHVRRPRSRSDS G DENEPIQERFFRPH	2.34
855 P46100	ATRX	1348	S	KPRYRHRLRHKLTV S DGESGEEKKTPKEH	2.27
856 Q13428	TCOF1	549	S	PSAQVGKWEDESESS S EESSDSSDGEVPTAV	2.24
857 P52701	MSH6	287	S	EGSSDEI S SGVGVDSE S EGLN S PVKVARKRKR	2.24
858 Q8NDI1	EHBP1	174	S	LREGKATDEDMQSLAS S LMSMKQADIGNLDDF	2.23
859 Q04637	EIF4G1	1596	S	SVTAFFKWLREAEES S DHN*****	2.19
860 Q6PD62	CTR9	1043	S	IADEGHPRNSNSNSD S DEDEQRKKCASSESD	2.16
861 Q96CB8	INTS12	378	S	PPPLTLGKTGLRSRSV S CNDNVSKVGLPSPSSL	2.16
862 Q96ST2	IWS1	25	S	QSDDGGATPVQDERD S GSDGEDDVNEQHSGS	2.16
863 P16383	GCFC2	117	S	KIHHSSESKDDQGLSS S DSSSLGEKELSSTV	2.15
864 P51858	HDGF	133	S	AAEGDGDKKGNAEGSS S DEEGKLVIDEPAKEK	2.15
865 O60832	DKC1	485	S	KKKSKKDKKAKAGLE S GAEPGDGSDTTKKK	2.14
866 P18583	SON	1929	S	NRKTVRARSRTPSRRS R SHTPSRRRSRSGV	2.14
867 Q8TF01	PNISR	286	S	TEDAEGGDGPRLPQRS K FDSDEEEEDTENVE	2.14
868 Q9Y3T9	NOC2L	673	S	DEDRKQFKDLFDLNS S EEDDT E FGSERGILR	2.13
869 O95382	MAP3K6	914	S	TLLGDPFLQPGKRSRS P SSPRHAPR PSDAPS	2.12
870 O00567	NOP56	570	S	GSKKKRKFSKEEPVSS S GPEEAVGKSSSSKKK	2.11
871 Q8TF01	PNISR	396	S	SLTGLGGLGGYGS G D S EDERSDRGS S ESSD T D	2.04
872 P18583	SON	1783	S	ASPVVSSMPERASES S SEEKD D YEIFVKVKD	2.04
873 Q5T3I0	GPATCH4	130	S	LTSGGEKPNKDL E SC S DDD N QGS K SPKILTD	2.04
874 Q96IZ7	RSRC1	116	S	SKSRTRRSRSRPLRS H SRSSSERSSHRRTRS	2.04
875 Q9NPQ8	RIC8A	436	S	AARGLMAGGRPEGQY S EDED T DTDEYKEAKA	2.00
876 Q9Y2W1	THRAP3	743	S	KERDLKRGKSRESV D RDSSHSRERSAEKTE	2.00
877 Q01433	AMPD2	188	T	LEPDILLR A QDFLKT T DSDSDLQLYKEQGEG	1769.91
878 O75400	PRPF40A	932	T	KSPKKKTGKDSGNWD T SGSEL S E G ELEKRRR	1140.61
879 Q9UQ35	SRRM2	1927	T	RSRASPVSRRRSRSRT T PPVTRRRSRSRTPTT	786.63
880 Q3YEC7	RABL6	599	T	RRADD F PVRDDPSDV T DEDEGP A E PPP PKL	570.30
881 P25205	MCM3	674	T	EKEKKRK R SEDESET T DEEE E KSQEDQE Q KR	550.64
882 Q6QNY0	BLOC1S3	63	T	RGRPTGLRVAGEAAET T DSEPEPEPEPTA APR	452.31
