

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
398364847	RTR1	NP_011066.3	0	0	0	0	0	0	0	0	0	0	0	0	8.85
6320061	RPB1	NP_010141.1	7.67	12	17	0	0	0	3.23	5	16	1.04	1	2	3.98
398365417	RPB2	NP_014794.3	6.05	7	8	1.14	1	2	2.45	2	5	2.70	3	7	3.02
398364469	RPB3	NP_012243.3	0	0	0	0	0	0	0	0	0	0	0	0	5.35
6322321	RPB4	NP_012395.1	8.60	2	2	0	0	0	0	0	0	0	0	0	0
6319630	RPB5	NP_009712.1	6.98	1	1	0	0	0	0	0	0	0	0	0	13.02
398366567	RPB7	NP_010692.3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365775	RPB10	NP_014853.3	0	0	0	0	0	0	0	0	0	37.14	1	1	0
6325445	RPB6	NP_015513.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321368	RPB9	NP_011445.1	8.20	1	2	8.20	1	1	0	0	0	0	0	0	0
6324798	RPB8	NP_014867.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324569	RPB11	NP_014638.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323632	SPT5	NP_013703.1	6.30	5	11	0	0	0	2.54	2	2	0	0	0	8.94
6321625	TFG1	NP_011702.1	3.27	2	4	0	0	0	0	0	0	0	0	0	1.36
6321937	RPB12	NP_012013.1	38.57	2	2	0	0	0	0	0	0	0	0	0	0
398365043	TFG2	NP_011519.3	5.00	2	2	0	0	0	0	0	0	0	0	0	2.00
154199603	IWR1	NP_010168.4	0	0	0	0	0	0	0	0	0	0	0	0	0
398365457	SPT4	NP_011577.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325128	TAF14	NP_015196.1	6.56	1	4	0	0	0	0	0	0	0	0	0	6.56
398365155	NPA3	NP_012606.3	11.95	4	8	0	0	0	0	0	0	0	0	0	8.83
6323272	GPN3	NP_013344.1	4.78	1	2	0	0	0	0	0	0	0	0	0	0
6322163	IRR1	NP_012238.1	0	0	0	0	0	0	1.04	1	1	0	0	0	0
398366099	RRP41	NP_011711.3	0	0	0	0	0	0	0	0	0	0	0	0	0
398366555	UBA2	NP_010678.3	0	0	0	0	0	0	0	0	0	0	0	0	2.20
6323670	TAP42	NP_013741.1	3.55	1	1	0	0	0	0	0	0	0	0	0	7.65
6323004	SSA2	NP_013076.1	57.90	38	914	24.57	10	28	12.21	6	72	31.46	14	134	56.65
6319505	RPL4A	NP_009587.1	45.30	15	225	19.89	6	26	12.43	3	60	44.20	11	360	37.02
398364991	RPL4B	NP_010295.3	44.75	14	218	19.89	6	26	12.43	3	60	0	0	0	0
398365303	RPL43B	NP_012628.3	63.04	7	386	29.35	2	21	0	0	0	29.35	2	15	43.48
398364349	RPL2A	NP_116688.3	60.24	14	523	4.33	1	2	4.33	1	1	24.02	4	128	27.17
6325337	TEF1	NP_015405.1	29.26	13	270	7.64	4	8	7.21	4	47	41.92	18	593	44.10
144228166	SSA1	NP_009396.2	57.48	37	875	20.87	9	29	8.26	4	45	31.31	14	135	57.79
6319559	RPL19A	NP_009641.1	31.22	8	413	20.63	6	18	6.35	3	54	15.34	3	73	29.63
6320065	RPL35A	NP_010145.1	39.17	6	40	34.17	4	25	7.50	1	1	0	0	0	30.83
398365603	RPS4A	NP_012679.3	43.30	15	97	19.54	4	9	7.28	2	82	30.27	8	18	32.57
398365321	RPS6B	NP_009740.3	30.51	6	86	19.92	5	22	21.61	4	148	23.73	4	39	33.90
398364655	SSA4	NP_011029.3	17.45	13	222	7.79	3	3	1.87	1	1	10.44	4	56	28.19

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			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
398366603	RPS17B	NP_010735.3	54.41	8	129	53.68	7	28	11.76	1	19	22.06	2	5	52.94
6324533	RPS15	NP_014602.1	16.90	2	22	21.13	2	56	7.75	1	2	35.92	3	20	30.28
6323104	RPL10	NP_013176.1	37.10	11	179	4.98	1	6	9.50	2	129	19.91	4	246	27.60
6319396	SSA3	NP_009478.1	17.87	13	193	6.63	3	9	1.69	1	6	10.17	4	7	19.72
6321786	RPL14B	NP_011862.1	51.45	8	135	44.20	7	54	5.80	1	6	0	0	0	0
6322847	RPL14A	NP_012920.1	51.45	8	141	44.20	7	54	5.80	1	6	41.30	5	141	45.65
6324670	RPS7A	NP_014739.1	26.84	5	78	29.47	6	21	0	0	0	51.05	8	32	31.05
6322668	RPL17A	NP_012741.1	49.46	8	271	22.83	3	17	12.50	2	20	52.72	7	209	36.41
398365285	CAX4	NP_011550.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322284	RPL17B	NP_012358.1	49.46	8	262	22.83	3	17	12.50	2	20	52.72	7	210	36.41
398364883	RPS7B	NP_014303.3	35.79	6	71	37.37	7	29	6.84	1	1	36.84	5	21	40.00
6323791	RPL13B	NP_013862.1	42.71	8	172	40.20	8	46	11.06	2	169	0	0	0	0
6320121	RPL13A	NP_010201.1	42.71	8	172	38.69	7	44	11.06	2	169	21.61	4	160	37.69
6324120	SSB2	NP_014190.1	54.98	32	913	21.53	9	46	14.03	8	104	46.82	23	291	58.73
6323577	RPS1B	NP_013648.1	39.22	9	77	21.96	5	14	11.37	3	32	18.43	3	7	35.29
6325157	ATG21	NP_015225.1	5.24	2	2	0	0	0	0	0	0	0	0	0	0
6323474	RPS1A	NP_013546.1	39.61	9	48	21.96	5	12	14.90	4	25	18.43	3	3	27.45
398364725	RPL23B	NP_011042.3	43.80	6	161	31.39	3	9	7.30	1	3	40.15	7	54	43.80
6319972	SSB1	NP_010052.1	54.98	32	881	23.65	10	47	14.03	8	104	46.82	23	301	58.73
154200011	RPL20B	NP_014957.2	38.95	7	150	23.84	4	14	26.74	5	199	24.42	5	375	30.23
6322989	UBI4	NP_013061.1	50.00	4	30	32.89	2	10	0	0	0	0	0	0	38.16
9755341	RPS16A	NP_013863.2	54.55	8	40	45.45	6	30	0	0	0	37.06	4	9	32.17
6319378	RPL32	NP_009460.1	44.62	6	99	16.15	2	5	6.15	1	1	6.15	1	4	23.85
6325061	APL5	NP_015129.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364505	RPS3	NP_014221.3	40.42	8	77	17.50	3	14	26.25	5	154	35.42	6	15	39.17
398365351	RPS13	NP_010349.3	56.29	8	94	41.06	5	10	13.25	2	4	23.84	4	84	45.70
6324260	RPL16B	NP_014330.1	31.82	8	94	13.64	2	5	8.08	2	40	27.78	5	15	27.27
398365605	RPS30B	NP_014825.3	30.16	2	16	15.87	1	5	0	0	0	0	0	0	15.87
6323481	RPL6B	NP_013553.1	32.39	7	56	14.20	2	4	29.55	6	49	20.45	3	19	37.50
398365377	RPS9B	NP_009748.3	34.36	9	62	0	0	0	12.82	3	126	18.46	3	18	30.26
6324265	YDJ1	NP_014335.1	30.56	12	81	0	0	0	6.85	2	5	29.10	7	65	27.87
398364745	ATP7	NP_012909.3	7.47	1	2	8.05	2	6	0	0	0	0	0	0	7.47
6321464	RPS25A	NP_011541.1	39.81	6	51	31.48	4	13	8.33	1	6	12.04	1	1	38.89
6323077	RPS0B	NP_013149.1	19.84	4	60	16.67	3	20	0	0	0	29.76	5	85	29.37
398365071	RPS11A	NP_010308.3	32.05	6	47	0	0	0	0	0	0	6.41	1	1	25.64
6321653	RPS0A	NP_011730.1	19.84	4	60	16.67	3	20	0	0	0	29.76	5	85	29.37
6323465	CNA1	NP_013537.1	3.62	2	2	1.99	1	1	1.99	1	1	0	0	0	7.23

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			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6321335	RPL28	NP_011412.1	33.56	5	87	6.04	1	4	10.07	1	33	10.07	1	8	16.11
6322505	SSC1	NP_012579.1	45.26	27	231	9.79	4	18	4.89	3	7	15.14	7	13	45.87
6323567	RPL6A	NP_013638.1	22.16	6	35	8.52	1	2	15.91	3	28	14.77	2	6	25.57
6321754	RPL8A	NP_011830.1	50.00	14	222	30.86	9	22	10.94	3	60	28.52	7	164	42.19
6322984	RPL8B	NP_013055.1	50.00	14	227	34.38	10	23	14.45	4	70	28.52	7	165	49.22
6323615	RPS18B	NP_013686.1	54.11	9	55	17.81	3	12	15.07	2	30	29.45	4	18	43.15
398366307	RPA190	NP_014986.3	6.49	9	14	0	0	0	1.38	2	2	0	0	0	8.71
6324637	RPL3	NP_014706.1	38.76	16	266	18.35	6	29	6.72	3	20	21.19	8	43	27.13
6320128	RPL31A	NP_010208.1	55.75	7	52	31.86	3	66	0	0	0	7.96	1	2	38.94
398366143	RPL31B	NP_013510.3	55.75	7	48	23.89	2	56	0	0	0	0	0	0	0
398365761	RPS23A	NP_011633.3	26.21	4	30	20.00	3	8	0	0	0	7.59	1	24	33.79
6322137	RPL34B	NP_012212.1	6.61	1	33	6.61	1	4	6.61	1	1	9.09	2	7	6.61
6321315	RPS2	NP_011392.1	16.14	4	27	4.33	1	5	4.33	1	3	16.93	3	26	11.02
6320949	RPS8B	NP_011028.1	39.00	6	100	0	0	0	12.00	3	50	20.00	3	24	35.50
6321798	RPL27A	NP_011874.1	38.97	6	115	10.29	1	1	10.29	1	23	0	0	0	0
6325176	RPS9A	NP_015244.1	35.03	8	30	0	0	0	13.71	2	22	19.29	3	6	21.83
86558907	RHR2	NP_012211.2	20.80	4	10	9.60	2	3	0	0	0	10.80	1	1	32.80
6322058	RPL16A	NP_012133.1	36.18	9	73	14.07	2	3	8.04	2	33	27.14	6	21	31.66
6324027	RPS19B	NP_014097.1	56.94	7	26	47.92	5	37	18.75	2	5	25.69	3	14	45.14
6323090	RPL22A	NP_013162.1	38.02	3	8	55.37	3	37	14.05	1	1	41.32	2	9	55.37
6320142	RPS29B	NP_010222.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324807	KIN4	NP_014876.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321522	RPL11B	NP_011599.1	36.78	8	43	24.71	4	11	12.64	2	13	24.71	4	81	31.03
6322583	RPS5	NP_012657.1	34.67	7	71	25.78	5	33	0	0	0	29.33	5	23	29.78
6325126	RPL5	NP_015194.1	29.97	7	92	8.75	2	7	3.37	1	1	22.90	3	7	19.19
6323073	PDC1	NP_013145.1	36.06	17	450	27.53	10	71	7.64	4	30	48.31	19	321	43.69
398366277	VMA4	NP_014977.3	53.65	14	154	32.62	7	31	13.73	3	21	23.18	5	9	53.22
6323824	HOT1	NP_013895.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325066	NAB3	NP_015134.1	1.12	1	1	0	0	0	0	0	0	0	0	0	0
398364517	RPS24A	NP_010997.3	31.85	5	127	17.78	2	19	6.67	1	3	17.78	2	39	31.85
6321362	RPL7A	NP_011439.1	39.75	8	112	18.03	3	5	9.02	2	7	38.52	8	58	43.44
6325058	RPL7B	NP_015126.1	39.75	8	54	18.03	3	5	9.02	2	8	38.52	8	49	43.44
37362649	GUS1	NP_011269.2	21.33	14	47	1.84	1	1	3.11	2	14	5.65	4	18	33.76
398365909	RPL24B	NP_011664.3	31.61	6	58	0	0	0	0	0	0	20.65	3	40	15.48
6324707	EFT1	NP_014776.1	44.30	34	751	20.19	14	35	11.64	8	62	11.28	8	93	48.93
6323376	RPL26A	NP_013448.1	48.82	8	59	29.92	6	13	6.30	1	24	0	0	0	0
116006491	RPL26B	NP_011548.2	49.61	9	52	22.05	5	9	6.30	1	24	32.28	7	43	35.43

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			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6325006	RPL36B	NP_015074.1	44.00	7	141	50.00	8	36	31.00	3	51	33.00	4	73	40.00
6323848	RPL36A	NP_013920.1	43.00	6	112	40.00	6	33	31.00	3	37	0	0	0	0
398366579	RPL12B	NP_010706.3	53.94	7	166	24.24	3	12	9.09	1	1	30.91	4	57	49.70
6321540	NOP7	NP_011617.1	5.45	2	2	0	0	0	0	0	0	0	0	0	1.49
398365599	PDC6	NP_011601.3	6.93	3	103	3.37	1	3	1.60	1	25	8.17	3	52	6.93
6323163	PDC5	NP_013235.1	5.15	2	53	5.15	2	5	1.78	1	2	12.79	4	20	5.15
6323583	CMP2	NP_013655.1	7.62	4	10	0	0	0	0	0	0	0	0	0	7.78
398365991	CLC1	NP_011683.3	25.32	5	8	12.45	2	10	0	0	0	0	0	0	6.87
37362619	RXT2	NP_009653.2	4.65	2	2	0	0	0	0	0	0	0	0	0	5.12
10383793	RPS14A	NP_009960.2	49.64	8	61	37.96	5	15	25.55	4	34	59.85	7	24	40.15
6322271	RPS22A	NP_012345.1	24.62	3	52	24.62	3	13	0	0	0	25.38	2	3	24.62
6321407	RPL24A	NP_011484.1	32.26	7	92	5.81	1	1	6.45	1	2	20.65	3	42	22.58
6323357	RPL38	NP_013429.1	41.03	3	42	28.21	2	6	0	0	0	35.90	2	10	32.05
398366183	HTB1	NP_010510.3	31.30	5	15	25.95	4	9	0	0	0	11.45	1	1	39.69
398364287	GCN1	NP_011320.3	3.29	9	15	0	0	0	1.27	3	11	1.98	4	8	8.16
6320016	VMA1	NP_010096.1	28.48	29	176	0.84	1	1	2.80	3	3	13.63	11	88	33.43
14318481	TUB2	NP_116616.1	20.57	8	70	0	0	0	1.97	1	3	29.98	9	88	27.79
398364585	KAR2	NP_012500.3	30.65	20	121	0	0	0	3.67	2	10	4.40	2	56	32.40
6323196	RPS31	NP_013268.1	10.53	1	3	0	0	0	0	0	0	0	0	0	10.53
6321013	PAB1	NP_011092.1	47.14	24	171	27.21	10	23	5.03	2	13	4.51	2	67	35.18
6324993	FUM1	NP_015061.1	2.66	1	1	0	0	0	0	0	0	0	0	0	2.66
398366075	RPS29A	NP_013492.3	14.29	1	2	14.29	1	5	0	0	0	0	0	0	14.29
6322755	YJU2	NP_012828.1	0	0	0	0	0	0	0	0	0	0	0	0	11.15
398366059	PLP2	NP_014924.3	8.39	2	3	0	0	0	0	0	0	0	0	0	7.34
6320718	SMT3	NP_010798.1	0	0	0	0	0	0	0	0	0	0	0	0	6.93
398365031	YBR139W	NP_009697.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6321850	FYV4	NP_011926.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366083	TDH3	NP_011708.3	41.27	14	233	55.12	20	643	12.05	4	85	62.65	14	151	49.70
6319279	CDC19	NP_009362.1	48.20	22	147	53.80	24	121	18.40	8	26	38.20	16	363	55.60
6322468	TDH2	NP_012542.1	41.27	13	231	49.70	16	639	12.05	4	85	34.64	10	102	47.59
6321249	RPS26A	NP_011326.1	26.05	3	41	20.17	2	2	20.17	2	7	0	0	0	0
6321772	RPS20	NP_011848.1	37.19	4	27	37.19	5	17	12.40	2	3	36.36	4	17	45.45
14318474	SEC53	NP_116609.1	22.05	5	8	0	0	0	0	0	0	0	0	0	31.10
6320589	YRA1	NP_010669.1	14.16	3	11	0	0	0	0	0	0	3.54	1	1	20.80
6320805	ECM10	NP_010884.1	12.27	9	93	0	0	0	1.24	1	1	0	0	0	0
6323776	ECM16	NP_013847.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320596	RVS167	NP_010676.1	5.60	2	5	0	0	0	0	0	0	6.43	3	3	4.98

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
398366359	SLH1	NP_011787.4	0.81	1	1	0	0	0	0	0	0	0	0	0	0.46
6324321	SIS1	NP_014391.1	42.33	11	69	17.05	4	10	3.41	1	1	0	0	0	26.14
6319447	NCL1	NP_009529.1	5.99	3	5	0	0	0	4.82	2	4	0	0	0	16.52
6323112	EMP70	NP_013184.1	8.25	4	4	0	0	0	0	0	0	0	0	0	2.40
6324595	SFM1	NP_014664.1	13.62	3	4	0	0	0	0	0	0	0	0	0	0
330443411	UBP13	NP_009486.3	0	0	0	0	0	0	0	0	0	0	0	0	1.87
6323868	EFR3	NP_013939.1	0	0	0	0	0	0	1.79	1	2	0	0	0	0
330443676	CDC34	NP_013418.2	5.00	2	7	0	0	0	0	0	0	0	0	0	10.58
6320615	TRS120	NP_010695.1	0	0	0	0	0	0	0	0	0	0	0	0	0.93
6323487	FMP27	NP_013559.1	0.49	1	1	0	0	0	0	0	0	0	0	0	0.34
6320231	REG1	NP_010311.1	0.89	1	5	1.28	1	1	0	0	0	0.89	1	1	0
6324585	AUS1	NP_014654.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325048	RKM1	NP_015116.1	6.86	4	5	0	0	0	0	0	0	0	0	0	0
6323732	SEG1	NP_013803.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364779	YGL041W	NP_00110617	5.19	1	1	18.83	2	6	0	0	0	0	0	0	0
398365827	CAF130	NP_011650.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325063	RSA1	NP_015131.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365559	MSH6	NP_010382.3	1.61	2	2	0	0	0	0	0	0	0	0	0	0
398365727	YNR034W	NP_061494.3	0	0	0	0	0	0	0	0	0	0	0	0	40.82
6323683	YET2	NP_013754.1	4.38	1	1	0	0	0	0	0	0	0	0	0	0
330443366	GPB2	NP_009345.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325012	HUT1	NP_015080.1	0	0	0	0	0	0	0	0	0	0	0	0	0
99030927	KKQ8	NP_012753.2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364567	MAD1	NP_011429.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6320763	RMD6	NP_010842.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365009	MEC1	NP_009694.3	17.02	39	241	0	0	0	0	0	0	0	0	0	28.25
398364407	RPL9A	NP_011368.3	29.32	5	44	19.90	3	13	6.28	1	12	35.08	6	74	48.69
6324298	HHF2	NP_014368.1	40.78	4	26	0	0	0	9.71	1	4	0	0	0	40.78
6324262	RPL9B	NP_014332.1	19.37	4	38	9.95	2	5	6.28	1	12	25.13	5	47	48.69
398364523	TDH1	NP_012483.3	22.59	6	83	31.63	8	273	9.34	3	84	24.70	8	69	28.92
398364951	VMA2	NP_009685.3	59.77	27	245	15.09	5	12	7.54	3	6	43.71	16	152	60.54
6321333	ARC1	NP_011410.1	9.84	3	7	0	0	0	6.91	2	7	2.66	1	1	17.29
6321408	RPL30	NP_011485.1	56.19	4	76	29.52	2	7	0	0	0	21.90	2	22	38.10
6324445	RPL25	NP_014514.1	42.96	6	106	24.65	3	14	24.65	3	21	18.31	2	19	35.21
6325114	RPL33A	NP_015182.1	25.23	3	34	0	0	0	9.35	1	3	9.35	1	2	24.30
398365881	RPL33B	NP_014877.3	24.30	2	25	0	0	0	9.35	1	3	9.35	1	2	24.30
398364339	THS1	NP_116578.3	17.03	11	31	2.18	1	1	1.91	1	1	5.59	3	5	31.74

