

Name	Uniprot AC	kDa	CE pre-B dimer			CE B-like dimer		
			PSMs	iBAQ	iBAQ RANK	PSMs	iBAQ	iBAQ RANK
PRP8	Q6P2Q9	273.6	2,283	2.15E+09	22	1,524	1.64E+09	35
BRR2	O75643	244.5	2,449	2.41E+09	18	1,676	1.81E+09	29
SNU114	Q15029	109.4	1,157	2.99E+09	10	709	2.71E+09	14
U5-40K	Q96DI7	39.3	329	1.20E+09	53	281	1.44E+09	41
PRP6	O94906	106.9	657	1.77E+09	32	505	1.49E+09	39
DIM1	P83876	16.8	202	8.98E+08	61	163	1.19E+09	47
PRP28	Q9BUQ8	95.6	715	2.09E+09	24	428	1.19E+09	46
LSm proteins								
LSm2	Q9Y333	10.8	84	1.15E+09	55	80	1.94E+09	24
LSm3	P62310	11.8	57	2.32E+09	19	87	3.42E+09	10
LSm4	Q9Y4Z0	15.4	119	6.66E+08	75	82	7.21E+08	63
LSm5	Q9Y4Y9	9.9	46	1.28E+09	48	40	1.69E+09	32
LSm6	P62312	9.1	42	2.01E+09	26	26	1.30E+09	44
LSm7	Q9UK45	11.6	28	2.67E+08	105	69	5.58E+08	72
LSm8	O95777	10.4	59	1.77E+09	31	126	2.50E+09	17
U4/U6								
PRP3	O43395	77.6	568	1.25E+09	51	426	1.06E+09	52
PRP4	O43172	58.5	486	1.69E+09	34	358	1.77E+09	30
PPIH	O43447	19.2	143	9.26E+08	60	101	7.51E+08	61
PRP31	Q8WWY3	55.5	397	1.86E+09	29	258	1.65E+09	34
SNU13	P55769	14.2	80	1.56E+09	37	59	1.85E+09	25
U4/U6.U5								
SNU66	O43290	90.3	700	1.94E+09	27	489	1.96E+09	23
SAD1	Q53GS9	65.4	339	2.03E+09	25	301	1.82E+09	28
SNRNP27	Q8WVK2	18.9	96	1.66E+09	35	48	1.16E+09	49
RBM42	Q9BTD8	50.4	209	1.53E+09	40	123	9.08E+08	58
PRP4 kinase	Q13523	117	587	9.52E+08	59	341	6.47E+08	67
B-specific proteins								
SNU23 (ZMAT2)	Q96NC0	23.6	27	3.20E+07	238	52	2.21E+08	96
PRP38A	Q8NAV1	37.5	116	4.98E+08	91	99	5.02E+08	74
SMU1 (fSAP57)	Q2TAY7	57.5	311	8.22E+08	65	244	9.77E+08	55
MFAP1	P55081	52	89	1.90E+08	122	76	2.76E+08	86
RED	Q13123	65.6	214	4.85E+08	92	167	4.55E+08	78
FBP21	O75554	42.5	36	3.73E+07	229	83	2.27E+08	95
UBL5	Q9BZL1	8.5	17	9.06E+07	175	24	4.83E+08	76
Cap binding complex								
NCBP2	P52298	18	47	5.49E+08	87	57	6.10E+08	70
NCBP1	Q09161	91.8	479	8.28E+08	64	508	9.12E+08	57
SR proteins								
SRSF1	Q07955	27.8	230	5.85E+09	4	200	4.42E+09	6
SRSF7	Q16629	27.4	114	4.02E+09	5	80	2.91E+09	13
SRSF3	P84103	19.3	53	1.55E+09	38	42	6.44E+08	68
SRSF5	Q13243	31.3	50	2.13E+08	115	41	2.44E+08	91
SRSF6	Q13247	39.6	70	7.62E+08	71	46	4.32E+08	79
SRSF2	Q01130	25.5	170	2.53E+09	15	116	2.26E+09	20
SFRS12	Q8WXA9	59.4	40	1.36E+08	144	53	2.52E+08	90