883 Q6PD62	CTR9	925	T	RSKKGG E DEFVN D TT D DLPI S KKKRRKG	393.31
884 Q4G0J3	LARP7	338	T	EASEASKENRDIEIS T EE E KDTGDLKDSSL	334.32
885 Q9NU22	MDN1	4898	T	DDLNLDSED K NGGED T DNEEGEEENPLEIKE	324.37
886 P55081	MFAP1	267	T	LEENKRS L AALDALNT T DDENDEEEYEAWKVR	314.65
887 Q13427	PPIG	358	T	TPSRSRSRDRFRRSET T PPHWRQEMQRAQRMR	255.65
888 P25205	MCM3	713	T	DAKGDGSYDPYDFSD T EEEMPQVHTPKTADS	235.99
889 Q8IZL8	PELP1	1090	T	KVQPPPETPAEEEMET T ETEA E ALQEKEQDDT	235.99
890 Q5C9Z4	NOM1	287	T	KAQEAEAQ S EDDD D ETEEEQGEEKEKG A QEK	196.66
891 Q8N5F7	NKAP	161	T	WGLSPKN P EPDSDEHT T PVEDEEPK K STSAS	186.90
892 Q8TF01	PNISR	410	T	DSEDE S DRGS S ESSD T DDEELRHRIRQKQEA	176.99
893 Q9NYF8	BCLAF1	402	T	KESGKQKFNDSEGDD T EETEDYRQFRKSVLA	174.01
894 Q8IWX8	CHERP	819	T	SYSPGRRRSRSRS R ST T PPSSAGLGSNS APP I	157.33
895 O43583	DENR	86	T	ENSPKQ E AGISEGQGT T AGEEEEKKQKRGGR	137.66
896 Q8WVC0	LEO1	188	T	EDKLQNS D DEKM Q NT T DDE P QL S DDER Q	117.99
897 P51116	FXR2	411	T	GSGRGSGGS D KAGYST T DESSSSSLHATRTYG	108.33
898 Q8N1G4	LRRC47	522	T	KKYTLENKEEGSLS T EADAVSGQLPDPTT N	105.23
899 Q9H501	ESF1	311	T	SDSGPDI A R G KGNIE T SSEDED D TADLF PEE	100.32
900 Q9UQ35	SRRM2	717	T	SRSRSLVRRGRSHS R TPQRGRGSSSERKN	98.33
901 P62269	RPS18	60	T	LRKADIDLT K RAGEL T EDEVERVITIMQNPR	88.66
902 Q13427	PPIG	748	T	VHEKNKKFDHES S PG T DEDKSG*****	88.06
903 Q13610	PWP1	55	T	KEKLQEE GGG SDEEE T GPS E DGMQSARTQA	84.66
904 Q14684	RRP1B	454	T	LKARVAEPGAEATSS T GEESGSEHPPAVPMH	83.54
905 Q5VZL5	ZMYM4	118	T	SSIHTDD S LEVERR T QHESDNENEIQIQNK	81.60
906 Q8WVC0	LEO1	302	T	KRKNAIASDSEADSD T EVPKDN S GTMDLF GG	80.61
907 Q9UQ35	SRRM2	1655	T	SSSKGRGPS E GS T ESSPEHPPKSRTARR	78.66
908 Q9UQ35	SRRM2	2022	T	RSRTSPVTRRRSRS R TPPAIRRRSRSRTPLL	72.11
909 Q9UQ35	SRRM2	2034	T	RSRTPPAIRRRSRS R TPLLPRKRSRSR S PLA	63.41
910 Q92733	PRCC	261	T	KAAAKSAALQVT K Q I T Q EEDDS D EVAPENF	59.00