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
398365335	YNL033W	NP_014365.3	0	0	0	0	0	0	0	0	0	0	0	0	0
33438893	YPL119C-	NP_878181.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322579	JHD2	NP_012653.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324141	KAR1	NP_014211.1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383756	LRE1	NP_009879.2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364917	NHP10	NP_010282.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323035	NSE1	NP_013107.1	0	0	0	0	0	0	0	0	0	0	0	0	0
50593425	OMA1	NP_013013.2	0	0	0	0	0	0	0	0	0	0	0	0	0
6321910	ORC6	NP_011986.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322494	PET191	NP_012568.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365057	SPO14	NP_012956.3	0	0	0	0	0	0	0	0	0	0	0	0	0
398364971	TRP1	NP_010290.3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365003	PSF1	NP_010296.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6321697	RAD2	NP_011774.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319525	REG2	NP_009606.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320271	RTR2	NP_010351.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324162	SKO1	NP_014232.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366633	SNM1	NP_010766.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322778	STB6	NP_012851.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325081	TRE1	NP_015149.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324447	TRM13	NP_014516.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323236	HRD3	NP_013308.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325184	UBP16	NP_015253.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364885	UBP5	NP_011071.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325071	UIP4	NP_015138.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365127	UIP5	NP_012970.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6319904	YIH1	NP_009985.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325266	SUT2	NP_015334.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324449	HRP1	NP_014518.1	9.74	3	6	4.31	1	1	0	0	0	0	0	0	3.18
6323278	YEF3	NP_013350.1	26.05	26	305	2.49	2	4	9.39	9	161	29.98	19	222	43.68
6322084	PFK26	NP_012159.1	22.73	18	84	0	0	0	5.68	3	25	1.21	1	1	37.85
6323057	RPL15A	NP_013129.1	26.47	6	72	3.43	1	1	0	0	0	20.59	4	46	25.49
6323769	RPL15B	NP_013840.1	26.47	6	46	0	0	0	0	0	0	20.59	4	10	25.49
6319282	FUN12	NP_009365.1	5.89	5	12	0	0	0	1.30	1	2	0.90	1	2	3.09
6320190	NOP1	NP_010270.1	11.01	4	7	0	0	0	0	0	0	0	0	0	35.17
6323654	CLU1	NP_013725.1	5.40	6	9	0	0	0	4.62	4	11	1.64	2	2	16.60
6324886	NOP58	NP_014955.1	12.72	6	14	0	0	0	5.87	2	8	0	0	0	20.74

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6321807	DED81	NP_011883.1	24.37	10	33	2.17	1	2	7.40	3	7	0	0	0	33.57
398364767	SUI2	NP_012540.3	9.54	2	5	0	0	0	3.62	1	2	0	0	0	18.75
6323011	DPS1	NP_013083.1	32.68	16	46	0	0	0	5.75	2	5	3.23	2	13	35.01
6323885	RRP5	NP_013956.1	4.80	8	12	0	0	0	1.91	3	4	0.52	1	1	8.62
398366547	RPP2B	NP_010670.3	98.18	8	114	60.00	3	19	23.64	2	30	23.64	2	3	81.82
6323288	HSP60	NP_013360.1	49.65	25	160	30.07	13	45	13.99	7	50	18.36	9	18	44.76
6321599	NSR1	NP_011675.1	17.87	7	10	0	0	0	3.38	1	15	0	0	0	13.04
6320863	GCD11	NP_010942.1	13.28	5	7	0	0	0	3.61	1	1	1.71	1	11	16.70
398366591	NPL3	NP_010720.3	13.77	4	16	6.52	2	2	3.38	1	2	11.35	3	18	27.05
6319483	IPP1	NP_009565.1	20.21	5	6	17.07	3	8	4.18	1	1	0	0	0	9.76
6325221	PMA2	NP_015289.1	17.53	16	229	1.37	1	1	4.01	3	71	0	0	0	0
6322745	GFA1	NP_012818.1	13.67	8	33	0	0	0	3.49	2	3	15.20	7	22	32.50
398364525	LSG1	NP_011416.3	5.16	3	3	0	0	0	2.19	1	1	0	0	0	3.91
6325038	SAR1	NP_015106.1	19.47	3	29	0	0	0	6.84	1	8	13.68	2	104	46.32
6323840	HSC82	NP_013911.1	48.09	34	428	9.22	5	17	19.15	13	109	11.21	5	8	41.28
6321881	GAR1	NP_011957.1	11.71	2	2	0	0	0	0	0	0	10.24	1	1	14.15
6320164	NAT1	NP_010244.1	1.41	1	1	0	0	0	3.16	2	4	0	0	0	5.62
398366403	RPS12	NP_015014.3	26.57	5	44	18.88	2	2	8.39	1	2	0	0	0	29.37
6322830	CDC16	NP_012903.1	20.24	15	92	0	0	0	0	0	0	0	0	0	24.52
330443397	ATP1	NP_009453.2	28.99	16	123	7.52	3	4	11.93	6	27	32.84	16	287	37.25
6324709	IDH2	NP_014779.1	3.79	2	2	4.88	1	2	0	0	0	8.13	2	3	14.91
6325178	RPL21B	NP_015246.1	39.38	7	198	16.88	3	5	16.25	3	89	0	0	0	0
6321903	UBA4	NP_011979.1	25.91	13	108	0	0	0	5.91	2	2	0	0	0	28.86
6325138	DBP1	NP_015206.1	8.10	5	36	0	0	0	0	0	0	0	0	0	0
6325248	TAE2	NP_015316.1	0	0	0	0	0	0	0	0	0	0	0	0	3.18
330443505	DBP10	NP_010253.2	0	0	0	0	0	0	0	0	0	0.90	1	1	1.31
6320148	PSA1	NP_010228.1	22.16	6	40	4.99	1	5	0	0	0	0	0	0	42.66
6325016	HSP82	NP_015084.1	46.12	32	408	6.77	4	15	17.63	12	108	6.91	4	6	42.60
6320874	ARB1	NP_010953.1	25.41	14	30	0	0	0	6.56	3	27	13.77	6	128	19.84
6319673	PGI1	NP_009755.1	7.40	4	7	7.40	3	37	2.71	1	4	0	0	0	7.04
398365251	GSC2	NP_011546.3	6.12	10	41	0	0	0	1.21	2	2	0	0	0	11.50
6325018	SUI3	NP_015087.1	14.74	4	14	4.21	1	2	6.67	2	2	0	0	0	33.68
6323374	FKS1	NP_013446.1	12.90	18	83	0	0	0	1.23	2	6	0	0	0	18.28
14318479	ACT1	NP_116614.1	40.00	13	158	7.73	3	4	8.80	3	6	28.80	6	52	37.33
6323554	TUB1	NP_013625.1	20.13	7	32	0	0	0	2.01	1	2	16.55	4	22	26.85
6321724	ZUO1	NP_011801.1	26.33	9	19	0	0	0	5.77	2	7	12.01	3	4	30.95
6320842	NUG1	NP_010921.1	13.27	5	6	0	0	0	0	0	0	1.92	1	2	21.15

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6321027	DMC1	NP_011106.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364483	HOR2	NP_010984.3	10.00	2	5	4.00	1	2	0	0	0	0	0	0	0
398365429	PDR5	NP_014796.3	12.97	17	59	0	0	0	0.73	1	1	0	0	0	11.25
6325245	RRP12	NP_015313.1	1.22	1	1	0	0	0	2.36	2	2	0	0	0	2.12
398364937	TOP2	NP_014311.3	0.84	1	2	0	0	0	0	0	0	0	0	0	5.25
6324660	TCB1	NP_014729.1	4.05	4	9	0	0	0	1.26	1	1	0	0	0	2.78
6320056	COP1	NP_010136.1	12.49	13	43	0	0	0	0	0	0	1.50	1	3	10.82
398365597	BMH2	NP_010384.3	36.26	11	47	23.08	5	13	5.13	1	1	4.03	1	1	21.98
6325030	NEW1	NP_015098.1	14.88	15	41	0	0	0	3.09	3	3	1.59	2	10	16.72
6321809	RPS27B	NP_011885.1	18.29	2	26	0	0	0	0	0	0	9.76	1	3	18.29
6321012	CHD1	NP_011091.1	0.89	1	1	0	0	0	0	0	0	0	0	0	0.89
6320499	SSD1	NP_010579.1	1.92	2	2	0	0	0	0	0	0	0	0	0	6.16
6323072	TRX1	NP_013144.1	66.02	6	40	12.62	1	1	33.98	3	5	0	0	0	66.02
6325238	VTC3	NP_015306.1	0.96	1	1	0	0	0	0	0	0	0	0	0	3.71
398365319	MAK21	NP_010345.3	0	0	0	0	0	0	1.27	1	2	0	0	0	2.34
398364327	LSB3	NP_219497.4	14.60	5	7	0	0	0	3.92	1	1	3.92	1	1	8.93
6319513	ATP3	NP_009595.1	31.51	10	41	10.61	2	9	12.22	3	32	18.65	5	11	40.84
6324140	SRP1	NP_014210.1	8.12	4	10	0	0	0	2.40	1	1	1.48	1	1	7.20
6324297	HHT2	NP_014367.1	11.76	2	10	0	0	0	0	0	0	0	0	0	0
6323306	DBP9	NP_013378.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324039	RFC3	NP_014109.1	0	0	0	0	0	0	3.82	1	1	0	0	0	0
6321193	RTF1	NP_011270.1	1.79	1	3	0	0	0	0	0	0	0	0	0	24.91
398366149	TRX2	NP_011725.3	66.35	6	50	11.54	1	5	34.62	3	6	0	0	0	66.35
6322452	CCT8	NP_012526.1	24.65	12	30	0	0	0	4.58	2	6	7.92	3	5	30.46
6321679	PFK1	NP_011756.1	16.11	12	31	2.03	1	1	2.63	2	5	1.11	1	1	26.65
6324219	NOP15	NP_014289.1	4.09	1	2	0	0	0	0	0	0	0	0	0	0
330443717	CTR9	NP_014496.2	0.84	1	1	0	0	0	0	0	0	0	0	0	0.84
6323755	ILV2	NP_013826.1	13.97	9	52	0	0	0	5.09	3	54	13.39	5	21	28.24
6320276	PAA1	NP_010356.1	17.80	3	10	0	0	0	0	0	0	0	0	0	16.23
6681849	RPL42A	NP_014237.2	15.09	2	3	0	0	0	0	0	0	0	0	0	22.64
14318517	SEC4	NP_116650.1	37.21	7	34	0	0	0	5.58	1	1	5.58	1	7	50.23
6320400	MSS116	NP_010480.1	0	0	0	0	0	0	3.77	2	4	0	0	0	14.16
6321389	TIF4632	NP_011466.1	0	0	0	0	0	0	0	0	0	0	0	0	0.98
6324081	RPA49	NP_014151.1	2.41	1	2	0	0	0	0	0	0	0	0	0	3.61
398366347	MES1	NP_011780.3	12.78	10	32	0	0	0	2.80	2	2	3.33	2	5	22.24
6320418	TCP1	NP_010498.1	14.49	7	30	0	0	0	3.94	2	21	0	0	0	23.61
398365595	PIL1	NP_011600.3	29.50	9	49	2.36	1	1	5.90	2	4	17.11	5	25	39.53

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
398364417	SAH1	NP_010961.3	13.14	5	12	7.80	3	10	2.67	1	1	0	0	0	5.12
6320396	RVB1	NP_010476.1	16.41	6	25	0	0	0	10.37	4	13	21.17	5	8	40.82
6320077	CDC48	NP_010157.1	31.62	20	138	12.22	8	17	3.23	2	7	12.22	5	9	35.33
6325025	FAS2	NP_015093.1	13.25	20	37	4.40	7	16	0.64	1	3	2.38	4	4	21.78
6324517	THI20	NP_014586.1	7.99	4	4	0	0	0	0	0	0	0	0	0	47.19
6323030	NOC3	NP_013102.1	1.66	1	1	0	0	0	0	0	0	0	0	0	2.41
6320143	TSR1	NP_010223.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321318	PRP43	NP_011395.1	2.87	2	4	0	0	0	0	0	0	0	0	0	3.26
6319618	MAK5	NP_009700.1	0	0	0	0	0	0	0	0	0	0	0	0	1.42
398364755	URB1	NP_012911.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6320780	POL5	NP_010859.1	2.54	3	3	0	0	0	0	0	0	0.98	1	1	1.47
6325021	RVB2	NP_015089.1	28.45	11	28	0	0	0	5.10	2	14	4.46	2	12	38.85
6322350	CCT7	NP_012424.1	16.55	8	14	0	0	0	9.64	4	7	2.18	1	1	34.91
14270688	UTP20	NP_009551.2	0.40	1	1	0	0	0	0	0	0	0	0	0	0.64
6324197	KRE33	NP_014267.1	3.31	3	3	0	0	0	0	0	0	0	0	0	3.31
6321816	RPN1	NP_011892.1	14.70	13	36	0.70	1	1	4.33	5	19	4.53	3	5	17.22
6322901	NAP1	NP_012974.1	31.41	9	55	9.59	2	4	4.08	1	2	0	0	0	36.93
6320030	GLT1	NP_010110.1	7.88	14	19	0	0	0	1.68	3	12	0.47	1	7	17.06
6323470	CAR2	NP_013542.1	3.07	1	1	0	0	0	0	0	0	0	0	0	6.84
6320956	KAP123	NP_011035.1	14.29	13	39	0	0	0	3.50	3	16	0.99	1	1	17.43
398365941	SEC7	NP_010454.3	3.58	7	12	0	0	0	0	0	0	0	0	0	0.45
6323324	GSP1	NP_013396.1	20.55	4	15	10.05	2	6	0	0	0	9.59	2	16	24.20
6322697	GPM1	NP_012770.1	48.99	11	213	46.56	10	75	0	0	0	23.89	5	20	32.39
6325013	SRP68	NP_015081.1	1.34	1	1	0	0	0	2.17	1	2	0	0	0	4.51
6322103	AVT7	NP_012178.1	1.63	1	1	0	0	0	1.63	1	1	1.63	1	5	1.63
6322049	CCT2	NP_012124.1	15.75	7	31	0	0	0	2.28	1	1	10.44	4	6	41.56
330443694	SLA2	NP_014156.2	2.07	2	14	0	0	0	0	0	0	0	0	0	15.19
6321804	YSC84	NP_011880.1	4.27	2	3	0	0	0	0	0	0	0	0	0	3.85
398366021	CCT6	NP_010474.3	6.96	4	9	0	0	0	0	0	0	2.93	1	1	21.79
6324042	SEC21	NP_014112.1	6.74	5	8	0	0	0	1.82	1	1	1.28	1	2	10.05
398364741	SCS2	NP_011046.3	17.21	3	7	0	0	0	11.48	2	9	0	0	0	17.21
6325253	LSP1	NP_015321.1	35.19	11	57	3.52	1	3	0	0	0	9.09	3	35	38.71
6321390	RPT6	NP_011467.1	15.80	6	8	0	0	0	6.67	2	4	11.36	3	6	17.04
398364807	PNC1	NP_011478.3	0	0	0	0	0	0	0	0	0	0	0	0	17.59
6325164	NOG1	NP_015232.1	5.26	3	6	0	0	0	0	0	0	0	0	0	8.81
6319814	GRX1	NP_009895.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322823	MAE1	NP_012896.1	12.56	7	12	0	0	0	4.04	2	8	18.98	8	49	18.09

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Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6320255	TPI1	NP_010335.1	23.79	5	15	18.15	4	20	0	0	0	0	0	0	14.52
6320983	GDI1	NP_011062.1	5.32	2	3	1.77	1	1	0	0	0	0	0	0	8.65
398365205	RPT5	NP_014760.3	30.88	10	22	0	0	0	8.29	3	6	24.19	9	21	36.87
6324992	KEL3	NP_015060.1	0	0	0	0	0	0	0	0	0	0	0	0	1.54
6323028	DNM1	NP_013100.1	5.94	4	5	0	0	0	1.59	1	2	0	0	0	3.43
398365977	RPT4	NP_014902.3	7.78	3	7	0	0	0	0	0	0	9.61	3	6	10.07
6320058	CCT4	NP_010138.1	9.66	4	12	0	0	0	5.30	2	23	2.84	1	6	17.05
6320379	HMO1	NP_010459.1	8.54	2	3	0	0	0	0	0	0	0	0	0	13.41
6322508	CYC1	NP_012582.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325137	VPS30	NP_015205.1	0	0	0	0	0	0	2.51	1	3	0	0	0	0
6321301	SEC27	NP_011378.1	13.39	11	57	0	0	0	3.04	2	12	1.12	1	1	13.39
6321673	YHB1	NP_011750.1	11.03	3	5	0	0	0	0	0	0	4.26	1	1	7.27
27469359	SYN8	NP_009388.2	2.75	1	2	0	0	0	0	0	0	0	0	0	0
6324433	CDC33	NP_014502.1	27.70	5	10	0	0	0	11.27	2	12	0	0	0	18.78
6323571	POB3	NP_013642.1	3.08	2	2	0	0	0	0	0	0	0	0	0	9.60
398365157	NOP2	NP_014338.3	3.72	2	2	0	0	0	0	0	0	0	0	0	8.09
6321228	YPT32	NP_011305.1	31.98	6	26	0	0	0	11.71	2	6	15.32	3	9	35.14
398366559	RPT3	NP_010682.3	2.80	1	2	0	0	0	2.80	1	1	12.62	3	13	18.69
6322704	RPT1	NP_012777.1	27.41	9	19	0	0	0	2.78	1	2	0	0	0	20.99
398366379	PRT1	NP_015006.3	13.50	10	19	0	0	0	4.98	3	12	1.31	1	1	27.65
398365125	PDA1	NP_011105.4	24.52	9	30	0	0	0	4.29	1	1	0	0	0	20.95
398364913	GND1	NP_012053.3	10.02	4	15	5.73	2	4	0	0	0	0	0	0	17.18
6322637	SAC1	NP_012710.1	13.48	6	9	0	0	0	0	0	0	0	0	0	6.74
398365933	CYS4	NP_011671.3	48.32	21	125	4.34	1	2	8.48	3	3	5.52	2	16	52.47
6323903	FAA4	NP_013974.1	7.49	5	20	0	0	0	1.44	1	6	1.73	1	2	11.10
6319806	PDI1	NP_009887.1	36.02	16	98	4.21	1	4	8.81	3	12	0	0	0	34.87
6323512	PHO84	NP_013583.1	0	0	0	0	0	0	0	0	0	2.90	1	1	0
6325367	RPC40	NP_015435.1	13.43	4	8	0	0	0	0	0	0	0	0	0	19.10
6319835	GBP2	NP_009916.1	2.11	1	1	0	0	0	0	0	0	0	0	0	5.85
6319290	MYO4	NP_009373.1	1.02	1	3	0	0	0	0	0	0	0	0	0	0
14318531	GCN20	NP_116664.1	5.19	4	6	0	0	0	3.32	2	6	1.20	1	4	6.38
6320869	YPT31	NP_010948.1	32.29	6	26	0	0	0	11.66	2	6	15.25	3	9	49.33
398365617	UTP22	NP_011604.3	0	0	0	0	0	0	0	0	0	0	0	0	1.62
398366209	RTN1	NP_010519.3	20.68	6	37	9.83	2	6	9.83	2	16	9.83	2	7	13.22
6325097	CDC60	NP_015165.1	17.43	17	40	2.39	1	1	2.11	2	7	2.02	1	1	17.61
6323294	RPS28B	NP_013366.1	31.34	2	45	41.79	3	7	13.43	1	15	0	0	0	31.34
6325151	SSE1	NP_015219.1	36.08	24	184	9.24	4	5	11.40	6	16	8.23	5	46	41.56

