

Homo sapiens	Arabidopsis thaliana	AGI locus	UniProt ID	Peptides	Sequence coverage [%]	Score	Intensity	MS/MS Count
<b>US snRNP</b>								
Brr2	BRR2/MAC14/EMB1507	AT1G20960	Q48534	29	14.6	79,507	444470000	43
US-40	MAC17	AT2G43770	O22B26	5	17.8	8,6494	59862000	5
Prp8	PRP8B	AT4G38780	Q95SD2	23	9.4	58,79	292690000	31
Snu114	GFA1/MEE5/CLO	AT1G06220	Q9LNC5	14	17.7	64,336	266600000	19
<b>US sm ring</b>								
Sm B/B'		A75G44500	Q9FH15	2	5.9	3,1189	14232000	2
Sm D2		A73G62840	Q8RUH0					
		A72G47640	Q8RUH0					
Sm D1	SmD1a	A73G07590	Q959F1					
	SmD1b	A74G02840	Q95Y09					
Sm D3		AT1G20580	Q9LMS2					
Sm E		A72G18740	Q9ZV45	1	14.1	10,073	10818000	1
Sm F	RUXF	A74G30220	F4JPK5					
Sm G		A72G23930	O82221	1	22.4	3,2787	5851700	1
		A73G11500	Q9CAX7					
<b>NTC</b>								
Cdc5	CDC5/MAC1	AT1G09770	P92948	18	21.6	39,897	228190000	19
Prp19	MAC3A	AT1G04510	F4EP3	12	27	32,736	130360000	10
	MAC3B	A72G33340	O22785	12	28.8	57,663	448850000	14
Sp127	MOS4	A73G18165	Q94959	1	4.3	3,2363	17435000	1
Syl1	MAC9	A75G28740	Q9LKL3	14	15.9	33,249	171140000	14
Syl2		A72G16860	Q9ZVX7					
Syl3	MAC10	A75G41770	F4JZK8	13	16.3	16,361	138040000	11
lay1	MAC1	A73G18790	Q9L597	1	3.7	1,3055	3306300	1
CTNBNL1		A73G02710	A3KPF3					
HSP73	HSP70	A73G09440	O65719	21	32.7	33,405	111960000	13
<b>NTR</b>								
RBM22	MAC5A	AT1G07360	Q9LNV5	6	12.5	26,372	42295000	6
SKIP	SKIP	AT1G77180	O80653	8	15.3	30,515	91681000	8
G10		A74G21110	O49553	1	11	14,075	4845200	1
PPL1		A72G36130	Q95H1					
Ad-002	CWC15	A73G13200	Q9LKS2	26	61.3	304.66	18999000000	56
PRL1	PRL1	A74G15900	Q42384	6	15.8	18,683	43612000	7
Aquarius	EMB2765	A72G38770	Q8L5Y4	12	8.9	28,811	122810000	12
<b>Splicing Factor</b>								
SRRM2/SRm300		A73G49601	B3H6J5					
Cwc22		AT1G80930	Q9SAG7	1	1.2	1,4812	1191100	1
RNF113A		A75G06420	Q9FNG6					
		A71G01350	Q8GK84					
Prp2	EMB2733/ESP3	AT1G32490	Q8VY00					
Prp17		AT1G10580	A4FVN8	4	7.9	9,6605	34635000	4