911 Q9NPQ8	RIC8A	441	T	MAGGRPEGQYSEDED T DTDEYKEAKASINPV	57.64
912 Q9UQ35	SRRM2	2069	T	SRSRTPRTARGKRSLT TRSPPAIRRSASGSS	54.12
913 Q9H501	ESF1	319	T	RGKGNIETSSSEDED T ADLFPEESGFHAWR	49.85
914 P51116	FXR2	451	T	RRTGGPAYGPSSDV T ASETESEKREEPNRA	49.33
915 Q4G0J3	LARP7	257	T	DTSNTSISKMKRSRP T SEGSDESTEPQKQC	47.33
916 Q7L590	MCM10	85	T	EKENLATLFGDMED T DEEEVPASQSTENRV	47.54
917 Q8IZL8	PELP1	1014	T	PEPEPGLLVEE P GTEEERGADTAPE	46.32
918 Q92541	RTF1	334	T	EEDDKSSEKSDRSSRT T SSSDEEEKEIPPK	46.22
919 Q96QU1	PCDH15	1863	T	CTTNLMPAEKIKSSMT T QLSTTVCKTDPQRE	42.38
920 O15021	MAST4	1821	T	GTLDIALLSGPQASK T ELPSPESAOQSPSPSG	42.79
921 Q9H0G5	NSRP1	263	T	SDFDAKSSADDEIEET TRVNCRREKVIETPEN	41.33
922 Q9UJV9	DDX41	13	T	***MEESEPERKRART T DEVAGGSRSEAED	39.33
923 Q96ST2	IWS1	317	T	SESEGPKGPASDSET TEDASRHKQKPESDDD	37.33
924 Q8WVC0	LEO1	629	T	SEEDKAQRLLKAKKL T SDEEGEPSGKRKAED	34.52
925 Q96QU1	PCDH15	1867	T	LMPAEKIKSSMTQLS TTVCKTDPQREP KGI	28.51
926 Q96QU1	PCDH15	1873	T	IKSSMTQLSTTVCKT TDPQREP KGILRHVKN	27.34
927 Q66PJ3	ARL6IP4	343	T	QWHRSGEEDGPVL T DEQKSRIQAMKPMTK	7.38
928 Q8NE71	ABCF1	108	T	EEKELMERLKKL SVPT TSDDEDEVAPKPRGG	6.90
929 P27361	MAPK3	202	T	LARIADPEHDHTGFL TEYVATRWYRAPEIML	5.02
930 P27361	MAPK3	207	T	DPEHDHTGFLTEYVA TRWYRAPEIMLNSKGY	4.88
931 O75528	TADA3	297	T	IPDMSGKESGADGAS T SPRNQNKPFSVPHTK	4.46
932 Q8IYB3	SRRM1	872	T	LAQEEPVAAPEPK TE SEEAEDNLDDLEKHL	4.42
933 Q13523	PRPF4B	847	T	LCDFGSASHVADND IT PYLVSRFYRAPEIII	3.60
934 Q9UQ35	SRRM2	577	T	SHSRSPATRGRSRSR T PARRGRSRSRTPARR	3.25
935 Q9UQ35	SRRM2	627	T	RGRSRSPATRGRSRSR T RSPVRRRSRSRSPAR	3.19
936 P17544	ATF7	53	T	GPARTDSVIIADQT TP TRFLKNCEEVGLFN	3.15
937 Q27J81	INF2	1179	T	PAAGPGGDEDEDE DT APEALDTSLDKSFS	3.15
938 O15042	U2SURP	931	T	DECTPTRKERKRRH ST SPSPSRSSGRRVKS	3.05
939 P17544	ATF7	51	T	KFGPARTDSVIIADQT TP PTPTRFLKNCEEVGL	3.05
940 Q9BVV6	KIAA0586	378	T	NVRLLEQILNNNDSL TRKSESSNTTSLTRSK	2.89
941 P85037	FOXK1	247	T	PDLRSMVSPVPSPT GT ISVPNSCPASPRGAG	2.87
942 Q9BW71	HIRIP3	358	T	EPTAKGSRKMARLG ST SGEESDLEREVSDSE	2.81
943 P28482	MAPK1	190	T	DPDHDTGFLTEYVA TRWYRAPEIMLNSKGY	2.75
944 Q9BW71	HIRIP3	84	T	REDKLDLTKKGKRPP T PCSDPERKRFRFNSE	2.67
945 Q13523	PRPF4B	576	T	MSVPSEPPSQSSTR T SPSPDDILERVAAD	2.64
946 Q9UQ35	SRRM2	499	T	SRSPATAKRGRSRSR T PTKRGHSRSRSPQWR	2.64
947 P42166	TMPO	74	T	SKGPPDFSSDEERE PT PVLGSGAAAAGRSRA	2.54
948 Q6UN15	FIP1L1	79	T	IEDETAENGVPKPKV T ETEDDSDDSDSDDDED	2.50
949 P51003	PAPOLA	544	T	SDNSMSVPSPTS ATKT TSPLNSSGSSQGRNSP	2.45
950 O95218-2	ZRANB2	303	T	SRSRSSSGDRKKR TR RSRSPESQVIGENTK	2.41
951 P35222	CTNNB1	551	T	LVQLLVRAHQDTQRR T SMGGTQQQFVEGVRM	2.37
952 P06493	CDK1	14	T	**MEDYTKIEKIGEGT TYGVVYKGRHKTTGQV	2.27
953 Q13017	ARHGAP5	1217	T	SPVETWKGGIDNP AI TSDQEELDDKKMKKTH	2.18
954 Q02156	PRKCE	710	T	VNNFDQDFTR T REEPVLT	2.14
955 P29692	EEF1D	147	T	VSPMRQVEPPAKK PA T	2.06
956 Q04759	PRKCQ	536	T	ADFGMCKENMLGDAK T NTFCGTPDYIAPEIL	2.04
957 Q7KZI7	MARK2	208	T	DFGFSNEFTFGNKLD T FCGSPPYAAPELFQG	2.00
958 Q13627	DYRK1A	321	Y	LGQRIY QY IQSRFYR	373.65
959 P25205	MCM3	705	Y	DAKGD SY DPYDFSD	117.99
960 P45973	CBX5	20	Y	SSEDEEE Y VVEKVLD	61.34
961 O60841	EIF5B	134	Y	AKPKVEM Y SGSDDDD	59.54
962 O15541	RNF113A	80	Y	SGKQKAA Y GDLSSEE	59.14
963	PREDICTED:	113	Y	MVSFIAV Y TSHLS--	59.00
964 P98175	IKFZp686E245	122	Y	SQEGKHD Y DDSSEQ	57.64
965 O15173	PGRMC2	179	Y	EGEEPTV Y SDEEPEK	49.31
966 P51116	FXR2	409	Y	GGSDKAG Y STDESSS	39.33
967 P06493	CDK1	15	Y	EKIGEGT Y GVVYKGR	8.63