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Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
398364441	NOT3	NP_012226.3	3.71	2	4	0	0	0	0	0	0	0	0	0	13.16
398366191	RPN13	NP_013525.3	15.38	2	4	0	0	0	8.33	1	1	0	0	0	8.33
6321453	YGR016W	NP_011530.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320930	ILV1	NP_011009.1	18.06	8	11	0	0	0	2.43	1	3	13.19	5	76	33.85
6323016	PUF3	NP_013088.1	3.19	2	4	0	0	0	0	0	0	0	0	0	1.14
37362700	SRP72	NP_015114.2	0	0	0	0	0	0	0	0	0	0	0	0	0
6321326	TAF6	NP_011403.1	3.49	1	1	0	0	0	0	0	0	0	0	0	8.91
6322565	ADO1	NP_012639.1	15.59	4	11	7.35	2	5	0	0	0	0	0	0	0
6319722	ISW1	NP_009804.1	1.06	1	1	0	0	0	0.89	1	1	0	0	0	4.34
330443605	SET2	NP_012367.2	0	0	0	0	0	0	0	0	0	0	0	0	3.68
398365101	ARP3	NP_012599.3	9.58	4	10	2.45	1	1	0	0	0	0	0	0	6.68
6321780	STE20	NP_011856.1	3.30	2	2	0	0	0	0	0	0	0	0	0	0
6321776	PRS3	NP_011852.1	14.69	5	13	0	0	0	0	0	0	10.31	3	9	22.19
6323214	RPL37A	NP_013286.1	21.59	2	7	13.64	1	1	11.36	1	1	0	0	0	0
6321757	ECM29	NP_011833.1	0.75	1	1	0	0	0	0.59	1	1	0	0	0	0
398365069	ADH5	NP_009703.3	5.70	2	16	2.28	1	1	0	0	0	2.28	1	1	5.70
398365695	CPR8	NP_014425.3	3.57	1	1	0	0	0	0	0	0	0	0	0	0
6319854	RVS161	NP_009935.1	10.19	3	3	0	0	0	3.40	1	2	7.92	1	2	19.62
6324620	DBP5	NP_014689.1	19.50	7	16	0	0	0	5.19	2	2	11.62	4	6	17.43
6321399	YGL039W	NP_011476.1	6.90	2	3	0	0	0	0	0	0	0	0	0	9.20
398365061	SUP45	NP_009701.3	16.25	8	26	0	0	0	2.97	1	4	2.97	1	11	40.27
6324673	KTR1	NP_014742.1	15.01	6	16	0	0	0	0	0	0	0	0	0	16.54
6323387	ILV5	NP_013459.1	13.67	5	9	0	0	0	0	0	0	0	0	0	28.10
6323795	TIF34	NP_013866.1	20.46	6	22	0	0	0	2.31	1	1	2.31	1	1	38.62
6322667	PRS1	NP_012740.1	5.15	2	4	0	0	0	3.28	1	18	2.58	1	2	16.39
6320910	RGI1	NP_010990.1	8.07	1	2	0	0	0	0	0	0	0	0	0	0
398365575	HOM6	NP_012673.3	0	0	0	10.03	2	3	0	0	0	0	0	0	0
6323916	TIF11	NP_013987.1	6.54	1	1	0	0	0	0	0	0	0	0	0	15.03
6321612	RBG2	NP_011689.1	2.72	1	2	0	0	0	0	0	0	0	0	0	14.95
6323225	PWP1	NP_013297.1	11.28	5	7	0	0	0	3.99	1	1	0	0	0	13.89
6324737	SSP1	NP_014806.1	12.23	2	5	0	0	0	0	0	0	0	0	0	0
6323450	CDC73	NP_013522.1	5.09	2	2	0	0	0	0	0	0	5.34	1	1	5.60
398365461	SRP40	NP_013018.3	8.62	2	5	0	0	0	0	0	0	0	0	0	2.96
6320708	RPL37B	NP_010788.1	0	0	0	0	0	0	18.18	2	2	0	0	0	0
398364957	MKT1	NP_014314.3	6.14	4	5	0	0	0	2.53	1	3	1.20	1	1	4.10
6320079	YDL124W	NP_010159.1	7.37	2	2	0	0	0	0	0	0	0	0	0	4.49
398364371	RSC8	NP_116695.3	6.82	3	6	0	0	0	1.97	1	1	2.51	1	1	5.21

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Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6325418	GPH1	NP_015486.1	1.55	1	1	0	0	0	0	0	0	1.11	1	2	11.97
6321859	DYS1	NP_011935.1	14.21	5	11	0	0	0	0	0	0	0	0	0	8.01
6323967	GAS1	NP_014038.1	13.06	8	67	0	0	0	0	0	0	0	0	0	17.53
6319695	PYC2	NP_009777.1	10.00	10	21	3.81	4	11	1.44	2	2	3.73	3	5	10.68
6323642	YPT7	NP_013713.1	51.44	8	63	0	0	0	5.29	1	1	28.85	5	67	60.10
6320552	HXT3	NP_010632.1	2.12	1	10	0	0	0	2.12	1	1	0	0	0	2.12
6325196	ALD6	NP_015264.1	9.20	4	8	6.60	3	7	0	0	0	2.20	1	2	6.60
6319315	EFB1	NP_009398.1	17.96	3	10	8.25	1	11	0	0	0	10.68	1	1	23.30
10383811	CDC39	NP_010017.2	0.71	1	1	0	0	0	0	0	0	0	0	0	1.42
6321753	SBP1	NP_011829.1	3.06	1	6	0	0	0	0	0	0	0	0	0	23.47
6320764	DLD3	NP_010843.1	2.82	1	1	0	0	0	0	0	0	0	0	0	2.22
398366413	ERV29	NP_011800.3	0	0	0	0	0	0	3.87	1	2	0	0	0	0
6320969	GLO3	NP_011048.1	7.51	4	9	0	0	0	0	0	0	1.42	1	7	1.42
6319602	TPS1	NP_009684.1	13.54	5	9	0	0	0	0	0	0	0	0	0	15.76
6579193	NMD2	NP_011944.2	0	0	0	0	0	0	0	0	0	0	0	0	0
6321990	UTP9	NP_012066.1	4.35	2	2	0	0	0	0	0	0	0	0	0	7.30
398366561	SXM1	NP_010683.3	2.33	2	2	0	0	0	0	0	0	0	0	0	0
6320157	NPC2	NP_010237.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364687	NOP9	NP_012524.3	1.50	1	1	0	0	0	0	0	0	0	0	0	0
22532997	BUD27	NP_683715.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366449	BFR2	NP_010585.3	8.43	4	4	0	0	0	0	0	0	0	0	0	5.62
398365717	TRM1	NP_010405.3	13.16	7	14	0	0	0	0	0	0	0	0	0	5.44
6323362	CHS5	NP_013434.1	3.58	2	3	0	0	0	0	0	0	0	0	0	0
398364573	PAN1	NP_012271.3	8.18	9	52	0	0	0	0	0	0	0	0	0	5.20
6320720	GRX2	NP_010801.1	6.99	1	1	0	0	0	0	0	0	0	0	0	6.99
6321595	PTI1	NP_011672.1	1.65	1	1	0	0	0	0	0	0	0	0	0	0
398364853	OCA1	NP_014300.3	5.46	1	1	0	0	0	0	0	0	0	0	0	5.46
6323852	YMR196	NP_013923.1	2.76	2	2	0	0	0	0	0	0	0.92	1	1	0
33438816	YI002W-	NP_878098.1	17.39	1	1	0	0	0	0	0	0	0	0	0	0
398366575	RRP17	NP_010700.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323359	TMA10	NP_013431.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365853	WTM1	NP_014873.3	19.91	8	36	12.13	4	16	0	0	0	2.29	1	1	12.59
6320181	GPD1	NP_010262.1	4.09	2	11	0	0	0	0	0	0	0	0	0	31.20
6323518	NAB6	NP_013589.1	1.23	1	1	0	0	0	1.23	1	1	0.88	1	4	3.35
6321221	KIP3	NP_011299.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365131	RFC2	NP_012602.3	7.08	2	2	0	0	0	0	0	0	0	0	0	0
6322294	ERG20	NP_012368.1	2.84	1	2	0	0	0	0	0	0	7.95	1	1	12.22

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Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6321024	ECM32	NP_011103.1	0	0	0	0	0	0	0	0	0	0	0	0	0
37362660	FUR1	NP_011996.2	15.74	4	7	0	0	0	0	0	0	0	0	0	22.69
398364665	SEC18	NP_009636.3	6.20	4	12	0	0	0	1.72	1	1	5.28	3	5	18.73
6323391	ADE13	NP_013463.1	1.87	1	2	4.15	2	3	0	0	0	0	0	0	0
6323699	STV1	NP_013770.1	4.83	4	8	0	0	0	0	0	0	0	0	0	3.37
6321412	TRP5	NP_011489.1	26.31	14	59	2.83	2	2	3.25	2	10	1.98	1	2	35.50
6321973	OYE2	NP_012049.1	11.75	4	5	0	0	0	4.25	1	1	0	0	0	0
6321801	ARD1	NP_011877.1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383761	HIS4	NP_009900.2	5.26	4	5	7.13	4	12	0	0	0	0	0	0	8.14
398365759	SAC6	NP_010414.3	15.42	9	25	0	0	0	2.02	1	1	2.34	1	1	18.85
6321721	BGL2	NP_011798.1	10.22	3	7	4.15	1	6	0	0	0	3.83	1	1	7.99
398364979	CCZ1	NP_009689.3	1.42	1	1	0	0	0	0	0	0	0	0	0	1.42
42742256	VTC4	NP_012522.2	4.85	3	5	0	0	0	0	0	0	0	0	0	2.36
6321814	THR1	NP_011890.1	1.96	1	1	0	0	0	0	0	0	0	0	0	5.32
10383798	PHO87	NP_009966.2	0	0	0	0	0	0	0	0	0	0	0	0	0
6319847	CDC10	NP_009928.1	6.83	2	2	0	0	0	2.80	1	1	0	0	0	15.53
398365363	URA8	NP_012637.4	4.15	2	2	0	0	0	0	0	0	0	0	0	11.94
6323138	AHP1	NP_013210.1	5.11	1	1	25.00	3	7	0	0	0	0	0	0	29.55
6320657	UTP6	NP_010737.1	1.59	1	2	0	0	0	0	0	0	0	0	0	0
6321919	YHR127W	NP_011995.1	0	0	0	0	0	0	0	0	0	0	0	0	0
41629688	PSP2	NP_013695.2	0	0	0	0	0	0	0	0	0	0	0	0	1.69
398364633	APS3	NP_012510.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322843	MRT4	NP_012916.1	13.14	3	8	0	0	0	0	0	0	0	0	0	16.10
330443486	PAT1	NP_010002.3	3.27	2	3	0	0	0	0	0	0	1.76	1	1	2.01
37362659	RRP3	NP_011932.2	2.59	1	2	0	0	0	0	0	0	4.39	2	13	0
6320845	SEC3	NP_010924.1	0	0	0	0	0	0	0	0	0	0.60	1	8	0
6324791	RFC1	NP_014860.1	0	0	0	0	0	0	1.51	1	2	0	0	0	0
6325369	MRD1	NP_015437.1	0	0	0	0	0	0	0	0	0	0	0	0	1.58
6322635	YRA2	NP_012708.1	5.42	1	1	0	0	0	0	0	0	0	0	0	0
6320368	SEC1	NP_010448.1	0	0	0	0	0	0	1.80	1	1	0	0	0	0
398364935	UBP3	NP_011078.3	1.32	1	1	0	0	0	1.21	1	1	0	0	0	5.04
6320789	GLY1	NP_010868.1	22.22	7	13	0	0	0	0	0	0	0	0	0	17.57
6325172	SEC16	NP_015240.1	0.46	1	1	0	0	0	0	0	0	0	0	0	0
398364311	TMA20	NP_010923.3	6.63	1	1	0	0	0	0	0	0	0	0	0	12.71
6320926	UTP7	NP_011005.1	3.97	2	3	0	0	0	0	0	0	0	0	0	0
330443568	ENP2	NP_011661.2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324328	CIT1	NP_014398.1	2.51	1	1	3.13	1	1	0	0	0	0	0	0	2.09

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6320843	PAC2	NP_010922.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365529	SIR1	NP_013027.4	0	0	0	0	0	0	0	0	0	0	0	0	0
14318435	TIP20	NP_116577.1	1.71	1	1	0	0	0	0	0	0	0	0	0	0
6323645	AIM34	NP_013716.1	0	0	0	0	0	0	0	0	0	0	0	0	0
330443386	BUD14	NP_009408.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6319751	CHK1	NP_009833.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319452	APN2	NP_009534.1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383783	YCR015C	NP_009941.2	0	0	0	0	0	0	0	0	0	0	0	0	0
33438820	YJL133C-A	NP_878102.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365435	ERG25	NP_011574.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6321363	MPS2	NP_011440.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323180	PCD1	NP_013252.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364799	PMD1	NP_011058.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322709	TGL1	NP_012782.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325305	TAH18	NP_015373.1	1.77	1	1	0	0	0	0	0	0	0	0	0	0
6322698	YKL151C	NP_012771.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322341	RPE1	NP_012414.1	0	0	0	5.46	1	1	0	0	0	0	0	0	8.40
6323336	STT4	NP_013408.1	1.21	2	2	0	0	0	0	0	0	0	0	0	0
6320542	MSN5	NP_010622.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325171	ELP3	NP_015239.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319274	ERV46	NP_009358.1	6.51	2	2	0	0	0	0	0	0	0	0	0	3.13
398366207	CRM1	NP_011734.3	2.40	2	6	0	0	0	1.29	1	1	0	0	0	4.24
6321198	CSE1	NP_011276.1	3.54	3	4	0	0	0	0	0	1.25	1	5	2.92	
398366491	SKP1	NP_010615.3	15.46	2	7	0	0	0	0	0	0	0	0	0	36.08
6323819	MLH1	NP_013890.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365033	IRA1	NP_009698.3	0	0	0	0	0	0	0	0	0	0	0	0	0
398366427	SRP101	NP_010578.3	0	0	0	0	0	0	0	0	1.45	1	1	0	
6320008	NUS1	NP_010088.1	4.27	1	1	0	0	0	0	0	0	0	0	0	4.00
6324098	PDR16	NP_014168.1	5.70	2	2	0	0	0	3.42	1	4	0	0	0	17.38
6324093	SIN4	NP_014163.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322063	MET18	NP_012138.1	0	0	0	0	0	0	0	0	1.26	1	1	0	
6319901	THR4	NP_009982.1	6.03	4	10	3.70	1	2	0	0	0	0	0	0	7.39
6321899	CDC12	NP_011975.1	17.20	5	6	0	0	0	0	0	0	0	0	0	16.71
398364305	YND1	NP_010920.3	0	0	0	0	0	0	0	0	1.27	1	2	1.27	
6325365	RPN7	NP_015433.1	12.35	6	12	0	0	0	3.03	1	2	0	0	0	19.35
6322288	RFA3	NP_012362.1	38.52	3	3	0	0	0	0	0	0	0	0	0	12.30
6323832	YMR178	NP_013903.1	0	0	0	0	0	0	6.20	1	1	0	0	0	16.42

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
14318538	YFR016C	NP_116671.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321647	SER2	NP_011724.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365769	SHG1	NP_009817.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6321009	SPT2	NP_011088.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365509	RRP1	NP_010372.3	11.51	3	6	0	0	0	0	0	0	0	0	0	10.43
6323756	MYO5	NP_013827.1	9.68	10	17	0	0	0	0.90	1	1	0.90	1	3	4.10
6324839	RBL2	NP_014908.1	11.32	1	3	0	0	0	0	0	0	0	0	0	0
6319685	DUR1,2	NP_009767.1	0	0	0	0	0	0	0.71	1	1	0	0	0	0
6325273	TIF6	NP_015341.1	14.69	3	5	0	0	0	0	0	0	0	0	0	11.43
6324938	PRE10	NP_015007.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324996	APM1	NP_015064.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324522	GAL11	NP_014591.1	1.20	1	1	0	0	0	0	0	0	0	0	0	0
6324113	RAP1	NP_014183.1	3.63	2	2	0	0	0	1.33	1	3	0	0	0	5.93
398366009	YGR169C	NP_878080.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6319564	POL30	NP_009645.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323546	UTP14	NP_013617.1	1.22	1	1	0	0	0	0	0	0	0.67	1	1	1.11
6324312	PUB1	NP_014382.1	1.99	1	13	0	0	0	0	0	0	0	0	0	3.75
6325286	APL4	NP_015354.1	0.96	1	1	0	0	0	0	0	0	0	0	0	0
6323401	SSQ1	NP_013473.1	6.70	3	5	0	0	0	0	0	0	0	0	0	0
6324048	HCH1	NP_014118.1	7.84	1	2	0	0	0	0	0	0	0	0	0	7.84
6319627	APD1	NP_009709.1	3.80	1	1	0	0	0	0	0	0	0	0	0	7.28
6320102	DUN1	NP_010182.1	2.92	1	1	0	0	0	0	0	0	0	0	0	1.75
10383806	SOL2	NP_009999.2	4.44	1	1	0	0	0	0	0	0	0	0	0	0
6320205	YRB1	NP_010285.1	5.47	1	8	5.47	1	4	0	0	0	0	0	0	0
398366613	TOM1	NP_010745.3	0.86	3	3	0	0	0	0.76	2	5	0	0	0	1.32
398365693	TMA64	NP_010402.3	1.77	1	1	0	0	0	0	0	0	0	0	0	1.77
6323893	BCH1	NP_013964.1	1.24	1	1	0	0	0	0	0	0	0	0	0	1.24
398366023	YTM1	NP_014915.3	3.91	2	3	0	0	0	3.70	1	1	0	0	0	4.35
6319437	STU1	NP_009519.1	0	0	0	0	0	0	0	0	0	0	0	0	0.66
398364833	MET4	NP_014296.4	0	0	0	0	0	0	0	0	0	0	0	0	2.08
6321424	PUF4	NP_011501.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322411	MTR4	NP_012485.1	1.21	1	1	0	0	0	0	0	0	0	0	0	10.16
14318501	LPD1	NP_116635.1	12.63	6	9	0	0	0	0	0	0	0	0	0	17.03
6321890	SFB3	NP_011966.1	8.29	7	14	0	0	0	0	0	0	0	0	0	0.86
398366661	GIN4	NP_010795.3	0.70	1	2	0	0	0	0	0	0	0	0	0	1.31
398365725	SOL1	NP_014432.3	4.36	1	1	0	0	0	0	0	0	0	0	0	0
6323570	DAK1	NP_013641.1	14.73	7	13	1.71	1	1	2.23	1	1	0	0	0	16.10

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6322260	ECM25	NP_012334.1	0	0	0	0	0	0	0	0	0	0	0	0	1.84
6319691	SDS24	NP_009773.1	1.90	1	1	0	0	0	0	0	0	0	0	0	0
6322202	SQT1	NP_012277.1	5.80	2	3	0	0	0	0	0	0	3.48	1	2	15.08
6321961	THP2	NP_012037.1	0	0	0	0	0	0	4.21	1	1	0	0	0	8.05
398364887	CKB1	NP_011496.3	3.60	1	5	0	0	0	0	0	0	0	0	0	3.60
6320131	YET3	NP_010211.1	13.30	3	10	0	0	0	0	0	0	0	0	0	0
398364267	URA3	NP_010893.3	0	0	0	0	0	0	0	0	0	0	0	0	0
398366549	NKP1	NP_010671.4	6.72	1	2	0	0	0	0	0	0	0	0	0	3.36
398366429	ATF1	NP_015022.3	3.24	1	1	0	0	0	0	0	0	3.81	1	1	4.95
398365899	ABP140	NP_014882.4	3.66	2	2	0	0	0	0	0	0	0	0	0	3.18
6324659	OST3	NP_014728.1	3.43	1	5	0	0	0	0	0	0	0	0	0	0
398365193	TRS33	NP_014758.3	5.22	1	1	0	0	0	0	0	0	0	0	0	5.22
6322746	APE1	NP_012819.1	0	0	0	0	0	0	0	0	0	0	0	0	5.06
398365017	YGR001C	NP_011515.4	7.26	2	2	0	0	0	0	0	0	0	0	0	6.05
6319455	FUS3	NP_009537.1	7.08	3	3	0	0	0	0	0	0	0	0	0	3.40
6321351	MMS2	NP_011428.1	0	0	0	0	0	0	0	0	0	0	0	0	21.90
6319435	YBL036C	NP_009517.1	0	0	0	0	0	0	0	0	0	0	0	0	10.51
6321450	SNU71	NP_011527.1	0	0	0	0	0	0	0	0	0	0	0	0	3.06
6325394	RRP9	NP_015463.1	3.84	2	2	0	0	0	0	0	0	0	0	0	1.57
269970294	GCV3	NP_009355.3	0	0	0	0	0	0	0	0	0	0	0	0	4.71
6323352	MMS22	NP_013424.1	0	0	0	0	0	0	0	0	0	0	0	0	0.55
330443445	PBY1	NP_009652.2	0	0	0	0	0	0	0	0	0	0	0	0	1.46
398364685	NAM9	NP_014262.3	2.26	1	1	0	0	0	0	0	0	0	0	0	2.06
6319999	YML11	NP_010079.1	4.02	1	1	0	0	0	0	0	0	0	0	0	4.02
6325336	MRL1	NP_015404.1	3.41	1	1	0	0	0	0	0	0	0	0	0	2.62
398365551	SGM1	NP_012668.3	1.84	1	1	0	0	0	0	0	0	0	0	0	2.12
6322985	FPS1	NP_013057.1	2.24	1	3	0	0	0	0	0	0	0	0	0	0
398365713	MRM1	NP_014844.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323001	ISA1	NP_013073.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323955	PRC1	NP_014026.1	0	0	0	0	0	0	0	0	0	0	0	0	1.88
6322095	YIL096C	NP_012170.1	0	0	0	0	0	0	0	0	0	0	0	0	1.79
6325139	MRP51	NP_015207.1	0	0	0	0	0	0	0	0	0	0	0	0	3.49
6323808	MRPS8	NP_013879.1	0	0	0	0	0	0	0	0	0	0	0	0	6.45
6322682	TPK3	NP_012755.1	0	0	0	0	0	0	0	0	0	0	0	0	2.51
6324978	YPL277C	NP_015046.1	0	0	0	0	0	0	0	0	0	2.05	1	2	0
6325296	ERV2	NP_015362.1	6.63	1	1	0	0	0	0	0	0	6.63	1	1	0
6323252	IFH1	NP_013324.1	0	0	0	1.11	1	1	0	0	0	0	0	0	0