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			PSMs	iBAQ	iBAQ RANK	PSMs	iBAQ	iBAQ RANK
SRS11	Q05519	53.5	117	6.48E+08	77	103	6.86E+08	65
TRA2B	P62995	33.7	73	6.08E+08	83	24	1.62E+08	109
SR-related proteins								
FLJ10154	Q9NWB6	33.2	37	8.05E+08	67	26	2.35E+08	94
hnRNP								
HNRNP A0	Q13151	30.9	119	1.19E+09	54	129	1.15E+09	50
HNRNP A1	P09651	38.7	432	2.46E+09	17	310	2.91E+09	12
HNRNP A3	P51991	39.6	229	6.58E+08	76	75	1.54E+08	110
HNRNP A2/B1	P22626	37.4	404	2.92E+09	11	192	9.49E+08	56
HNRNP C	P07910	33.7	180	1.32E+09	47	21	4.48E+07	151
HNRNP H1	P31943	49.2	178	5.23E+08	88	76	1.83E+08	101
HNRNP L	P14866	64.1	192	4.26E+08	94	37	2.26E+07	196
PCBP1	Q15365	37.5	162	6.10E+08	82	105	8.09E+08	59
Other factors								
MSI2	Q96DH6	35.2	90	6.30E+08	80	62	4.65E+08	77
PRP38B	Q5VTL8	64.5	118	2.12E+08	116	109	2.41E+08	93
ERH	P84090	12.3	93	3.84E+09	6	37	7.37E+08	62
MGN2	Q96A72	17.3	91	3.66E+08	98	3	3.98E+06	310
RSRC2	Q7L4I2	50.6	117	7.68E+08	70	89	6.76E+08	66
TOE1	Q96GM8	56.5	72	9.22E+07	172	139	4.22E+08	80
EIF4A3	P38919	46.9	254	1.11E+09	56	23	7.50E+06	264
MAGOH	Q96A72	17.2	91	3.66E+08	98	3	3.98E+06	310
RBM8A (Y14)	Q9Y5S9	19.9	128	7.48E+08	72	5	3.21E+06	327
PININ	Q9H307	81.6	204	3.51E+08	99	59	3.68E+07	165
RNPS1	Q15287	34.2	152	5.73E+08	85	32	1.41E+08	111
SAP18	O00422	17.6	94	1.01E+09	58	36	1.72E+08	104
THOC4 (ALY/REF)	Q86V81	26.9	137	5.67E+08	86	18	4.53E+07	150
UAP56	Q13838	49	224	7.87E+08	68	42	3.18E+07	172
PABPN1, PAB2	Q86U42	32.8	75	5.20E+08	89	2	3.92E+06	311
SRRM2	Q9UQ35	300	600	4.19E+08	95	289	2.56E+08	89
BUB3	O43684	37.2	177	1.47E+09	42	192	1.84E+09	26
ZNF207	O43670	50.8	99	5.08E+08	90	72	4.93E+08	75
SF4 (F23858)	Q8IWZ8	72.5	235	2.60E+08	107	182	2.44E+08	92
PRP19	Q9UMS4	55.2	302	1.04E+09	57	130	2.68E+08	88
SPF27	O75934	26	94	3.78E+08	96	42	1.27E+08	117
FUS	P35637	53.4	108	4.66E+08	93	95	5.74E+08	71
PPP1R8 (NIPP1)	Q12972	38.5	85	9.05E+07	176	84	2.14E+08	97
ZC3H18 (NHN1)	Q86VM9	106.4	294	3.66E+08	97	219	2.69E+08	87
SRRT (ASR2B)	Q9BXP5	100	598	1.54E+09	39	487	1.45E+09	40
DDX5 (p68)	P17844	69.2	309	6.94E+08	73	127	2.13E+08	98
AGGF1	Q8N302	80.977	92	9.19E+07	173	155	1.97E+08	99
BUD31	P41223	17.0	50	2.00E+8	118	34	1.84E+08	100
MBP-MS2*		56.9	1,338	1.32E+10	1	1,287	1.16E+10	1