968	B4DWW4	MCM3	753	Y	DGDSYDPYDFSDTEE	7.42
969	Q9HCE7	SMURF1	113	Y	SRLKDTGYQRQLDLCK	7.19
970	P27361	MAPK3	204	Y	HTGFLTEYVATRWYR	4.74
971	Q13523	PRPF4B	849	Y	ADNDITPYLVSRFYR	4.61
972	P49840	GSK3A	279	Y	RGEPNVSYICSRYYR	2.84
973	G3V5T9	CDK2	15	Y	EKIGEGTYGVVYKAR	2.27
974	Q8IZT6	ASPM	2564	Y	SIVIQSTYRMYRQYC	2.11

Table S1: Identified phosphorylation sites are list with accession information, phosphorylation site with positions in the proteins, spanning amino acids sequences and fold of changes. Protein identifications overlapped with protein array analysis (Keller *et al.*, *Mol Cell* 2014) were highlighted in blue.

Supplemental Table S2

No.	Sample ID	Gender	Age	Histology (WHO Grade)
1	2014-06517	M	51	Renal Clear Cell Carcinoma (II)
2	2014-03303	M	38	Renal Clear Cell Carcinoma (II)
3	2014-01577	F	64	Renal Clear Cell Carcinoma (II -III)
4	2014-05726	F	78	Renal Clear Cell Carcinoma (I - II)
5	2014-02433	M	69	Renal Clear Cell Carcinoma (II)
6	2013-05583	F	40	Renal Clear Cell Carcinoma (II)
7	2013-05739	F	62	Renal Clear Cell Carcinoma (II -III)
8	2013-06867	M	32	Renal Clear Cell Carcinoma (I - II)
9	2014-00532	M	77	Renal Clear Cell Carcinoma (II -III)
10	2014-00862	M	58	Renal Clear Cell Carcinoma (II)
1	2015-05563	M	38	Breast Infiltrating Ductal Carcinoma (II)
2	2015-06795	M	51	Breast Infiltrating Ductal Carcinoma (II -III)
3	2015-19494	M	45	Breast Infiltrating Ductal Carcinoma (III)
4	2015-19651	M	39	Breast Infiltrating Ductal Carcinoma (II)
5	2015-19745	M	53	Breast Infiltrating Ductal Carcinoma (III)
6	2015-19926	M	49	Breast Infiltrating Apocrine Carcinoma (II -III)
7	2015-20046	M	69	Breast Infiltrating Ductal Carcinoma (II)
8	2015-20132	M	47	Breast Infiltrating Ductal Carcinoma (II)
9	2015-20133	M	62	Breast Infiltrating Ductal Carcinoma (II)
10	2015-20135	M	36	Breast Infiltrating Ductal Carcinoma (II)

Supplemental Table S2, Information of RCC and breast cancer samples were obtained and listed (related to Figure 7A&B and Figure S6).