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Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6324988	ACM1	NP_015056.1	0	0	0	0	0	0	0	0	0	0	0	0	0
330443417	RRN6	NP_009539.2	0	0	0	0	0	0	0	0	0	0	0	0	0
6325398	TAZ1	NP_015466.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321828	MSC7	NP_011904.1	4.66	3	4	0	0	0	0	0	0	0	0	0	0
6323320	GUF1	NP_013392.1	0	0	0	0	0	0	0	0	0	0	0	0	0
16740526	GYP8	NP_444296.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324783	NPT1	NP_014852.1	5.13	2	2	0	0	0	0	0	0	0	0	0	0
6319974	HOP	NP_010054.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319931	ABP1	NP_010012.1	24.66	13	34	0	0	0	2.20	1	1	0	0	0	14.02
6321787	OSH7	NP_011863.1	11.21	6	11	0	0	0	2.97	1	1	3.66	2	9	16.02
6319512	CHS2	NP_009594.1	0.83	1	1	0	0	0	0	0	0	0	0	0	2.18
6324912	KRE5	NP_014981.1	0	0	0	0	0	0	0	0	0	0	0	0	1.32
398365337	SFL1	NP_014783.4	0	0	0	0	0	0	0	0	0	0	0	0	0
398364433	CAJ1	NP_010967.3	14.32	4	6	3.58	1	4	0	0	0	0	0	0	23.02
6322547	EMC2	NP_012621.1	3.77	1	1	0	0	0	0	0	0	0	0	0	0
6324225	LEU4	NP_014295.1	3.07	2	2	0	0	0	0	0	0	0	0	0	2.42
6322537	MIR1	NP_012611.1	14.47	4	7	0	0	0	0	0	0	0	0	0	9.65
6321795	ERG11	NP_011871.1	18.87	7	12	0	0	0	0	0	0	0	0	0	10.38
6323953	YMR295C	NP_014024.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366013	YDR186C	NP_010472.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323943	DSS1	NP_014014.1	0	0	0	0	0	0	0	0	0	0	0	0	1.14
6320851	HEM14	NP_010930.1	2.23	1	2	0	0	0	0	0	0	0	0	0	0
6322116	SER33	NP_012191.1	16.63	6	15	0	0	0	2.13	1	1	2.35	1	1	17.70
83578105	TMA46	NP_014734.2	2.90	1	1	0	0	0	0	0	0	0	0	0	0
6321960	CDC23	NP_012036.1	0	0	0	0	0	0	0	0	0	0	0	0	2.24
6320470	AKR1	NP_010550.1	0	0	0	0	0	0	0	0	0	0	0	0	1.57
398365797	YNR048W	NP_014446.3	0	0	0	0	0	0	0	0	0	0	0	0	3.05
6322387	SMC3	NP_012461.1	0	0	0	1.22	1	1	0	0	0	0	0	0	0
6321806	ARG4	NP_011882.1	1.94	1	1	0	0	0	0	0	0	0	0	0	0
6324055	GOR1	NP_014125.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324554	YOL019W	NP_014623.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324897	PMT3	NP_014966.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323553	ALO1	NP_013624.1	19.58	8	22	0	0	0	0	0	0	0	0	0	15.78
6323958	ADE4	NP_014029.1	7.45	3	12	0	0	0	0	0	0	0	0	0	4.12
6322585	ENT3	NP_012659.1	7.84	3	48	0	0	0	2.70	1	1	0	0	0	26.47
6322732	SBA1	NP_012805.1	16.20	3	6	4.17	1	1	0	0	0	0	0	0	0
6319767	BSD2	NP_009849.1	0	0	0	0	0	0	4.05	1	1	0	0	0	4.05

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6323630	ERV25	NP_013701.1	6.16	1	2	0	0	0	0	0	0	0	0	0	0
6323716	TVP18	NP_013787.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319289	SNC1	NP_009372.1	11.11	1	3	0	0	0	0	0	0	0	0	0	0
398364281	EMP24	NP_011315.3	12.81	3	10	0	0	0	4.43	1	2	0	0	0	6.90
6320298	DNF2	NP_010378.1	2.23	3	3	0	0	0	0	0	0	0	0	0	2.30
330443597	MNN5	NP_012349.2	7.85	4	7	0	0	0	2.39	1	1	0	0	0	6.48
6320897	FCY2	NP_010976.1	2.44	2	2	0	0	0	0	0	0	0	0	0	1.69
330443542	SAP155	NP_116698.3	0	0	0	0	0	0	0	0	0	0	0	0	2.99
14318521	SPB4	NP_116654.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323367	NUP2	NP_013439.1	13.19	7	9	0	0	0	0	0	0	0	0	0	0
6319303	TPD3	NP_009386.1	9.29	4	4	0	0	0	0	0	0	0	0	0	4.57
6324166	RIA1	NP_014236.1	0.72	1	1	0	0	0	0	0	0	0	0	0	0.99
398364557	AIM21	NP_012268.3	0	0	0	0	0	0	1.18	1	1	0	0	0	0
6322369	SRS2	NP_012443.1	0.68	1	1	0	0	0	0	0	0	0	0	0	0.68
6324088	ZWF1	NP_014158.1	0	0	0	0	0	0	0	0	0	0	0	0	3.37
6319890	TAF2	NP_009971.1	0	0	0	0	0	0	0	0	0	0	0	0	0.57
6320033	SFA1	NP_010113.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325429	BSP1	NP_015497.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323022	YLL007C	NP_013094.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324918	YOR342C	NP_014987.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320091	TRM3	NP_010171.1	0.63	1	1	0	0	0	0	0	0	0	0	0	0.63
6322925	SIS2	NP_012998.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321999	SCH9	NP_012075.1	1.21	1	1	0	0	0	1.82	1	4	1.82	1	2	0
6320106	RPN6	NP_010186.1	2.07	1	1	0	0	0	0	0	0	0	0	0	25.58
6320934	PTC2	NP_011013.1	18.32	8	31	0	0	0	0	0	0	0	0	0	10.13
6322545	STE18	NP_012619.1	15.45	1	1	0	0	0	0	0	0	0	0	0	15.45
27808714	LAP3	NP_014160.2	0	0	0	0	0	0	0	0	0	0	0	0	3.08
6322410	IRC8	NP_012484.1	0	0	0	0	0	0	0	0	0	0	0	0	0
14318532	UBP6	NP_116665.1	13.23	5	7	0	0	0	5.41	2	3	0	0	0	17.03
6320129	BRE1	NP_010209.1	1.29	1	2	0	0	0	0	0	0	0	0	0	0
6324287	COG6	NP_014357.1	1.67	1	1	0	0	0	0	0	0	0	0	0	0
6320846	NTF2	NP_010925.1	6.40	1	2	0	0	0	0	0	0	0	0	0	12.80
6324508	MET22	NP_014577.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319975	GCS1	NP_010055.1	44.32	13	69	0	0	0	0	0	0	0	0	0	53.69
63535562	CHA1	NP_00101803	4.44	2	2	0	0	0	0	0	0	0	0	0	6.67
6322141	NEO1	NP_012216.1	2.00	2	2	0	0	0	0	0	0	0	0	0	1.56
398365175	CDC28	NP_009718.3	13.42	4	5	0	0	0	4.70	1	1	9.06	1	2	11.07

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6324946	MRS6	NP_015015.1	6.47	3	4	0	0	0	0	0	0	0	0	0	2.32
6325052	HRR25	NP_015120.1	3.85	1	2	0	0	0	0	0	0	0	0	0	1.82
6324631	SGT1	NP_014700.1	2.53	1	1	0	0	0	0	0	0	0	0	0	2.78
6323651	ADI1	NP_013722.1	0	0	0	0	0	0	0	0	0	0	0	0	7.26
398365629	DBF2	NP_011606.3	3.50	2	2	0	0	0	0	0	0	0	0	0	0
6320725	GRH1	NP_010805.1	4.03	1	1	0	0	0	0	0	0	0	0	0	0
6324056	TOF1	NP_014126.1	0	0	0	0	0	0	0	0	0	0	0	0	1.05
6321823	PIH1	NP_011899.1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383767	KCC4	NP_009907.2	1.64	1	1	0	0	0	0	0	0	0	0	0	0
6319556	UBC4	NP_009638.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325007	GYP5	NP_015075.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322121	SEC6	NP_012196.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323292	TMA7	NP_013364.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325227	TRM44	NP_015295.1	2.29	1	1	0	0	0	0	0	0	0	0	0	8.11
6322526	TOR1	NP_012600.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324085	SUI1	NP_014155.1	40.74	4	15	0	0	0	0	0	0	0	0	0	70.37
398366199	IVY1	NP_010515.3	2.43	1	2	0	0	0	0	0	0	0	0	0	6.62
6322839	ARC19	NP_012912.1	9.36	1	1	0	0	0	0	0	0	0	0	0	14.04
6321914	CIA2	NP_011990.1	0	0	0	0	0	0	0	0	0	0	0	0	5.19
6319737	RGD1	NP_009819.1	1.50	1	1	0	0	0	0	0	0	0	0	0	0
6322419	MHP1	NP_012493.1	0	0	0	0	0	0	0	0	0	0	0	0	0.50
398366245	LDB19	NP_014967.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325011	YPL245W	NP_015079.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364697	LSM4	NP_011037.3	4.81	1	1	5.88	1	5	0	0	0	0	0	0	10.70
6320840	FMP52	NP_010919.1	0	0	0	0	0	0	0	0	0	0	0	0	18.61
6324794	RCN2	NP_014863.1	4.91	1	1	0	0	0	0	0	0	0	0	0	4.53
6321800	VPS29	NP_011876.1	4.26	1	1	0	0	0	0	0	0	0	0	0	8.16
398364967	SOK1	NP_010289.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6320413	UME6	NP_010493.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322379	IML2	NP_012453.1	0	0	0	0	0	0	0	0	0	0	0	0	1.23
6325362	COG4	NP_015430.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322156	BCY1	NP_012231.1	8.65	4	8	0	0	0	0	0	0	0	0	0	11.06
6322424	SNX4	NP_012498.1	2.13	1	1	0	0	0	0	0	0	0	0	0	3.78
398365471	MCM7	NP_009761.4	1.07	1	3	0	0	0	0	0	0	0	0	0	1.54
398365269	SMY2	NP_009731.4	2.43	1	1	0	0	0	0	0	0	0	0	0	0
6323616	YML6	NP_013687.1	0	0	0	0	0	0	0	0	0	0	0	0	2.45
6325343	SUA7	NP_015411.1	0	0	0	6.96	1	1	3.77	1	1	0	0	0	8.70

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Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
6324068	ORC5	NP_014138.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325213	NOP4	NP_015282.1	0	0	0	0	0	0	0	0	0	0	0	0	2.48
6324041	CAF40	NP_014111.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320086	CYK3	NP_010166.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321762	SNF6	NP_011838.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365087	GRX4	NP_011101.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323218	ATG26	NP_013290.1	0	0	0	0	0	0	0	0	0	0	0	0	0
41629693	PRP18	NP_011520.2	0	0	0	0	0	0	0	0	0	0	0	0	0
330443528	PTP3	NP_010998.2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364733	YIR035C	NP_012301.3	3.54	1	1	0	0	0	0	0	0	0	0	0	3.54
6321766	AIM17	NP_011842.1	0	0	0	0	0	0	0	0	0	0	0	0	0
14318483	MOB2	NP_116618.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324527	PSK2	NP_014597.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365421	PRP16	NP_013012.3	0	0	0	0	0	0	0	0	0	0	0	0	0
41629685	RSC30	NP_011923.2	1.70	1	1	0	0	0	0	0	0	0	0	0	0
6320907	VHR2	NP_010986.1	0	0	0	0	0	0	0	0	0	0	0	0	2.97
398366405	GCN2	NP_010569.3	0	0	0	0	0	0	0	0	0	0	0	0	0
398364549	GYP6	NP_012491.3	0	0	0	0	0	0	0	0	0	0	0	0	0
330443696	YNL193W	NP_014206.2	0	0	0	0	0	0	0	0	0	0	0	0	0
6325243	YPL014W	NP_015311.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325327	MED1	NP_015395.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320160	MTF2	NP_010240.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319287	PRP45	NP_009370.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365919	SEC5	NP_010450.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6320433	SIR4	NP_010513.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321725	BIO2	NP_011802.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324177	INN1	NP_014247.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319534	ORC2	NP_009616.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320887	SAP1	NP_010966.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321204	SEC15	NP_011281.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323111	SRL2	NP_013183.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366601	SSN2	NP_010731.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323541	VPS9	NP_013612.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321955	YAP1801	NP_012031.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323215	EMG1	NP_013287.1	8.73	3	3	0	0	0	0	0	0	0	0	0	15.08
6319868	HTL1	NP_009949.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321537	MDR1	NP_011614.1	0	0	0	0	0	0	1.68	1	1	0	0	0	0.84

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Accession	Acronym	NP_value	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC
			BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_1	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2	BY4741_TAP_2
398364647	SLM4	NP_009633.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6324450	SMF1	NP_014519.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320105	SNU23	NP_010185.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366077	SPC19	NP_010487.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323175	SPE4	NP_013247.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322832	SPT23	NP_012905.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366615	TFB3	NP_010748.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6320032	UGX2	NP_010112.1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323960	YME2	NP_014031.1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365297	YOS9	NP_010342.3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323821	ALD3	NP_013892.1	0	0	0	0	0	0	0	0	0	0	0	0	1.38
6325392	MSS18	NP_015460.1	0	0	0	0	0	0	0	0	0	3.36	1	1	0
6321895	SBE22	NP_011971.1	0	0	0	0	0	0	0	0	0	0.94	1	1	0
6323844	SGS1	NP_013915.1	0	0	0	0	0	0	0	0	0	1.04	1	1	0
6321230	SIP2	NP_011307.1	0	0	0	0	0	0	0	0	0	2.41	1	1	0
6319783	VBA3	NP_009864.1	0	0	0	0	0	0	0	0	0	6.77	1	1	0
6320295	YDR090C	NP_010375.1	0	0	0	0	0	0	0	0	0	9.03	1	1	0
6320812	YEL023C	NP_010891.1	0	0	0	0	0	0	0	0	0	2.05	1	1	0
330443522	YEN1	NP_010959.2	0	0	0	0	0	0	0	0	0	1.19	1	1	0
6323731	YMR085W	NP_013802.1	0	0	0	0	0	0	0	0	0	3.70	1	1	0
6323761	YMR114C	NP_013832.1	0	0	0	0	0	0	0	0	0	4.08	1	1	0
6323599	CAT2	NP_013670.1	0	0	0	0	0	0	0	0	0	1.79	1	1	0
6320221	YDR018C	NP_010301.1	0	0	0	0	0	0	0	0	0	3.54	1	1	0
6324392	YNR064C	NP_014462.1	0	0	0	0	0	0	0	0	0	3.45	1	1	0
6324597	AHC1	NP_014666.1	0	0	0	0	0	0	0	0	0	2.65	1	1	0
6323774	DLT1	NP_013845.1	0	0	0	0	0	0	0	0	0	2.92	1	1	0
6323532	MDM1	NP_013603.1	0	0	0	0	0	0	0	0	0	0.98	1	1	0
6323098	MEF1	NP_013170.1	0	0	0	0	0	0	0	0	0	2.50	1	1	0
6322467	MHO1	NP_012541.1	0	0	0	0	0	0	0	0	0	8.28	1	1	0
6322177	PDR11	NP_012252.1	0	0	0	0	0	0	0.64	1	1	0	0	0	0
6323099	XYL2	NP_013171.1	0	0	0	0	0	0	2.81	1	1	0	0	0	0
14318495	EPL1	NP_116629.1	0	0	0	0	0	0	0.96	1	1	0	0	0	0
398364429	FLC3	NP_011376.3	0	0	0	0	0	0	1.37	1	1	0	0	0	0
6320193	GRX6	NP_010274.1	0	0	0	0	0	0	5.63	1	1	0	0	0	0
398364915	HUL4	NP_012570.3	0	0	0	0	0	0	1.35	1	1	0	0	0	0
6325457	ARR1	NP_015525.1	0	0	0	4.76	1	1	0	0	0	0	0	0	0
6322660	PXA2	NP_012733.1	0	0	0	1.64	1	1	0	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398364847	RTR1	2	3	8.85	2	3	0	0	0	19.47	7	661	69.03	18
6320061	RPB1	7	9	3.98	7	9	2.19	3	4	9.87	20	188	33.81	50
398365417	RPB2	3	3	3.02	3	3	0.74	1	1	7.84	10	298	36.60	38
398364469	RPB3	1	1	5.35	1	1	0	0	0	9.43	3	63	38.68	8
6322321	RPB4	0	0	0	0	0	13.12	2	4	0	0	0	21.72	3
6319630	RPB5	2	2	13.02	2	2	0	0	0	3.72	1	4	41.86	6
398366567	RPB7	0	0	0	0	0	0	0	0	0	0	0	0	0
398365775	RPB10	0	0	0	0	0	15.71	1	2	0	0	0	15.71	1
6325445	RPB6	0	0	0	0	0	0	0	0	0	0	0	26.45	2
6321368	RPB9	0	0	0	0	0	0	0	0	0	0	0	20.49	2
6324798	RPB8	0	0	0	0	0	0	0	0	6.85	1	21	25.34	3
6324569	RPB11	0	0	0	0	0	0	0	0	13.33	2	23	39.17	4
6323632	SPT5	8	10	8.94	8	10	4.33	3	3	1.22	1	8	0	0
6321625	TFG1	1	2	1.36	1	2	0	0	0	1.50	1	1	0	0
6321937	RPB12	0	0	0	0	0	28.57	1	7	10.00	1	30	40.00	3
398365043	TFG2	1	1	2.00	1	1	2.75	1	1	0	0	0	0	0
154199603	IWR1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365457	SPT4	0	0	0	0	0	0	0	0	0	0	0	0	0
6325128	TAF14	1	1	6.56	1	1	6.56	1	4	0	0	0	0	0
398365155	NPA3	3	4	8.83	3	4	6.23	2	2	9.09	3	17	14.81	5
6323272	GPN3	0	0	0	0	0	4.78	1	1	0	0	0	12.87	2
6322163	IRR1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366099	RRP41	0	0	0	0	0	0	0	0	0	0	0	0	0
398366555	UBA2	1	1	2.20	1	1	7.23	4	4	0	0	0	0	0
6323670	TAP42	3	5	7.65	3	5	3.55	1	2	0	0	0	0	0
6323004	SSA2	29	541	56.65	29	541	80.13	48	1842	13.62	8	69	51.96	22
6319505	RPL4A	12	120	37.02	12	120	50.00	15	302	9.12	4	599	24.59	6
398364991	RPL4B	0	0	32.60	10	117	50.00	14	286	9.12	4	599	20.72	5
398365303	RPL43B	5	83	43.48	5	83	39.13	5	340	9.78	2	238	44.57	5
398364349	RPL2A	4	79	27.17	4	79	48.03	11	239	20.08	7	655	18.50	5
6325337	TEF1	18	438	44.10	18	438	41.05	16	226	13.54	8	221	25.55	7
144228166	SSA1	31	516	57.79	31	516	76.64	46	1665	17.13	10	75	50.31	25
6319559	RPL19A	6	75	29.63	6	75	31.22	8	293	16.40	6	130	20.63	4
6320065	RPL35A	5	27	30.83	5	27	40.83	9	135	0	0	0	37.50	6
398365603	RPS4A	8	32	32.57	8	32	48.66	14	81	16.09	6	487	23.75	6
398365321	RPS6B	7	38	33.90	7	38	56.36	18	190	21.61	5	161	34.32	10
398364655	SSA4	15	134	28.19	15	134	21.18	18	510	5.76	4	9	15.42	8

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

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		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1_ _WT_1	TAP_Rtr1_ _WT_1
398366603	RPS17B	6	47	52.94	6	47	54.41	8	158	7.35	1	4	55.15	10
6324533	RPS15	3	29	30.28	3	29	30.28	3	48	16.90	2	60	38.03	4
6323104	RPL10	7	118	27.60	7	118	38.91	10	103	33.03	12	421	15.38	3
6319396	SSA3	11	117	19.72	11	117	22.65	18	436	5.70	4	39	13.56	7
6321786	RPL14B	0	0	39.13	6	52	44.93	8	152	12.32	4	174	18.84	3
6322847	RPL14A	7	53	45.65	7	53	51.45	9	159	12.32	4	174	18.84	3
6324670	RPS7A	6	61	31.05	6	61	39.47	10	224	15.79	4	51	53.16	9
6322668	RPL17A	6	82	36.41	6	82	34.78	5	328	20.11	3	56	28.26	4
398365285	CAX4	0	0	0	0	0	0	0	0	0	0	0	0	0
6322284	RPL17B	6	85	36.41	6	85	34.78	5	306	20.11	3	56	28.26	4
398364883	RPS7B	7	33	40.00	7	33	57.89	12	213	15.79	4	51	43.68	9
6323791	RPL13B	0	0	37.69	6	25	44.72	10	113	26.63	9	1125	39.20	7
6320121	RPL13A	6	25	37.69	6	25	44.72	10	104	26.63	9	1125	37.69	6
6324120	SSB2	25	473	58.73	25	473	66.56	39	906	9.30	8	111	38.34	15
6323577	RPS1B	8	46	35.29	8	46	55.69	13	199	16.47	5	87	38.82	8
6325157	ATG21	0	0	0	0	0	0	0	0	0	0	0	0	0
6323474	RPS1A	6	41	27.45	6	41	55.69	12	179	16.47	5	87	38.82	8
398364725	RPL23B	6	114	43.80	6	114	46.72	8	216	5.11	1	1	32.85	4
6319972	SSB1	25	472	58.73	25	472	66.39	38	900	10.60	9	112	38.34	15
154200011	RPL20B	4	47	30.23	4	47	39.53	8	156	23.84	6	773	29.65	5
6322989	UBI4	2	8	38.16	2	8	50.00	3	11	17.11	1	1	50.00	3
9755341	RPS16A	4	24	32.17	4	24	58.04	8	58	31.47	6	80	49.65	7
6319378	RPL32	3	35	23.85	3	35	36.92	5	169	35.38	6	212	17.69	2
6325061	APL5	0	0	0	0	0	1.61	1	1	0	0	0	0	0
398364505	RPS3	8	40	39.17	8	40	52.92	11	51	25.42	5	74	47.08	10
398365351	RPS13	6	44	45.70	6	44	58.28	9	206	7.28	1	12	45.70	6
6324260	RPL16B	6	28	27.27	6	28	32.83	10	115	22.73	7	48	6.06	1
398365605	RPS30B	1	3	15.87	1	3	30.16	2	6	0	0	0	17.46	2
6323481	RPL6B	6	14	37.50	6	14	53.98	9	84	30.68	8	237	4.55	1
398365377	RPS9B	7	18	30.26	7	18	43.08	12	44	21.03	5	79	23.08	4
6324265	YDJ1	8	77	27.87	8	77	25.92	11	47	5.13	2	13	6.60	2
398364745	ATP7	1	1	7.47	1	1	7.47	1	1	0	0	0	8.05	2
6321464	RPS25A	5	13	38.89	5	13	39.81	6	56	27.78	3	60	31.48	5
6323077	RPS0B	5	37	29.37	5	37	29.37	5	45	9.92	3	135	28.97	4
398365071	RPS11A	4	6	25.64	4	6	42.95	7	56	19.23	3	123	18.59	3
6321653	RPS0A	5	37	29.37	5	37	19.84	4	44	9.92	3	135	28.97	4
6323465	CNA1	3	3	7.23	3	3	2.35	1	1	0	0	0	26.40	11

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6321335	RPL28	2	22	16.11	2	22	35.57	7	67	16.78	3	178	22.15	3
6322505	SSC1	25	166	45.87	25	166	55.05	32	338	5.81	4	7	14.53	6
6323567	RPL6A	5	13	25.57	5	13	48.86	9	72	31.82	9	215	4.55	1
6321754	RPL8A	11	91	42.19	11	91	56.25	19	186	27.73	11	405	32.81	8
6322984	RPL8B	13	93	49.22	13	93	57.03	20	189	27.73	11	513	36.33	9
6323615	RPS18B	6	61	43.15	6	61	59.59	9	114	13.70	4	67	43.15	5
398366307	RPA190	12	16	8.71	12	16	4.21	5	7	3.61	6	25	0	0
6324637	RPL3	11	43	27.13	11	43	42.38	16	245	30.49	15	789	0	0
6320128	RPL31A	4	28	38.94	4	28	61.95	8	85	31.86	5	399	30.09	3
398366143	RPL31B	0	0	30.97	3	27	53.98	7	78	31.86	5	354	30.09	3
398365761	RPS23A	3	27	33.79	3	27	34.48	4	24	19.31	3	18	18.62	1
6322137	RPL34B	1	1	6.61	1	1	6.61	1	14	6.61	1	20	6.61	1
6321315	RPS2	2	20	11.02	2	20	35.43	7	67	9.45	3	30	6.69	1
6320949	RPS8B	5	55	35.50	5	55	34.50	6	62	12.00	3	5	34.50	5
6321798	RPL27A	0	0	37.50	6	26	42.65	7	137	14.71	3	475	15.44	2
6325176	RPS9A	5	12	21.83	5	12	43.15	11	34	12.69	4	75	14.72	3
86558907	RHR2	7	12	32.80	7	12	26.40	6	22	3.60	1	14	26.40	6
6322058	RPL16A	7	23	31.66	7	23	37.69	12	100	31.16	10	97	0	0
6324027	RPS19B	5	45	45.14	5	45	60.42	10	89	7.64	2	12	53.47	9
6323090	RPL22A	3	12	55.37	3	12	61.16	4	40	0	0	0	61.16	4
6320142	RPS29B	0	0	0	0	0	0	0	0	0	0	0	32.14	2
6324807	KIN4	0	0	0	0	0	0	0	0	0	0	0	0	0
6321522	RPL11B	6	26	31.03	6	26	36.78	7	42	11.49	2	41	25.86	5
6322583	RPS5	6	149	29.78	6	149	30.22	7	86	13.78	3	10	34.22	8
6325126	RPL5	5	58	19.19	5	58	45.79	9	88	20.20	7	162	3.03	1
6323073	PDC1	19	234	43.69	19	234	49.20	21	433	8.88	6	441	22.56	8
398366277	VMA4	13	247	53.22	13	247	54.51	16	199	17.60	4	31	21.89	5
6323824	HOT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325066	NAB3	0	0	0	0	0	2.99	2	2	0	0	0	0	0
398364517	RPS24A	5	58	31.85	5	58	48.89	8	120	20.00	3	30	40.00	6
6321362	RPL7A	9	58	43.44	9	58	49.18	12	128	28.69	8	621	18.03	3
6325058	RPL7B	9	49	43.44	9	49	49.18	12	124	28.69	8	628	18.03	3
37362649	GUS1	20	96	33.76	20	96	22.18	13	90	8.76	7	61	2.12	1
398365909	RPL24B	2	18	15.48	2	18	27.10	6	140	12.90	2	4	9.03	1
6324707	EFT1	32	590	48.93	32	590	56.41	37	629	13.90	13	486	14.61	8
6323376	RPL26A	0	0	34.65	6	13	51.18	10	59	22.05	6	177	44.09	8
116006491	RPL26B	7	18	35.43	7	18	51.18	11	61	22.05	6	186	44.09	8

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		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6325006	RPL36B	4	23	40.00	4	23	49.00	6	88	34.00	5	120	33.00	4
6323848	RPL36A	0	0	31.00	3	15	50.00	6	79	34.00	4	70	33.00	4
398366579	RPL12B	6	95	49.70	6	95	56.36	7	139	4.85	1	1	33.33	4
6321540	NOP7	1	1	1.49	1	1	4.30	2	5	0	0	0	0	0
398365599	PDC6	3	22	6.93	3	22	6.93	3	93	1.60	1	10	3.37	1
6323163	PDC5	2	14	5.15	2	14	9.59	3	77	1.78	1	18	3.37	1
6323583	CMP2	4	4	7.78	4	4	4.97	2	3	0	0	0	11.42	5
398365991	CLC1	1	2	6.87	1	2	6.87	1	2	0	0	0	15.45	3
37362619	RXT2	2	2	5.12	2	2	0	0	0	0	0	0	0	0
10383793	RPS14A	5	37	40.15	5	37	65.69	10	135	34.31	6	105	43.80	7
6322271	RPS22A	3	52	24.62	3	52	24.62	3	24	13.08	2	33	25.38	3
6321407	RPL24A	4	23	22.58	4	23	27.74	7	163	10.97	2	4	16.77	2
6323357	RPL38	2	14	32.05	2	14	50.00	4	67	0	0	0	33.33	3
398366183	HTB1	6	56	39.69	6	56	31.30	5	77	8.40	1	1	18.32	2
398364287	GCN1	20	35	8.16	20	35	2.10	5	6	2.96	7	61	0	0
6320016	VMA1	33	228	33.43	33	228	28.38	26	154	9.62	10	244	2.80	2
14318481	TUB2	10	72	27.79	10	72	35.01	11	62	4.38	3	99	8.10	3
398364585	KAR2	20	48	32.40	20	48	31.82	21	127	1.61	1	1	3.96	2
6323196	RPS31	1	2	10.53	1	2	32.89	2	11	0	0	0	0	0
6321013	PAB1	16	60	35.18	16	60	49.74	22	128	7.45	5	24	28.08	13
6324993	FUM1	1	2	2.66	1	2	5.33	2	2	0	0	0	0	0
398366075	RPS29A	1	2	14.29	1	2	14.29	1	3	0	0	0	32.14	2
6322755	YJU2	2	2	11.15	2	2	0	0	0	0	0	0	0	0
398366059	PLP2	2	5	7.34	2	5	0	0	0	0	0	0	0	0
6320718	SMT3	1	1	6.93	1	1	6.93	1	1	0	0	0	0	0
398365031	YBR139W	0	0	0	0	0	0	0	0	0	0	0	3.35	1
6321850	FYV4	0	0	0	0	0	0	0	0	0	0	0	0	0
398366083	TDH3	14	416	49.70	14	416	63.86	15	408	24.40	10	401	46.08	12
6319279	CDC19	21	122	55.60	21	122	58.60	25	144	18.20	10	410	31.60	11
6322468	TDH2	13	389	47.59	13	389	50.30	13	399	21.99	9	397	45.78	11
6321249	RPS26A	0	0	26.05	3	11	26.05	3	37	13.45	2	68	20.17	2
6321772	RPS20	6	16	45.45	6	16	33.88	6	25	27.27	5	22	9.92	1
14318474	SEC53	7	20	31.10	7	20	24.02	5	14	5.51	1	12	5.91	1
6320589	YRA1	4	17	20.80	4	17	19.91	4	9	3.54	1	1	19.91	4
6320805	ECM10	0	0	12.27	9	64	12.27	9	74	4.19	3	4	2.80	1
6323776	ECM16	0	0	0	0	0	0	0	0	0	0	0	0	0
6320596	RVS167	2	3	4.98	2	3	0	0	0	2.07	1	6	0	0

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		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398366359	SLH1	1	2	0.46	1	2	0	0	0	0	0	0	0	0
6324321	SIS1	6	9	26.14	6	9	24.72	6	16	4.55	1	1	10.80	2
6319447	NCL1	9	11	16.52	9	11	10.23	5	17	4.24	2	16	0	0
6323112	EMP70	1	1	2.40	1	1	4.05	2	2	0	0	0	0	0
6324595	SFM1	0	0	0	0	0	16.90	3	4	0	0	0	0	0
330443411	UBP13	1	1	1.87	1	1	0	0	0	0	0	0	0	0
6323868	EFR3	0	0	0	0	0	0	0	0	0	0	0	0	0
330443676	CDC34	3	5	10.58	3	5	6.92	3	3	1.92	1	1	0	0
6320615	TRS120	1	1	0.93	1	1	0	0	0	0	0	0	0	0
6323487	FMP27	1	1	0.34	1	1	0	0	0	0	0	0	0	0
6320231	REG1	0	0	0	0	0	4.64	3	4	0.89	1	8	0	0
6324585	AUS1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325048	RKM1	0	0	0	0	0	7.03	3	6	0	0	0	0	0
6323732	SEG1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364779	YGL041W-	0	0	0	0	0	0	0	0	0	0	0	13.64	1
398365827	CAF130	0	0	0	0	0	0	0	0	0	0	0	0	0
6325063	RSA1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365559	MSH6	0	0	1.05	1	1	0	0	0	1.13	1	2	0	0
398365727	YNR034W-	3	10	40.82	3	10	0	0	0	0	0	0	0	0
6323683	YET2	0	0	0	0	0	0	0	0	0	0	0	0	0
330443366	GPB2	0	0	0	0	0	0	0	0	0	0	0	0	0
6325012	HUT1	0	0	0	0	0	0	0	0	0	0	0	0	0
99030927	KKQ8	0	0	0	0	0	0	0	0	0	0	0	0	0
398364567	MAD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320763	RMD6	0	0	0	0	0	0	0	0	0	0	0	0	0
398365009	MEC1	56	325	28.25	56	325	23.48	49	148	0	0	0	0	0
398364407	RPL9A	7	109	48.69	7	109	51.31	8	248	11.52	3	257	25.13	4
6324298	HHF2	4	30	40.78	4	30	41.75	5	28	0	0	0	29.13	3
6324262	RPL9B	7	98	48.69	7	98	50.26	7	177	16.23	4	260	15.18	3
398364523	TDH1	8	312	28.92	8	312	26.51	6	212	12.65	5	270	35.24	9
398364951	VMA2	26	348	60.54	26	348	61.32	26	192	10.44	6	281	28.43	10
6321333	ARC1	6	17	17.29	6	17	14.89	6	21	7.98	3	10	0	0
6321408	RPL30	3	80	38.10	3	80	67.62	7	108	8.57	1	5	39.05	3
6324445	RPL25	5	22	35.21	5	22	45.07	8	116	16.90	2	16	19.01	2
6325114	RPL33A	2	11	24.30	2	11	32.71	4	30	15.89	2	82	0	0
398365881	RPL33B	2	9	24.30	2	9	31.78	3	14	15.89	2	82	0	0
398364339	THS1	20	43	31.74	20	43	29.02	17	58	2.45	2	12	6.13	3