Supplementary Experimental Procedures

Antibodies

Ser202, S203 and S202/203 specific phosphorylation antibodies were homemade. AKT1S1 antibody (#2610), PKM2 antibody (#3198), p70 S6 kinase antibody (#9202), Phospho-p70 S6 kinase (Thr389) antibody (#9205), LC3B antibody (#2775), 4E-BP1 (53H11) rabbit mAb (#9644) and Phospho-4E-BP1 (Thr37/46) (236B4) Rabbit mAb (#2855) were purchased from Cell Signaling Technology. Anti-SQSTM1 / p62 antibody (ab56416) was from abcam.

***In vitro* PKM2 phosphorylation of HEK293T proteome**

HEK293T proteome were prepared as following: HEK293T cells were cultured in DMEM medium with 10% newborn bovine serum, 100 units/ml penicillin and 100 µg/ml streptomycin in a humidified atmosphere of 5% CO₂ at 37°C. Harvested cells were collected, washed with PBS (pH7.5) and lysed in 50 mM HEPES buffer (pH 7.5) buffer containing 1% Triton (v/v) , 65 mM DTT , 1 mM PMSF, 2% proteases inhibitors cocktail (Roche) for 30 minutes. After centrifugation at 12000 rpm at 4°C for 30 minutes, the supernatant was collected and subject to either acetone precipitation followed by PKM2 phosphorylation or immobilization to CNBr-activated Sepharose for dephosphorylation and phosphorylation. For proteome dephosphorylation, 50U alkaline phosphatase (Sigma) was incubated with immobilized proteins in 1mM MgCl₂, 1mM ZnCl₂, 50 mM HEPES (pH 7.5). The resulted dephosphorylated immobilized proteome were washed with PBS (pH7.5) overnight to remove alkaline phosphatase and any small molecules that may serve as phosphor-donor.

***In vitro* kinase reaction**

The proteins or dephosphorylated immobilized proteome were preincubated in kinase buffer (100mM KCl, 50mM MgCl₂, 1mM DTT, 1mM NaVO₄, 5% glycerol, 30mM HEPES (pH 7.6)) at 30°C for 30min. *In vitro* kinase reaction was triggered by adding 50 µg recombinant PKM2 and 0.5mM PEP, while in the control experiment only

contains 0.5 mM PEP. Both reactions were allowed to process at 30°C for 2 hours before 100 mM ammonium bicarbonate (NH_4HCO_3) was added to stop the reaction.

Trypsin digestion and phosphopeptides enrichment

The washed immobilized proteome were incubated with 20 mM DTT for 2 hours at 37°C, followed by addition of 60 mM iodoacetic acid to the reaction mixture. After incubation in the dark for 1 hour at 25°C, trypsin was added at 50 µg/ mg protein in 100mM NH_4HCO_3 buffer (pH 8.0). After digestion for 18h at 37°C, the released peptides were collected and acidized by 5% trifluoroacetic acid before they were subjected to phosphopeptides enrichment by Ti41-IMAC microspheres.

Mass Spectrometry

The mass spectrometer was set as below: ion transfer capillary 200 °C, spray voltage 1.8 kV, full MS range 400-2000, and full mass spectra was acquired in the Orbitrap at a resolution of 60,000 with the target ion setting of 10^6 . One full MS scan was followed by ten MS/MS scans, and the multistage activation was enabled. The dynamic exclusion function was set as follows: repeat count 2, repeat duration 30 s, and exclusion duration 60 s. Each sample was analyzed five times by the RPLC-MS/MS.

Gene silence

Synthetic oligos were used for *siRNA* silencing of *AKT1S1* and *PKM2*, the sequence used for silencing genes are:

AKT1S1: -GCTGAGTTCTAAGCTCTAAAT-

PKM2: -CTGCAGCGTTGTTAGCAAATAAT-

For *shRNA* silencing of genes, above sequences were cloned into PMKO vector, the plasmids constructed were cotransfected into HEK293T cells together with the *gag* and *vsvg* genes expressing plasmids to produce retrovirus. Retroviral supernatant was harvested 36 hr after initial plasmid transfection and mixed with polybrene (8 mg/ml) to increase the infection efficiency. Cells were infected with retrovirus and selected in puromycin (1 µg/ml) for 2 weeks. Knockdown efficiencies were analyzed by either western blots or Q-PCR.