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
330443609	URA2	47	95	26.65	47	95	16.76	30	81	5.92	13	143	5.19	8
6323209	SAM1	7	20	17.80	7	20	55.24	16	123	5.76	3	95	0	0
398364415	HXK1	5	11	13.81	5	11	5.98	3	4	3.51	2	3	7.22	2
6321232	CHC1	17	23	12.89	17	23	8.29	10	13	1.33	2	4	0.97	1
6322352	UTP10	3	3	2.09	3	3	0	0	0	0	0	0	0	0
398366655	SAM2	7	18	17.45	7	18	44.53	15	85	8.33	3	33	0	0
6322115	RPN2	13	20	15.03	13	20	6.67	6	7	3.07	3	14	1.59	1
6320073	RPP1B	1	13	15.09	1	13	47.17	2	139	0	0	0	15.09	1
6320122	RPP1A	2	8	28.30	2	8	58.49	3	7	0	0	0	0	0
6323693	ERB1	2	3	2.48	2	3	6.69	5	9	1.36	1	1	0	0
6320784	VMA8	0	0	0	0	0	8.59	2	2	12.50	4	32	8.59	2
6323087	SHM2	12	70	26.87	12	70	15.57	6	26	12.79	7	84	6.18	2
6320119	SUB2	12	37	29.60	12	37	31.61	13	24	11.21	4	29	7.62	2
6320573	ESF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323179	STM1	1	1	2.93	1	1	32.60	6	13	15.75	3	5	16.12	2
398366241	SEN1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323763	ASC1	5	9	18.18	5	9	20.69	6	14	4.70	1	1	3.76	1
6324772	BFR1	10	21	22.55	10	21	26.38	11	27	2.34	1	1	11.49	4
6320175	ARP2	5	12	17.14	5	12	16.37	4	9	2.81	1	3	0	0
398366669	EMI2	1	1	2.40	1	1	0	0	0	0	0	0	0	0
6323389	RSC2	3	4	3.60	3	4	0	0	0	1.35	1	1	0	0
6324663	VPS21	6	18	31.90	6	18	60.48	10	48	12.38	2	4	0	0
6319543	TAT1	1	1	1.94	1	1	0	0	0	0	0	0	0	0
6325417	KRE6	3	11	5.97	3	11	0	0	0	2.08	1	4	0	0
6324021	KRI1	2	2	4.06	2	2	0	0	0	0	0	0	0	0
6323454	YLR422W	0	0	0	0	0	0	0	0	0.47	1	2	0	0
6319756	PAF1	0	0	0	0	0	5.39	1	1	0	0	0	0	0
6323725	SEC14	13	101	53.62	13	101	60.53	14	127	0	0	0	0	0
6323368	SGD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322598	IML1	1	1	0.69	1	1	0	0	0	0	0	0	0	0
398366503	YDR341C	7	12	12.19	7	12	12.03	7	10	1.65	1	5	0	0
6320423	RAD9	0	0	0	0	0	0	0	0	0	0	0	0	0
6325147	GDE1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319585	CMD1	4	11	38.10	4	11	8.84	1	2	6.12	1	7	0	0
6322946	PTR2	2	4	4.16	2	4	0	0	0	0	0	0	0	0
398366323	GND2	0	0	4.47	2	2	2.03	1	2	0	0	0	2.03	1
6324231	RAS2	2	2	11.49	2	2	15.84	3	12	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6320637	TIF35	6	14	31.02	6	14	16.42	3	12	7.30	2	6	0	0
398364223	NPR2	1	1	2.28	1	1	0	0	0	0	0	0	0	0
6325390	SPN1	8	15	24.88	8	15	3.66	1	2	3.66	2	3	3.41	1
6324578	DNL4	0	0	0	0	0	0	0	0	0	0	0	0	0
6319291	FRT2	0	0	0	0	0	0	0	0	0	0	0	0	0
6322242	HXT9	0	0	0	0	0	0	0	0	0	0	0	0	0
6322272	RPL39	0	0	0	0	0	0	0	0	0	0	0	0	0
14318518	VTC2	3	4	3.38	3	4	0	0	0	1.09	1	5	0	0
330443752	THI22	0	0	3.15	2	8	1.75	1	1	1.75	1	7	0	0
6325175	MOT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322796	DEF1	0	0	0	0	0	0	0	0	0	0	0	8.67	2
398365059	PTK2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320345	DOP1	3	4	1.83	3	4	0	0	0	0	0	0	0	0
6323863	HFA1	0	0	0	0	0	0.42	1	2	0.52	1	18	0	0
6322948	MLP1	6	8	4.16	6	8	4.16	5	6	0	0	0	0	0
6322999	FRA1	1	1	1.74	1	1	1.47	1	1	1.07	1	1	0	0
6324491	IRA2	1	1	0.49	1	1	0.23	1	1	0.36	1	1	0	0
6324215	RPC19	0	0	0	0	0	10.56	1	1	8.45	1	5	10.56	1
6320419	UPC2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364627	RDH54	2	2	2.30	2	2	0	0	0	1.04	1	1	0	0
6321752	VMR1	0	0	0	0	0	0.50	1	1	0	0	0	0	0
6321793	TIM10	0	0	0	0	0	15.05	1	1	0	0	0	0	0
6322032	BNR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325074	RTT10	0	0	0	0	0	0	0	0	0	0	0	0	0
6325235	RAD1	0	0	0	0	0	0	0	0	1.27	1	1	0	0
6323869	CEF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323917	TPS3	1	2	1.61	1	2	1.04	1	1	0	0	0	0	0
6321552	SPT6	7	10	5.17	7	10	2.83	3	4	0.55	1	1	0	0
6320853	BIM1	0	0	0	0	0	0	0	0	4.94	1	2	0	0
398364243	YEL043W	1	3	1.15	1	3	0	0	0	0	0	0	0	0
6320027	DLD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323356	PEX30	1	2	3.44	1	2	0	0	0	1.53	1	1	0	0
6320799	ANP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324393	YNR065C	0	0	1.88	2	2	1.88	2	2	0	0	0	0	0
6323266	THI7	0	0	0	0	0	0	0	0	0	0	0	0	0
6323445	YLR413W	1	1	1.93	1	1	0	0	0	1.48	1	1	0	0
398364667	THO2	0	0	0	0	0	0.88	1	2	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1_ _WT_1	TAP_Rtr1_ _WT_1
37362637	JIP4	0	0	0	0	0	1.03	1	1	0	0	0	0	0
398364707	AAC3	0	0	7.49	2	6	3.58	1	2	0	0	0	3.58	1
6322570	YMR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324879	ISW2	0	0	0	0	0	0	0	0	0	0	0	0	0
10383772	LEU2	0	0	0	0	0	0	0	0	0	0	0	6.32	1
6322079	HOS4	0	0	0.65	1	1	0	0	0	0	0	0	0	0
6321210	VID30	1	1	1.46	1	1	0	0	0	0.84	1	1	0	0
398365701	YNR029C	1	1	2.80	1	1	0	0	0	0	0	0	0	0
6320692	VPS52	1	1	2.50	1	1	0	0	0	0	0	0	0	0
398365525	SMD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322086	SLM1	1	1	1.75	1	1	0	0	0	0	0	0	2.92	1
398364897	CBP6	4	9	21.60	4	9	27.78	6	8	0	0	0	6.79	1
6323726	NAM7	1	1	1.13	1	1	1.85	1	1	0	0	0	0	0
6324982	SAM4	1	1	3.38	1	1	0	0	0	0	0	0	0	0
6320324	VBA4	0	0	0	0	0	0	0	0	0	0	0	0	0
6324612	HIR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324981	SAM3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325379	AXL1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321020	BRR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324694	GCY1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322312	DAS1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364625	YKL033W-	2	2	8.47	2	2	0	0	0	3.39	1	1	0	0
6320034	NRP1	1	1	1.67	1	1	0	0	0	0	0	0	0	0
6325374	YPR117W	0	0	0	0	0	0	0	0	0	0	0	0	0
6324546	MDM38	0	0	0	0	0	0	0	0	1.75	1	1	3.14	1
6323289	LCB5	0	0	0	0	0	0	0	0	0	0	0	0	0
6321205	TAN1	0	0	0	0	0	6.57	1	2	0	0	0	0	0
6319351	PHO11	0	0	0	0	0	0	0	0	0	0	0	0	0
6319523	REB1	1	1	1.23	1	1	3.46	2	5	1.60	1	1	0	0
6320921	YER077C	1	1	1.02	1	1	0	0	0	0	0	0	0	0
6320826	GLC3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365141	INP53	1	1	0.99	1	1	0.72	1	1	0	0	0	0	0
398366583	SIP1	1	1	1.47	1	1	0	0	0	0	0	0	0	0
398364669	MAD3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322700	DBR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323317	YLR287C	0	0	0	0	0	0	0	0	0	0	0	0	0
398365819	SAN1	1	3	3.28	1	3	1.48	1	1	2.95	1	1	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398365335	YNL033W	0	0	0	0	0	0	0	0	0	0	0	0	0
33438893	YPL119C-	0	0	0	0	0	0	0	0	0	0	0	0	0
6322579	JHD2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324141	KAR1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383756	LRE1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364917	NHP10	0	0	0	0	0	0	0	0	0	0	0	0	0
6323035	NSE1	0	0	0	0	0	0	0	0	0	0	0	0	0
50593425	OMA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321910	ORC6	0	0	0	0	0	0	0	0	0	0	0	0	0
6322494	PET191	0	0	0	0	0	0	0	0	0	0	0	0	0
398365057	SPO14	0	0	0	0	0	0	0	0	0	0	0	0	0
398364971	TRP1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365003	PSF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321697	RAD2	0	0	0	0	0	0	0	0	0	0	0	0	0
6319525	REG2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320271	RTR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324162	SKO1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366633	SNM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322778	STB6	0	0	0	0	0	0	0	0	0	0	0	0	0
6325081	TRE1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324447	TRM13	0	0	0	0	0	0	0	0	0	0	0	0	0
6323236	HRD3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325184	UBP16	0	0	0	0	0	0	0	0	0	0	0	0	0
398364885	UBP5	0	0	0	0	0	0	0	0	0	0	0	0	0
6325071	UIP4	0	0	0	0	0	0	0	0	0	0	0	0	0
398365127	UIP5	0	0	0	0	0	0	0	0	0	0	0	0	0
6319904	YIH1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325266	SUT2	0	0	0	0	0	0	0	0	7.46	1	1	0	0
6324449	HRP1	1	1	3.18	1	1	7.12	3	8	1.50	1	3	2.43	1
6323278	YEF3	36	340	43.68	36	340	29.50	26	113	15.13	18	452	5.94	5
6322084	PFK26	28	347	37.85	28	347	28.66	18	54	1.21	1	2	0	0
6323057	RPL15A	5	18	25.49	5	18	36.27	7	179	15.20	4	413	6.86	1
6323769	RPL15B	5	12	25.49	5	12	31.37	6	162	15.20	4	347	6.86	1
6319282	FUN12	2	2	3.09	2	2	4.89	4	4	3.19	2	9	0	0
6320190	NOP1	9	25	35.17	9	25	6.42	2	7	11.31	5	16	4.89	1
6323654	CLU1	16	31	16.60	16	31	0.70	1	1	5.95	7	20	0	0
6324886	NOP58	8	17	20.74	8	17	21.33	8	21	8.22	3	11	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6323226	NOP56	9	15	23.21	9	15	22.22	10	14	4.37	2	18	0	0
37362658	SSZ1	13	54	35.50	13	54	36.06	15	41	7.06	4	51	7.43	2
6324452	RPL18A	4	49	24.19	4	49	27.96	5	98	30.65	6	380	6.99	1
398366653	LCD1	16	43	26.37	16	43	19.68	12	41	0	0	0	0	0
6322313	RPA34	7	45	37.34	7	45	43.35	11	215	0	0	0	0	0
398364437	RPL1B	9	85	38.25	9	85	38.71	10	111	7.37	2	35	0	0
6321968	ENO2	14	178	42.11	14	178	48.97	17	272	13.73	6	88	41.65	11
398365729	DED1	21	115	43.71	21	115	28.81	15	52	8.11	6	60	35.60	14
6323566	FPR3	2	6	6.81	2	6	6.81	2	2	5.11	2	4	0	0
398366315	ENO1	10	95	29.52	10	95	34.32	11	140	13.73	6	88	23.11	6
10383781	PGK1	16	45	48.08	16	45	45.67	16	47	3.85	2	30	38.22	10
6321874	KSP1	6	12	7.68	6	12	0	0	0	0	0	0	0	0
6321531	VAS1	31	74	33.42	31	74	23.73	22	57	3.89	4	16	0	0
6322769	TEF4	9	71	27.43	9	71	19.17	8	31	11.89	5	30	8.01	2
398366187	HTA1	1	4	6.82	1	4	12.12	2	12	5.30	1	18	12.12	2
330443451	GRS1	27	206	45.73	27	206	44.98	25	136	19.49	16	261	6.15	3
6321203	ADE5,7	12	31	19.58	12	31	20.07	11	14	4.24	4	16	2.24	1
6324486	ADH1	14	654	47.99	14	654	55.75	19	463	13.79	5	143	13.79	4
6323371	RPP0	9	100	33.33	9	100	31.73	8	113	8.65	3	114	3.53	1
6323886	RPS10B	1	1	17.14	1	1	38.10	3	51	20.95	3	5	0	0
6320009	ARF1	8	735	56.35	8	735	56.35	8	336	12.71	2	100	27.62	4
6321429	LEU1	23	320	35.82	23	320	39.41	22	122	2.82	2	19	0	0
6323204	CBF5	2	4	6.42	2	4	0	0	0	4.97	2	4	0	0
6320064	ARF2	8	326	56.35	8	326	56.35	8	129	12.71	2	100	14.36	2
6321265	XRN1	14	21	11.58	14	21	2.75	3	5	0	0	0	0	0
6323904	RKR1	24	63	19.53	24	63	1.02	1	1	0	0	0	0	0
6322790	FBA1	8	60	36.21	8	60	31.75	7	80	6.13	2	36	9.47	2
6324562	HTZ1	2	13	14.18	2	13	22.39	3	11	12.69	2	27	29.85	4
6321231	SPT16	5	10	5.60	5	10	4.93	4	5	0.97	1	1	0	0
398365455	TEF3	10	61	10.44	10	61	6.80	6	16	9.39	10	100	1.05	1
398366333	FPR4	0	0	4.34	1	2	9.18	3	5	0	0	0	0	0
398364607	DBP3	1	1	2.29	1	1	4.40	2	2	2.10	2	6	0	0
6324534	RPP2A	5	76	79.25	5	76	79.25	5	85	11.32	1	26	28.30	2
6321430	PMA1	16	71	20.26	16	71	24.51	20	84	10.89	11	303	1.63	1
398364809	DBP2	14	66	28.39	14	66	9.16	4	16	0	0	0	7.14	3
6319668	RPL21A	5	19	22.50	5	19	26.25	6	32	26.25	6	462	11.88	2
6323981	FET4	6	13	13.22	6	13	8.33	2	4	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6321807	DED81	15	54	33.57	15	54	36.82	16	65	9.75	5	11	2.17	1
398364767	SUI2	4	9	18.75	4	9	9.87	2	5	9.54	3	25	0	0
6323011	DPS1	16	55	35.01	16	55	42.37	21	70	7.18	4	41	6.28	3
6323885	RRP5	12	16	8.62	12	16	3.64	6	8	2.14	3	4	0.69	1
398366547	RPP2B	6	80	81.82	6	80	86.36	7	111	12.73	1	11	22.73	2
6323288	HSP60	22	82	44.76	22	82	52.80	26	155	8.22	4	40	31.82	14
6321599	NSR1	4	7	13.04	4	7	13.04	4	7	7.97	3	14	0	0
6320863	GCD11	6	14	16.70	6	14	22.20	9	19	6.83	3	5	0	0
398366591	NPL3	9	88	27.05	9	88	13.77	4	40	0	0	0	11.84	3
6319483	IPP1	3	4	9.76	3	4	10.45	2	4	0	0	0	0	0
6325221	PMA2	0	0	14.15	12	55	18.27	15	68	8.24	8	288	1.58	1
6322745	GFA1	17	74	32.50	17	74	19.39	10	26	2.37	2	38	2.93	1
398364525	LSG1	2	3	3.91	2	3	12.03	6	16	0	0	0	0	0
6325038	SAR1	7	50	46.32	7	50	24.74	4	15	11.05	2	21	0	0
6323840	HSC82	30	257	41.28	30	257	42.13	34	374	11.63	8	55	21.13	11
6321881	GAR1	2	9	14.15	2	9	6.34	1	5	0	0	0	0	0
6320164	NAT1	4	4	5.62	4	4	3.40	2	2	0	0	0	0	0
398366403	RPS12	4	26	29.37	4	26	33.57	5	34	23.08	3	162	6.99	1
6322830	CDC16	18	92	24.52	18	92	14.40	11	30	0	0	0	0	0
330443397	ATP1	18	193	37.25	18	193	43.30	22	165	11.38	8	215	17.80	8
6324709	IDH2	4	6	14.91	4	6	9.76	2	2	5.42	2	8	2.98	1
6325178	RPL21B	0	0	22.50	5	19	26.25	6	32	33.13	7	474	11.88	2
6321903	UBA4	14	116	28.86	14	116	31.82	13	37	0	0	0	0	0
6325138	DBP1	0	0	8.10	5	31	8.10	5	16	2.59	2	15	8.43	4
6325248	TAE2	3	3	3.18	3	3	0	0	0	0	0	0	0	0
330443505	DBP10	1	1	1.31	1	1	0	0	0	0	0	0	0	0
6320148	PSA1	10	132	42.66	10	132	27.42	7	51	0	0	0	8.31	2
6325016	HSP82	29	236	42.60	29	236	38.79	29	337	11.42	8	55	21.02	11
6320874	ARB1	9	22	19.84	9	22	9.84	4	14	11.31	6	35	0	0
6319673	PGI1	3	4	7.04	3	4	1.81	1	2	0	0	0	2.71	1
398365251	GSC2	18	49	11.50	18	49	3.69	7	13	1.21	1	3	0	0
6325018	SUI3	8	19	33.68	8	19	23.16	5	18	3.51	2	3	8.42	1
6323374	FKS1	26	81	18.28	26	81	8.00	11	24	1.23	1	2	0	0
14318479	ACT1	11	77	37.33	11	77	49.87	13	139	20.27	7	165	13.33	3
6323554	TUB1	9	84	26.85	9	84	23.49	8	60	5.37	2	4	10.96	3
6321724	ZUO1	9	22	30.95	9	22	27.02	8	19	8.55	5	29	3.70	1
6320842	NUG1	8	14	21.15	8	14	13.65	5	5	4.42	3	7	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1_ _WT_1	TAP_Rtr1_ _WT_1
6322652	YKT6	6	11	32.00	6	11	13.00	3	4	13.00	2	10	7.00	1
6321706	FOL2	6	6	27.16	6	6	24.28	5	6	0	0	0	0	0
6319546	HSP26	4	7	27.10	4	7	15.89	2	5	8.88	2	5	27.10	5
398364355	RPL29	1	2	15.25	1	2	13.56	1	3	13.56	1	2	13.56	1
398365973	TIF4631	2	2	2.21	2	2	1.26	1	1	5.57	5	9	4.31	2
6321929	ARO9	1	1	1.95	1	1	0	0	0	0	0	0	0	0
6321808	YHR020W	19	138	35.76	19	138	31.98	17	70	7.85	5	55	7.41	3
6323729	ADH3	4	21	9.33	4	21	11.47	4	6	1.87	1	1	0	0
6319395	ILS1	13	26	15.11	13	26	16.51	13	29	2.33	2	14	1.31	1
6323969	NIP1	6	7	8.99	6	7	8.87	5	6	3.94	3	6	3.57	2
6322666	FAS1	27	53	16.53	27	53	14.19	21	42	2.63	6	27	2.83	4
6323568	TCB3	6	9	5.05	6	9	2.27	2	4	0.84	1	1	0	0
398366505	HXT7	1	1	2.46	1	1	0	0	0	0	0	0	0	0
6322289	CPS1	3	4	5.73	3	4	3.65	2	2	0	0	0	0	0
398366285	ALA1	18	39	23.59	18	39	25.37	18	55	3.24	3	4	4.07	3
6319553	RPG1	8	14	11.00	8	14	5.71	3	4	10.37	8	38	4.36	3
6325298	TIF5	9	16	24.20	9	16	32.59	11	31	5.19	2	22	0	0
398366229	FAA1	12	35	23.14	12	35	9.29	4	5	4.14	4	40	0	0
6323613	TSA1	4	30	23.98	4	30	34.18	7	71	0	0	0	20.41	3
6323183	RNH203	7	113	82.73	7	113	82.73	9	347	0	0	0	0	0
6320359	CPR1	0	0	0	0	0	20.37	2	7	8.02	1	7	0	0
6321236	ARO8	5	9	14.20	5	9	19.00	6	9	0	0	0	0	0
6323861	PFK2	18	50	25.13	18	50	19.50	13	35	5.01	5	17	14.91	9
398365253	TIF1	17	138	50.63	17	138	50.89	18	146	9.62	4	67	19.24	6
6323511	TUB3	6	64	18.65	6	64	19.10	6	48	3.82	2	2	8.09	2
37362625	SRO9	6	10	20.97	6	10	20.51	5	9	3.92	1	2	4.61	1
398365115	BMH1	8	23	26.97	8	23	41.57	8	27	5.24	1	7	10.11	2
398366063	TYS1	5	13	15.23	5	13	12.18	5	10	2.03	1	5	0	0
6323002	HSP104	31	97	38.99	31	97	16.30	11	23	10.90	10	34	2.09	1
6324902	MYO2	9	9	6.48	9	9	0	0	0	2.10	2	2	0	0
6323947	HAS1	4	6	9.11	4	6	4.95	2	2	1.98	1	2	0	0
6320376	HSP42	0	0	0	0	0	0	0	0	2.93	1	10	0	0
398364255	HYP2	4	21	37.58	4	21	22.93	3	15	0	0	0	0	0
6320809	SNU13	2	5	13.49	2	5	30.95	3	4	5.56	1	1	0	0
6323021	DRS1	2	2	3.19	2	2	0	0	0	0	0	0	0	0
398366329	PYK2	0	0	1.98	1	10	1.98	1	5	0	0	0	1.98	1
398365741	NOC2	1	1	1.41	1	1	0.99	1	1	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6321027	DMC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364483	HOR2	0	0	13.60	3	6	10.80	3	14	3.60	1	14	10.80	3
398365429	PDR5	13	26	11.25	13	26	5.29	6	7	1.79	2	9	1.13	1
6325245	RRP12	2	4	2.12	2	4	0	0	0	0.98	1	3	0	0
398364937	TOP2	6	9	5.25	6	9	0	0	0	0	0	0	0	0
6324660	TCB1	2	2	2.78	2	2	0	0	0	0	0	0	0	0
6320056	COP1	10	15	10.82	10	15	11.41	10	19	2.25	3	17	0	0
398365597	BMH2	6	19	21.98	6	19	39.56	8	23	5.13	1	3	9.89	2
6325030	NEW1	16	24	16.72	16	24	12.29	11	13	4.85	5	16	1.34	1
6321809	RPS27B	2	2	18.29	2	2	29.27	3	19	10.98	1	1	0	0
6321012	CHD1	1	1	0.89	1	1	0	0	0	0	0	0	0	0
6320499	SSD1	6	9	6.16	6	9	4.80	4	5	1.12	1	1	0	0
6323072	TRX1	6	23	66.02	6	23	58.25	6	28	22.33	2	6	0	0
6325238	VTC3	3	3	3.71	3	3	0	0	0	1.20	1	1	1.56	1
398365319	MAK21	2	2	2.34	2	2	1.27	1	1	1.27	1	1	0	0
398364327	LSB3	3	3	8.93	3	3	1.74	1	1	7.84	2	8	0	0
6319513	ATP3	11	43	40.84	11	43	52.09	13	48	4.18	1	9	20.58	5
6324140	SRP1	4	6	7.20	4	6	0	0	0	0	0	0	0	0
6324297	HHT2	0	0	0	0	0	0	0	0	11.76	2	11	0	0
6323306	DBP9	0	0	0	0	0	0	0	0	0	0	0	0	0
6324039	RFC3	0	0	2.65	1	1	0	0	0	0	0	0	0	0
6321193	RTF1	11	55	24.91	11	55	0	0	0	0	0	0	0	0
398366149	TRX2	6	20	66.35	6	20	58.65	6	39	9.62	1	3	0	0
6322452	CCT8	15	39	30.46	15	39	29.58	12	26	17.43	9	89	4.40	2
6321679	PFK1	21	58	26.65	21	58	22.70	17	37	5.67	5	66	7.19	4
6324219	NOP15	0	0	0	0	0	10.00	2	3	0	0	0	0	0
330443717	CTR9	1	1	0.84	1	1	0	0	0	0	0	0	0	0
6323755	ILV2	13	53	28.24	13	53	33.77	20	132	7.13	5	104	4.66	2
6320276	PAA1	3	7	16.23	3	7	7.33	1	2	0	0	0	0	0
6681849	RPL42A	3	3	22.64	3	3	0	0	0	8.49	2	5	0	0
14318517	SEC4	8	38	50.23	8	38	50.23	8	69	5.58	1	28	0	0
6320400	MSS116	7	15	14.16	7	15	3.77	2	3	6.02	3	10	0	0
6321389	TIF4632	1	1	0.98	1	1	0	0	0	1.20	1	1	5.80	2
6324081	RPA49	1	1	3.61	1	1	0	0	0	0	0	0	0	0
398366347	MES1	14	32	22.24	14	32	19.44	11	22	3.99	3	6	0	0
6320418	TCP1	11	31	23.61	11	31	21.29	9	30	13.77	7	39	4.83	2
398365595	PIL1	10	69	39.53	10	69	27.43	8	58	10.32	5	8	10.32	2

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398364417	SAH1	2	9	5.12	2	9	14.48	6	18	0	0	0	2.67	1
6320396	RVB1	14	42	40.82	14	42	19.65	7	15	9.72	4	18	0	0
6320077	CDC48	22	110	35.33	22	110	35.57	23	95	0.96	1	1	1.44	1
6325025	FAS2	31	53	21.78	31	53	15.10	23	43	5.03	5	13	1.43	2
6324517	THI20	20	117	47.19	20	117	23.59	9	18	1.81	1	7	0	0
6323030	NOC3	1	1	2.41	1	1	0	0	0	0	0	0	0	0
6320143	TSR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321318	PRP43	2	3	3.26	2	3	6.65	4	5	0	0	0	0	0
6319618	MAK5	1	1	1.42	1	1	0	0	0	0	0	0	0	0
398364755	URB1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320780	POL5	1	1	1.47	1	1	0.68	1	1	0	0	0	0	0
6325021	RVB2	14	43	38.85	14	43	14.01	5	8	5.10	2	30	6.37	2
6322350	CCT7	15	35	34.91	15	35	16.00	7	10	6.91	3	23	2.36	1
14270688	UTP20	1	1	0.64	1	1	0	0	0	0	0	0	0	0
6324197	KRE33	3	5	3.31	3	5	0	0	0	1.14	1	1	0	0
6321816	RPN1	14	22	17.22	14	22	4.53	3	5	3.73	3	86	0	0
6322901	NAP1	10	25	36.93	10	25	31.89	8	15	0	0	0	8.15	2
6320030	GLT1	27	48	17.06	27	48	11.79	17	43	3.36	6	56	1.72	2
6323470	CAR2	2	2	6.84	2	2	7.31	2	3	0	0	0	4.95	1
6320956	KAP123	15	48	17.43	15	48	9.16	9	25	4.67	5	41	1.80	1
398365941	SEC7	1	1	0.45	1	1	0.60	1	1	1.59	3	8	0	0
6323324	GSP1	5	10	24.20	5	10	15.53	3	8	5.02	1	35	0	0
6322697	GPM1	6	45	32.39	6	45	42.11	8	39	8.10	2	68	37.65	7
6325013	SRP68	2	4	4.51	2	4	0	0	0	0	0	0	0	0
6322103	AVT7	1	2	1.63	1	2	0	0	0	1.63	1	27	0	0
6322049	CCT2	16	90	41.56	16	90	43.83	17	86	0	0	0	10.25	3
330443694	SLA2	10	14	15.19	10	14	3.00	2	2	0	0	0	0	0
6321804	YSC84	2	3	3.85	2	3	1.71	1	1	0	0	0	0	0
398366021	CCT6	9	16	21.79	9	16	21.98	9	36	5.31	3	62	2.75	1
6324042	SEC21	8	14	10.05	8	14	8.66	5	6	0.96	1	8	0	0
398364741	SCS2	3	6	17.21	3	6	13.93	2	2	9.84	2	14	0	0
6325253	LSP1	11	77	38.71	11	77	38.42	10	70	10.26	4	33	22.29	5
6321390	RPT6	6	10	17.04	6	10	10.37	3	5	2.22	1	9	2.96	1
398364807	PNC1	3	8	17.59	3	8	0	0	0	4.17	1	3	0	0
6325164	NOG1	6	10	8.81	6	10	12.06	7	10	1.39	1	2	0	0
6319814	GRX1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322823	MAE1	9	26	18.09	9	26	10.31	4	6	13.45	8	83	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1_ _WT_1	TAP_Rtr1_ _WT_1
6320255	TPI1	3	4	14.52	3	4	23.79	4	8	3.63	1	1	8.06	1
6320983	GDI1	4	6	8.65	4	6	3.10	1	1	0	0	0	0	0
398365205	RPT5	12	44	36.87	12	44	30.41	10	23	8.99	4	34	0	0
6324992	KEL3	1	1	1.54	1	1	0	0	0	0	0	0	0	0
6323028	DNM1	2	3	3.43	2	3	1.85	1	1	0	0	0	0	0
398365977	RPT4	4	8	10.07	4	8	2.75	1	1	0	0	0	2.75	1
6320058	CCT4	7	24	17.05	7	24	17.61	7	13	8.14	3	36	3.03	1
6320379	HMO1	3	4	13.41	3	4	16.67	3	11	0	0	0	0	0
6322508	CYC1	0	0	0	0	0	0	0	0	7.34	1	1	0	0
6325137	VPS30	0	0	0	0	0	0	0	0	0	0	0	0	0
6321301	SEC27	9	18	13.39	9	18	24.30	17	37	7.99	7	25	0	0
6321673	YHB1	2	5	7.27	2	5	11.78	3	18	0	0	0	0	0
27469359	SYN8	0	0	0	0	0	0	0	0	0	0	0	0	0
6324433	CDC33	4	13	18.78	4	13	21.13	5	11	21.60	5	16	0	0
6323571	POB3	4	4	9.60	4	4	6.52	2	2	0	0	0	0	0
398365157	NOP2	4	5	8.09	4	5	16.18	7	8	0	0	0	0	0
6321228	YPT32	7	40	35.14	7	40	36.49	8	74	4.95	1	17	0	0
398366559	RPT3	6	16	18.69	6	16	6.78	2	2	0	0	0	2.80	1
6322704	RPT1	9	17	20.99	9	17	12.63	5	10	14.56	6	61	2.57	1
398366379	PRT1	19	42	27.65	19	42	17.69	10	15	4.85	4	7	1.70	1
398365125	PDA1	8	12	20.95	8	12	19.29	7	21	7.86	3	7	7.14	2
398364913	GND1	6	11	17.18	6	11	5.52	2	6	0	0	0	9.20	4
6322637	SAC1	3	6	6.74	3	6	4.82	2	2	1.61	1	8	0	0
398365933	CYS4	20	194	52.47	20	194	66.47	30	154	9.86	5	77	24.85	8
6323903	FAA4	6	11	11.10	6	11	7.06	4	5	4.18	3	47	0	0
6319806	PDI1	14	70	34.87	14	70	37.74	15	73	7.85	3	10	7.66	2
6323512	PHO84	0	0	0	0	0	0	0	0	0	0	0	0	0
6325367	RPC40	5	5	19.10	5	5	13.43	4	7	0	0	0	0	0
6319835	GBP2	2	2	5.85	2	2	4.22	1	1	0	0	0	0	0
6319290	MYO4	0	0	0	0	0	0	0	0	0	0	0	0	0
14318531	GCN20	4	9	6.38	4	9	2.79	2	2	7.98	6	19	0	0
6320869	YPT31	9	46	49.33	9	46	68.16	11	96	4.93	1	17	0	0
398365617	UTP22	2	2	1.62	2	2	1.13	1	1	0	0	0	0	0
398366209	RTN1	3	5	13.22	3	5	14.24	4	13	14.24	4	87	9.83	2
6325097	CDC60	15	30	17.61	15	30	22.20	18	38	0	0	0	0	0
6323294	RPS28B	2	22	31.34	2	22	37.31	3	30	0	0	0	13.43	1
6325151	SSE1	25	178	41.56	25	178	48.92	29	256	8.37	6	92	11.69	6

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398365213	ARL1	2	3	13.66	2	3	0	0	0	7.65	1	1	0	0
6319587	YSA1	2	2	11.69	2	2	0	0	0	0	0	0	0	0
14318480	YPT1	8	54	49.51	8	54	53.40	11	81	6.80	1	4	0	0
398366227	SEC26	12	16	14.18	12	16	5.76	5	7	3.80	3	11	0	0
6322381	SCP160	40	155	39.12	40	155	30.69	32	78	7.20	10	107	4.99	4
6325131	NAN1	3	5	3.79	3	5	7.48	5	6	2.46	2	13	0	0
6320226	SES1	9	13	20.56	9	13	21.00	7	14	3.03	1	1	3.25	1
6322853	VPS1	12	21	23.44	12	21	8.10	5	7	7.95	5	70	1.70	1
6321184	HXK2	10	18	25.31	10	18	29.42	11	26	6.17	3	8	4.32	1
6323259	CDC42	3	8	19.90	3	8	5.24	1	5	5.76	1	2	0	0
398366139	GCD6	5	5	8.57	5	5	0	0	0	0	0	0	0	0
6325267	RPA135	7	8	8.40	7	8	7.73	6	7	1.41	2	2	1.50	1
6323722	PDS5	2	2	2.58	2	2	0	0	0	1.49	2	2	0	0
6319809	GLK1	3	5	6.60	3	5	7.80	3	6	0	0	0	0	0
6321857	SSF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324720	PNO1	5	11	22.63	5	11	13.87	3	7	20.44	6	16	0	0
398364619	UGP1	7	11	18.24	7	11	7.01	3	3	0	0	0	3.21	1
6319582	PHO88	4	7	26.06	4	7	22.34	3	7	11.17	2	2	0	0
6325103	PEP4	7	31	24.44	7	31	10.86	3	4	4.20	1	1	20.49	6
6319499	OLA1	15	43	39.85	15	43	34.01	11	27	13.96	6	143	7.87	2
6322254	LAA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322385	NET1	2	3	2.10	2	3	1.26	1	2	0	0	0	0	0
82795243	NHP2	0	0	0	0	0	9.62	1	1	0	0	0	0	0
6325039	BMS1	0	0	0	0	0	1.10	1	1	0	0	0	0	0
398365715	ENP1	0	0	0	0	0	3.52	1	1	6.63	2	2	0	0
398366507	SVF1	6	19	17.46	6	19	28.69	10	127	0	0	0	0	0
6322866	YPT52	7	28	42.74	7	28	60.68	10	37	3.85	1	1	0	0
6320197	RPT2	8	19	24.94	8	19	5.49	2	2	1.83	1	9	0	0
398364435	GVP36	11	56	38.65	11	56	49.39	12	45	10.12	3	25	0	0
6324273	POR1	8	13	36.04	8	13	12.72	3	5	9.54	2	33	0	0
83578109	JIP5	0	0	0	0	0	0	0	0	0	0	0	0	0
154199622	HRB1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383791	RIM1	3	4	28.89	3	4	17.04	2	4	0	0	0	0	0
6325345	SRP54	3	6	6.47	3	6	4.07	2	3	3.33	1	5	0	0
398366281	SDA1	1	1	1.83	1	1	0.91	1	1	0	0	0	0	0
398366061	IKI3	5	5	4.08	5	5	5.34	6	9	0.74	1	1	0	0
6321940	CRP1	2	2	5.38	2	2	3.23	1	3	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1_ _WT_1	TAP_Rtr1_ _WT_1
398366337	APL6	1	1	0.99	1	1	2.22	1	1	0	0	0	0	0
6320242	KRS1	10	43	18.78	10	43	28.26	14	50	5.25	3	5	0	0
6324705	YOR131C	2	2	9.17	2	2	0	0	0	5.50	1	1	0	0
6319760	SSH1	2	3	4.69	2	3	2.86	1	2	1.84	1	4	0	0
6323335	ACO1	3	4	5.66	3	4	2.83	2	3	1.67	1	1	1.67	1
330443748	GLN1	6	14	17.84	6	14	17.30	5	15	15.14	6	40	0	0
6319464	SLA1	2	2	1.53	2	2	1.05	1	1	0	0	0	0	0
14627173	BBC1	1	2	0.95	1	2	0	0	0	0.95	1	4	0	0
398364931	URB2	0	0	0	0	0	0	0	0	0	0	0	0	0
6325423	RHO1	5	8	42.58	5	8	35.41	5	9	0	0	0	0	0
6321837	AAP1	1	1	1.05	1	1	0	0	0	0	0	0	0	0
6319281	RBG1	2	2	7.86	2	2	10.30	2	2	4.61	2	2	0	0
6323379	KAP95	2	3	3.14	2	3	3.37	2	3	0	0	0	0	0
398366047	RNR4	10	47	37.10	10	47	33.33	9	52	4.06	1	6	13.62	3
6322446	CCT3	12	31	28.84	12	31	18.73	8	25	14.04	7	24	0	0
398365063	LAT1	5	13	15.15	5	13	12.45	5	6	0	0	0	14.32	5
6322427	HCA4	0	0	0	0	0	1.82	1	1	0	0	0	0	0
6322073	RHO3	4	7	19.91	4	7	0	0	0	0	0	0	0	0
6324630	NOB1	2	2	4.36	2	2	0	0	0	1.96	1	2	0	0
398364409	RET2	0	0	0	0	0	2.38	1	1	0	0	0	0	0
6319796	SPB1	3	5	4.88	3	5	0	0	0	0	0	0	0	0
6323968	PSE1	1	1	1.01	1	1	1.56	1	5	3.67	2	10	0	0
6322934	RPF2	1	3	2.62	1	3	6.10	2	2	0	0	0	0	0
398365233	RPS21A	0	0	26.44	2	4	36.78	3	18	14.94	1	3	0	0
6322325	RPS21B	3	6	44.83	3	6	55.17	4	20	14.94	1	3	0	0
398366473	SSF2	0	0	0	0	0	0	0	0	0	0	0	0	0
86559466	CTT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325449	QCR2	6	8	19.02	6	8	7.34	2	2	0	0	0	0	0
6323537	TSL1	5	5	5.56	5	5	1.18	1	1	0	0	0	0	0
6320914	RNR1	13	33	17.34	13	33	18.92	12	30	4.95	5	37	2.03	1
6323182	ACS2	13	27	24.60	13	27	22.99	14	32	6.44	5	107	0	0
6319432	URA7	15	44	27.81	15	44	18.31	8	15	15.89	10	179	0	0
6323135	MDN1	1	2	0.22	1	2	0	0	0	0	0	0	0	0
398365477	ATP2	14	101	38.16	14	101	30.92	11	48	5.87	3	11	18.59	7
6322988	VPS13	8	8	2.96	8	8	0.54	1	1	1.62	3	7	0	0
6322794	TMA19	5	12	29.94	5	12	34.13	6	20	0	0	0	17.96	2
6323037	RLP24	1	1	6.53	1	1	11.56	2	2	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398364441	NOT3	8	12	13.16	8	12	7.54	4	5	0	0	0	0	0
398366191	RPN13	1	1	8.33	1	1	22.44	3	6	0	0	0	0	0
6321453	YGR016W	0	0	0	0	0	0	0	0	0	0	0	0	0
6320930	ILV1	14	90	33.85	14	90	31.60	12	43	7.99	5	33	3.65	1
6323016	PUF3	1	1	1.14	1	1	0	0	0	0	0	0	0	0
37362700	SRP72	0	0	0	0	0	5.47	2	3	0	0	0	0	0
6321326	TAF6	3	5	8.91	3	5	0	0	0	0	0	0	0	0
6322565	ADO1	0	0	0	0	0	10.59	2	2	3.53	1	3	0	0
6319722	ISW1	4	6	4.34	4	6	0	0	0	1.95	1	3	0	0
330443605	SET2	2	2	3.68	2	2	0	0	0	0	0	0	0	0
398365101	ARP3	3	3	6.68	3	3	4.90	2	6	5.12	2	3	0	0
6321780	STE20	0	0	0	0	0	1.28	1	2	1.92	1	1	0	0
6321776	PRS3	6	8	22.19	6	8	24.69	7	9	9.06	3	64	0	0
6323214	RPL37A	0	0	0	0	0	13.64	1	1	39.77	5	36	13.64	1
6321757	ECM29	0	0	0	0	0	1.12	1	3	1.12	2	3	0	0
398365069	ADH5	2	10	5.70	2	10	2.28	1	23	3.42	1	1	2.28	1
398365695	CPR8	0	0	0	0	0	0	0	0	0	0	0	0	0
6319854	RVS161	4	12	19.62	4	12	2.64	1	1	6.42	2	6	0	0
6324620	DBP5	6	10	17.43	6	10	5.81	2	4	6.22	3	8	0	0
6321399	YGL039W	2	3	9.20	2	3	4.60	1	1	0	0	0	0	0
398365061	SUP45	14	34	40.27	14	34	16.48	6	9	3.66	1	1	4.35	1
6324673	KTR1	6	10	16.54	6	10	3.31	1	1	0	0	0	0	0
6323387	ILV5	9	15	28.10	9	15	29.87	9	15	0	0	0	0	0
6323795	TIF34	10	16	38.62	10	16	27.09	7	12	10.09	3	9	0	0
6322667	PRS1	5	15	16.39	5	15	18.03	6	15	3.28	1	14	7.73	2
6320910	RGI1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365575	HOM6	0	0	0	0	0	0	0	0	0	0	0	0	0
6323916	TIF11	2	2	15.03	2	2	26.80	3	3	0	0	0	0	0
6321612	RBG2	4	8	14.95	4	8	2.99	1	3	8.15	3	5	8.97	2
6323225	PWP1	6	7	13.89	6	7	7.12	4	6	1.56	1	1	0	0
6324737	SSP1	0	0	0	0	0	6.38	1	1	6.38	1	4	0	0
6323450	CDC73	2	2	5.60	2	2	0	0	0	0	0	0	0	0
398365461	SRP40	1	1	2.96	1	1	0	0	0	0	0	0	0	0
6320708	RPL37B	0	0	0	0	0	0	0	0	26.14	4	15	0	0
398364957	MKT1	2	2	4.10	2	2	1.33	1	1	1.20	1	1	0	0
6320079	YDL124W	1	1	4.49	1	1	8.33	2	2	0	0	0	0	0
398364371	RSC8	2	2	5.21	2	2	4.31	2	2	5.21	2	2	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6325418	GPH1	8	8	11.97	8	8	0	0	0	0	0	0	0	0
6321859	DYS1	3	4	8.01	3	4	17.31	5	15	0	0	0	3.10	1
6323967	GAS1	8	20	17.53	8	20	7.87	4	9	4.11	3	10	2.68	1
6319695	PYC2	11	19	10.68	11	19	6.44	6	9	6.53	8	30	6.10	5
6323642	YPT7	9	82	60.10	9	82	65.38	10	106	5.29	1	17	0	0
6320552	HXT3	1	1	2.12	1	1	3.70	1	1	0	0	0	0	0
6325196	ALD6	3	4	6.60	3	4	13.20	5	6	2.20	1	13	0	0
6319315	EFB1	4	32	23.30	4	32	21.84	4	28	0	0	0	0	0
10383811	CDC39	2	2	1.42	2	2	0.71	1	1	0.52	1	3	0	0
6321753	SBP1	5	18	23.47	5	18	27.55	5	24	0	0	0	4.42	1
6320764	DLD3	1	1	2.22	1	1	0	0	0	0	0	0	0	0
398366413	ERV29	0	0	0	0	0	0	0	0	0	0	0	0	0
6320969	GLO3	1	3	1.42	1	3	0	0	0	1.42	1	59	0	0
6319602	TPS1	6	15	15.76	6	15	10.71	4	9	0	0	0	2.22	1
6579193	NMD2	0	0	0	0	0	0	0	0	0	0	0	0	0
6321990	UTP9	3	4	7.30	3	4	6.09	3	4	0	0	0	0	0
398366561	SXM1	0	0	0	0	0	1.06	1	1	0	0	0	0	0
6320157	NPC2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364687	NOP9	0	0	0	0	0	0	0	0	0	0	0	0	0
22532997	BUD27	0	0	0	0	0	0	0	0	0	0	0	0	0
398366449	BFR2	3	4	5.62	3	4	8.99	4	6	0	0	0	0	0
398365717	TRM1	3	4	5.44	3	4	9.82	4	9	0	0	0	4.21	2
6323362	CHS5	0	0	0	0	0	2.24	1	2	0	0	0	0	0
398364573	PAN1	6	8	5.20	6	8	10.00	11	27	0	0	0	3.78	3
6320720	GRX2	1	1	6.99	1	1	6.99	1	2	0	0	0	0	0
6321595	PTI1	0	0	0	0	0	6.12	1	1	0	0	0	0	0
398364853	OCA1	1	1	5.46	1	1	0	0	0	4.20	1	1	0	0
6323852	YMR196	0	0	0	0	0	0	0	0	0	0	0	0	0
33438816	YILO02W-	0	0	0	0	0	0	0	0	0	0	0	0	0
398366575	RRP17	0	0	0	0	0	0	0	0	0	0	0	0	0
6323359	TMA10	0	0	0	0	0	0	0	0	0	0	0	0	0
398365853	WTM1	4	5	12.59	4	5	8.01	3	4	4.58	2	10	0	0
6320181	GPD1	9	61	31.20	9	61	14.83	5	7	4.09	2	12	0	0
6323518	NAB6	3	3	3.35	3	3	0	0	0	0	0	0	0	0
6321221	KIP3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365131	RFC2	0	0	0	0	0	4.82	1	2	0	0	0	0	0
6322294	ERG20	4	4	12.22	4	4	2.84	1	1	2.84	1	1	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6321024	ECM32	0	0	0	0	0	0	0	0	0	0	0	0	0
37362660	FUR1	5	17	22.69	5	17	34.72	8	25	4.63	1	1	0	0
398364665	SEC18	11	19	18.73	11	19	4.75	3	5	2.51	2	3	1.58	1
6323391	ADE13	0	0	0	0	0	0	0	0	0	0	0	0	0
6323699	STV1	3	4	3.37	3	4	1.24	1	1	0	0	0	0	0
6321412	TRP5	17	186	35.50	17	186	28.85	15	83	0	0	0	8.77	4
6321973	OYE2	0	0	0	0	0	4.75	1	1	2.25	1	2	0	0
6321801	ARD1	0	0	0	0	0	4.62	1	1	0	0	0	0	0
10383761	HIS4	5	8	8.14	5	8	2.13	1	2	0	0	0	0	0
398365759	SAC6	9	18	18.85	9	18	11.37	5	11	1.56	1	1	0	0
6321721	BGL2	2	3	7.99	2	3	7.03	2	2	0	0	0	0	0
398364979	CCZ1	1	1	1.42	1	1	0	0	0	0	0	0	0	0
42742256	VTC4	2	2	2.36	2	2	2.50	2	2	4.85	3	8	0	0
6321814	THR1	2	2	5.32	2	2	0	0	0	3.08	1	1	0	0
10383798	PHO87	0	0	0.76	1	1	0	0	0	0	0	0	0	0
6319847	CDC10	3	5	15.53	3	5	2.48	1	1	9.32	3	6	0	0
398365363	URA8	6	11	11.94	6	11	2.42	1	1	6.23	4	7	0	0
6323138	AHP1	3	8	29.55	3	8	14.77	2	2	0	0	0	0	0
6320657	UTP6	0	0	0	0	0	0	0	0	0	0	0	0	0
6321919	YHR127W	0	0	0	0	0	0	0	0	0	0	0	0	0
41629688	PSP2	1	2	1.69	1	2	1.69	1	1	2.87	2	6	0	0
398364633	APS3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322843	MRT4	3	6	16.10	3	6	21.19	4	4	4.24	1	1	0	0
330443486	PAT1	2	2	2.01	2	2	0	0	0	4.15	3	5	2.76	1
37362659	RRP3	0	0	0	0	0	0	0	0	2.59	1	11	0	0
6320845	SEC3	0	0	0	0	0	0	0	0	0	0	0	0	0
6324791	RFC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325369	MRD1	1	1	1.58	1	1	0	0	0	1.58	1	1	0	0
6322635	YRA2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320368	SEC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364935	UBP3	4	6	5.04	4	6	0	0	0	0	0	0	0	0
6320789	GLY1	5	10	17.57	5	10	31.27	9	26	0	0	0	0	0
6325172	SEC16	0	0	0	0	0	0	0	0	0	0	0	0	0
398364311	TMA20	2	4	12.71	2	4	0	0	0	0	0	0	0	0
6320926	UTP7	0	0	0	0	0	0	0	0	0	0	0	0	0
330443568	ENP2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324328	CIT1	1	1	2.09	1	1	0	0	0	2.09	1	1	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6324550	IFM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324950	ALD4	5	5	12.33	5	5	6.94	2	2	2.12	1	4	3.47	1
6324635	CKA2	0	0	0	0	0	6.78	2	2	0	0	0	0	0
6319849	YCP4	4	16	20.65	4	16	21.46	4	9	0	0	0	0	0
6325202	LGE1	1	1	4.52	1	1	0	0	0	0	0	0	0	0
398366027	SUR4	0	0	0	0	0	4.93	1	2	5.22	1	2	0	0
6321641	PCT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325280	EAF3	1	1	2.00	1	1	0	0	0	3.24	1	1	0	0
6320803	MCM3	3	5	4.02	3	5	0	0	0	4.33	4	10	0	0
6323872	SKY1	1	1	1.48	1	1	0	0	0	0	0	0	0	0
6325140	IDI1	2	6	9.03	2	6	9.38	2	5	4.86	1	6	0	0
6320049	SAS10	3	3	6.56	3	3	4.10	2	3	1.31	1	1	0	0
6324989	DIM1	3	4	11.01	3	4	3.14	1	1	0	0	0	0	0
398366039	PBP1	1	2	2.22	1	2	0	0	0	0	0	0	9.70	4
6323088	REX2	1	1	4.46	1	1	0	0	0	0	0	0	0	0
6324899	PRO2	1	1	2.41	1	1	0	0	0	0	0	0	0	0
398365467	YNL010W	2	7	7.47	2	7	9.13	1	1	3.73	1	1	0	0
41629671	SEA4	3	3	2.99	3	3	0	0	0	2.41	2	3	0	0
6320704	PUF6	1	1	1.83	1	1	5.18	3	3	0	0	0	0	0
6323650	PLB1	1	1	1.66	1	1	0	0	0	0	0	0	0	0
6323158	DIP2	1	1	1.48	1	1	0	0	0	0	0	0	0	0
6322727	SRP21	0	0	0	0	0	0	0	0	0	0	0	0	0
14318504	MDJ1	0	0	0	0	0	3.33	1	1	0	0	0	0	0
6322878	GCN3	0	0	0	0	0	0	0	0	3.61	1	7	0	0
330443380	PSK1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364789	UFD4	0	0	0	0	0	0	0	0	0	0	0	0	0
6319903	PWP2	2	3	2.17	2	3	1.08	1	1	0	0	0	0	0
6324613	CKB2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324568	TOP1	0	0	0	0	0	1.17	1	1	0	0	0	0	0
398365393	MDM32	0	0	0	0	0	1.61	1	1	0	0	0	0	0
6322731	VPH2	0	0	0	0	0	4.19	1	1	0	0	0	0	0
6323653	HXT2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320236	MRH1	1	1	5.00	1	1	5.00	1	6	3.75	2	8	0	0
398364559	NSP1	2	4	3.40	2	4	23.33	13	23	0	0	0	2.07	1
6321498	ADE6	4	5	3.90	4	5	0	0	0	2.80	3	14	0	0
6319326	ADE1	1	4	3.92	1	4	7.84	2	3	3.27	1	3	0	0
6321812	MYO1	1	1	0.47	1	1	0.78	1	1	0	0	0	5.39	7