Bioinformatics analysis

All analysis were performed using published software. The secondary structures and solvent accessibilities of HPSRPs were predicted with NetSurfP 1.1 (Petersen et al., 2009), while the disorder regions and sub-cellular localizations were predicted by ESpritz (Walsh et al., 2012) and WoLF PSORT (Qi et al., 2007), repectively. The gene ontology annotations were downloaded from QuickGO database (Barrell et al., 2009), while the enrichment analyses were performed with a hypergeometric distribution as previously described (Qi et al., 2014). The enrichment analyses of KEGG pathways were carried out using DAVID (Huang da et al., 2009), which also presented the pathways with annotated HPSPs. All the heatmaps were visualized with the ggplot2 program (<http://had.co.nz/ggplot2/>) in the R package (<http://www.r-project.org/>). The amino acid preferences were visualized with WebLogo (Crooks et al., 2004), while the comparisons of amino acid preferences were visualized with Two Sample Logo (Vacic et al., 2006).

Immunohistochemistry

Tissue sections were prepared form the formalin-fixed paraffin embedded specimens. Antigen retrieval of renal cell carcinoma or breast cancer specimens was performed by incubating the slides in Tris-EDTA buffer (pH 8.4) at 99°C for 60 mins. The endogenous peroxidase activity was inactivated in solution of methanol with 3% H₂O₂. The slides were incubated with primary antibody for 60 mins and secondary antibody for 8 mins, then stained with DAB Chromagen for 8 mins. All procedures were performed using stainer (BenchMark XT, Ventana) and the slides were scaned by scanner (Ventana iScan Coreo). The quantification of IHC results were performed by an experienced pathologist. The intensity was calculated according to positive areas and positive degree. Sections were staining with PKM2 (1:100), P-S202/203-AKT1S1 (1:30), P-S2448-mTOR (1:100), P-T37/46-4EBP1 (1:500), p62 (1:200) and LC3B (1:100) antibody using a Ultraview Detection Kit.

Supplementary References

- Qi, H., P., Park, K.J., Obayashi, T., Fujita, N., Harada, H., Adams-Collier, C.J., and Nakai, K. (2007). WoLF PSORT: protein localization predictor. *Nucleic acids research* 35, W585-587.
- Barrell, D., Dimmer, E., Huntley, R.P., Binns, D., O'Donovan, C., and Apweiler, R. (2009). The GOA database in 2009—an integrated Gene Ontology Annotation resource. *Nucleic acids research* 37, D396-403.
- Crooks, G.E., Hon, G., Chandonia, J.M., and Brenner, S.E. (2004). WebLogo: a sequence logo generator. *Genome research* 14, 1188-1190.
- Huang da, W., Sherman, B.T., and Lempicki, R.A. (2009). Systematic and integrative analysis of large gene lists using DAVID bioinformatics resources. *Nature protocols* 4, 44-57.
- Petersen, B., Petersen, T.N., Andersen, P., Nielsen, M., and Lundegaard, C. (2009). A generic method for assignment of reliability scores applied to solvent accessibility predictions. *BMC structural biology* 9, 51.
- Qi, L., Liu, Z., Wang, J., Cui, Y., Guo, Y., Zhou, T., Zhou, Z., Guo, X., Xue, Y., and Sha, J. (2014). Systematic Analysis of the Phosphoproteome and Kinase-substrate Networks in the Mouse Testis. *Molecular & cellular proteomics : MCP* 13, 3626-3638.
- Vacic, V., Iakoucheva, L.M., and Radivojac, P. (2006). Two Sample Logo: a graphical representation of the differences between two sets of sequence alignments. *Bioinformatics* 22, 1536-1537.
- Walsh, I., Martin, A.J., Di Domenico, T., and Tosatto, S.C. (2012). ESpritz: accurate and fast prediction of protein disorder. *Bioinformatics* 28, 503-509.