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6324734	MTR10	0	0	0	0	0	0	0	0	0	0	0	0	0
6320284	PET100	0	0	0	0	0	0	0	0	0	0	0	0	0
6323411	SEC61	0	0	0	0	0	0	0	0	0	0	0	0	0
6320883	ERG28	0	0	0	0	0	0	0	0	7.43	1	10	0	0
6324593	YOR019W	0	0	0	0	0	0	0	0	1.23	1	2	0	0
6320432	ADK1	5	7	22.97	5	7	31.08	6	11	4.05	1	1	0	0
6322919	CCP1	1	1	3.60	1	1	11.63	3	7	0	0	0	0	0
6320204	NTH1	5	9	6.66	5	9	0	0	0	0	0	0	3.33	2
6320279	TPS2	9	11	12.61	9	11	1.23	1	1	0	0	0	7.37	5
6323681	CCS1	3	4	13.25	3	4	10.04	2	2	0	0	0	0	0
398366097	ADH6	1	1	3.06	1	1	16.39	4	5	0	0	0	0	0
6324381	NOG2	4	4	9.67	4	4	2.88	1	3	1.85	1	3	0	0
6323974	PRE5	2	3	9.40	2	3	0	0	0	0	0	0	0	0
6320090	ATG20	1	1	2.34	1	1	0	0	0	0	0	0	0	0
37362683	SCJ1	2	2	8.22	2	2	8.22	2	4	0	0	0	0	0
6323676	ARP9	1	2	2.57	1	2	0	0	0	0	0	0	0	0
398364547	SER3	5	7	14.29	5	7	19.19	7	13	6.40	3	6	0	0
6319439	HEK2	1	1	3.94	1	1	2.10	1	1	14.96	4	81	0	0
6321964	NMD3	6	8	12.55	6	8	13.13	6	12	5.41	2	4	0	0
6321984	ERG9	2	4	4.50	2	4	0	0	0	2.25	1	1	0	0
6323768	ADE17	15	27	37.67	15	27	4.22	2	3	0	0	0	0	0
398365401	CPA2	6	11	7.96	6	11	12.34	10	22	0	0	0	0	0
42742253	ATG27	4	6	17.34	4	6	0	0	0	0	0	0	0	0
6324191	SRV2	7	28	15.78	7	28	6.65	3	5	5.70	1	1	6.27	2
6322476	ILV3	10	25	22.74	10	25	20.85	8	20	3.08	2	8	0	0
6321900	GGA2	2	4	4.79	2	4	9.91	4	6	1.37	1	3	2.91	1
330443535	STE2	1	2	3.25	1	2	2.32	1	1	0	0	0	0	0
6324343	ACC1	28	64	15.18	28	64	13.57	23	40	3.72	8	46	3.00	5
6319434	APL3	0	0	0	0	0	0	0	0	0	0	0	0	0
398364593	MET6	0	0	0	0	0	0	0	0	0	0	0	0	0
6323751	YPK2	0	0	0	0	0	0	0	0	1.92	1	1	0	0
330443495	AIM6	2	2	5.90	2	2	0	0	0	0	0	0	0	0
6323752	PGM2	0	0	0	0	0	0	0	0	0	0	0	0	0
82795256	SEC28	1	1	3.38	1	1	8.78	2	3	0	0	0	0	0
6322215	YIR024C	0	0	0	0	0	0	0	0	0	0	0	0	0
6323291	YPT6	3	8	16.28	3	8	5.12	1	18	0	0	0	0	0
6321632	PDX1	1	1	3.66	1	1	0	0	0	3.66	1	2	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398365005	RIX1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323975	YMR315	3	3	8.31	3	3	0	0	0	0	0	0	2.01	1
6323740	UTP15	2	2	5.46	2	2	11.70	4	4	0	0	0	0	0
6322743	AAT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323033	SSL1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365521	SWT1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365403	NUP133	3	3	3.11	3	3	1.38	1	1	1.04	1	2	0	0
6319276	CDC24	0	0	0	0	0	0	0	0	3.04	2	5	0	0
6323251	UTP13	4	4	7.34	4	4	15.79	9	23	2.94	2	6	0	0
6321913	LSM12	0	0	0	0	0	0	0	0	0	0	0	17.11	2
6322257	RCY1	1	3	1.07	1	3	1.43	1	1	0	0	0	0	0
6319900	RSC6	0	0	0	0	0	0	0	0	0	0	0	0	0
398364743	IRC2	5	7	21.29	5	7	0	0	0	7.60	2	3	0	0
6320070	CDC53	0	0	0	0	0	0	0	0	1.10	1	3	1.84	1
398364601	NFU1	1	1	4.69	1	1	17.97	4	8	0	0	0	0	0
6323229	YKE2	0	0	0	0	0	0	0	0	0	0	0	14.91	1
6322154	CKA1	0	0	0	0	0	2.42	1	1	0	0	0	0	0
6323687	IOC4	0	0	0	0	0	0	0	0	0	0	0	0	0
6325372	RGC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325234	MET12	2	2	3.65	2	2	2.28	1	1	1.67	1	1	0	0
10383790	RHB1	0	0	0	0	0	0	0	0	5.26	1	2	0	0
10383792	SYP1	4	5	4.83	4	5	0	0	0	0	0	0	8.05	4
398365079	SEC9	0	0	0	0	0	0	0	0	0	0	0	0	0
398365907	APM3	1	1	2.28	1	1	0	0	0	0	0	0	0	0
6322783	YNK1	1	1	7.84	1	1	18.95	2	6	0	0	0	0	0
6324470	TPT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321843	CIC1	2	2	5.59	2	2	3.46	1	1	0	0	0	0	0
6319298	CCR4	3	3	4.78	3	3	1.91	1	1	2.75	2	7	0	0
6321257	GTS1	1	3	3.28	1	3	4.04	1	1	0	0	0	0	0
6321341	SRM1	2	2	4.15	2	2	0	0	0	0	0	0	0	0
398365245	PFY1	1	1	7.94	1	1	7.94	1	1	0	0	0	0	0
6324375	FPK1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364253	AYR1	1	1	3.37	1	1	0	0	0	2.69	1	1	0	0
398365397	YGR054W	1	1	2.34	1	1	3.58	2	2	1.87	1	3	0	0
398365527	RLP7	1	1	4.04	1	1	4.04	1	2	11.18	3	22	0	0
6319976	SHS1	2	3	3.63	2	3	4.17	2	2	0	0	0	0	0
398364769	RSP5	5	7	7.66	5	7	5.93	3	4	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
14318538	YFR016C	0	0	0	0	0	0	0	0	0	0	0	0	0
6321647	SER2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365769	SHG1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321009	SPT2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365509	RRP1	3	3	10.43	3	3	3.60	1	1	2.88	1	1	0	0
6323756	MYO5	4	4	4.10	4	4	0	0	0	2.05	2	2	0	0
6324839	RBL2	0	0	0	0	0	23.58	2	2	0	0	0	0	0
6319685	DUR1,2	0	0	0	0	0	0.76	1	1	0	0	0	0.76	1
6325273	TIF6	2	3	11.43	2	3	12.65	2	4	0	0	0	0	0
6324938	PRE10	0	0	0	0	0	0	0	0	0	0	0	0	0
6324996	APM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324522	GAL11	0	0	0	0	0	0	0	0	0	0	0	1.02	1
6324113	RAP1	4	8	5.93	4	8	1.69	1	1	0	0	0	0	0
398366009	YGR169C-	0	0	0	0	0	0	0	0	0	0	0	0	0
6319564	POL30	0	0	0	0	0	0	0	0	0	0	0	0	0
6323546	UTP14	1	1	1.11	1	1	0	0	0	0	0	0	0	0
6324312	PUB1	2	2	3.75	2	2	4.86	2	9	0	0	0	0	0
6325286	APL4	0	0	0	0	0	0	0	0	0	0	0	0	0
6323401	SSQ1	0	0	0	0	0	5.02	2	3	0	0	0	0	0
6324048	HCH1	1	2	7.84	1	2	0	0	0	0	0	0	0	0
6319627	APD1	2	2	7.28	2	2	0	0	0	0	0	0	0	0
6320102	DUN1	1	1	1.75	1	1	0	0	0	0	0	0	0	0
10383806	SOL2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320205	YRB1	0	0	0	0	0	14.93	2	8	5.47	1	1	0	0
398366613	TOM1	4	4	1.32	4	4	0	0	0	0.80	2	2	0	0
398365693	TMA64	1	1	1.77	1	1	0	0	0	0	0	0	0	0
6323893	BCH1	1	1	1.24	1	1	0	0	0	1.24	1	1	0	0
398366023	YTM1	2	2	4.35	2	2	0	0	0	4.57	2	3	0	0
6319437	STU1	1	1	0.66	1	1	0	0	0	0	0	0	0	0
398364833	MET4	1	1	2.08	1	1	0	0	0	0	0	0	0	0
6321424	PUF4	0	0	0	0	0	0	0	0	0.90	1	3	0	0
6322411	MTR4	8	9	10.16	8	9	1.49	1	3	0.84	1	1	0	0
14318501	LPD1	6	9	17.03	6	9	16.43	5	6	0	0	0	0	0
6321890	SFB3	1	1	0.86	1	1	1.94	1	1	0	0	0	0	0
398366661	GIN4	1	1	1.31	1	1	0	0	0	0	0	0	0	0
398365725	SOL1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323570	DAK1	7	20	16.10	7	20	16.61	7	15	1.71	1	1	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
398364717	OLE1	0	0	0	0	0	2.35	1	1	0	0	0	0	0
398364715	OST1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320639	CYM1	0	0	0	0	0	0	0	0	1.01	1	1	0	0
6324109	ADE12	6	22	18.01	6	22	7.39	2	5	0	0	0	0	0
330443698	CHS1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365123	AQR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324745	LCB4	3	3	7.37	3	3	0	0	0	2.08	1	4	0	0
398366587	RPN9	6	8	16.03	6	8	10.43	3	9	4.83	2	7	0	0
6320837	MNN1	2	2	2.23	2	2	0	0	0	0	0	0	0	0
6324594	HSP10	3	3	34.91	3	3	65.09	6	39	0	0	0	0	0
6320859	RPN3	8	19	19.31	8	19	7.27	3	3	3.63	2	2	0	0
398366611	NHX1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320395	SLY1	5	12	10.06	5	12	4.05	2	2	1.50	1	14	2.10	1
41629692	YIP4	1	1	5.11	1	1	0	0	0	0	0	0	0	0
330443374	DRS2	1	1	0.74	1	1	0	0	0	0	0	0	1.77	1
6325207	MNN9	1	1	2.53	1	1	0	0	0	0	0	0	0	0
6323858	ERG20	0	0	0	0	0	0	0	0	0	0	0	0	0
6319489	MNN2	3	3	6.87	3	3	0	0	0	0	0	0	0	0
6319726	ARO4	3	5	9.46	3	5	11.08	3	10	0	0	0	0	0
398364279	VAC8	2	2	4.67	2	2	4.67	2	2	2.94	2	6	0	0
6324555	TLG2	2	2	5.04	2	2	0	0	0	0	0	0	0	0
398366239	GNT1	2	20	3.05	2	20	1.22	1	1	0	0	0	0	0
6322249	OPT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319454	PEP1	14	17	11.15	14	17	4.12	5	6	1.14	2	3	0	0
6322787	YKL063C	0	0	0	0	0	0	0	0	0	0	0	0	0
6324645	NRT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323270	YLR241W	1	1	1.28	1	1	0	0	0	0	0	0	0	0
14318577	RPN12	6	12	31.39	6	12	31.39	8	16	8.03	1	2	0	0
6324061	LYP1	0	0	2.29	1	1	0	0	0	0	0	0	0	0
6320993	FTR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320705	ITR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319319	ERP1	2	4	9.59	2	4	0	0	0	0	0	0	0	0
6324828	SEC63	2	2	3.02	2	2	4.98	2	2	7.24	5	10	0	0
6319429	FUI1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325293	VMA13	9	17	22.38	9	17	10.67	4	4	2.72	1	8	0	0
10383762	RNQ1	1	1	4.20	1	1	4.20	1	1	2.22	1	23	0	0
6320691	KRE2	4	4	10.41	4	4	0	0	0	5.20	1	1	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6323630	ERV25	0	0	0	0	0	0	0	0	0	0	0	0	0
6323716	TVP18	0	0	0	0	0	0	0	0	0	0	0	0	0
6319289	SNC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364281	EMP24	1	1	6.90	1	1	6.90	1	1	0	0	0	0	0
6320298	DNF2	3	3	2.30	3	3	0	0	0	0	0	0	0	0
330443597	MNN5	3	5	6.48	3	5	0	0	0	0	0	0	0	0
6320897	FCY2	1	1	1.69	1	1	0	0	0	0	0	0	0	0
330443542	SAP155	3	3	2.99	3	3	0	0	0	0	0	0	0	0
14318521	SPB4	0	0	0	0	0	0	0	0	0	0	0	0	0
6323367	NUP2	0	0	0	0	0	8.47	3	4	0	0	0	0	0
6319303	TPD3	2	3	4.57	2	3	4.72	3	4	0	0	0	0	0
6324166	RIA1	1	1	0.99	1	1	0	0	0	1.17	1	2	0	0
398364557	AIM21	0	0	0	0	0	2.50	1	3	0	0	0	0	0
6322369	SRS2	1	1	0.68	1	1	0.68	1	1	0	0	0	0	0
6324088	ZWF1	1	1	3.37	1	1	0	0	0	0	0	0	0	0
6319890	TAF2	1	1	0.57	1	1	0	0	0	0	0	0	0	0
6320033	SFA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325429	BSP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323022	YLL007C	0	0	0	0	0	0	0	0	0	0	0	0	0
6324918	YOR342C	0	0	0	0	0	0	0	0	0	0	0	0	0
6320091	TRM3	1	1	0.63	1	1	0	0	0	0.77	1	3	0	0
6322925	SIS2	0	0	0	0	0	0	0	0	0	0	0	0	0
6321999	SCH9	0	0	0	0	0	0	0	0	4.25	3	17	0	0
6320106	RPN6	9	13	25.58	9	13	2.30	1	4	1.84	1	2	0	0
6320934	PTC2	4	4	10.13	4	4	4.31	2	3	0	0	0	0	0
6322545	STE18	1	1	15.45	1	1	0	0	0	0	0	0	0	0
27808714	LAP3	1	1	3.08	1	1	0	0	0	0	0	0	0	0
6322410	IRC8	0	0	0	0	0	0	0	0	0	0	0	0	0
14318532	UBP6	7	9	17.03	7	9	9.02	3	5	2.61	1	11	0	0
6320129	BRE1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324287	COG6	0	0	0	0	0	0	0	0	0	0	0	0	0
6320846	NTF2	1	1	12.80	1	1	12.80	1	1	0	0	0	0	0
6324508	MET22	0	0	0	0	0	0	0	0	0	0	0	0	0
6319975	GCS1	13	55	53.69	13	55	55.97	17	78	0	0	0	0	0
63535562	CHA1	2	2	6.67	2	2	1.94	1	1	0	0	0	0	0
6322141	NEO1	1	1	1.56	1	1	0	0	0	0	0	0	0	0
398365175	CDC28	3	4	11.07	3	4	3.69	1	1	11.41	3	7	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6322378	TAX4	0	0	0	0	0	0	0	0	0	0	0	0	0
6320848	TIR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320093	TMA17	0	0	0	0	0	0	0	0	0	0	0	0	0
6322332	TRK1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320738	QCR7	0	0	0	0	0	0	0	0	0	0	0	0	0
6321543	VOA1	0	0	0	0	0	0	0	0	0	0	0	0	0
41629668	VPS8	0	0	0	0	0	0	0	0	0	0	0	0	0
6323150	YPS3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365307	EHT1	7	10	18.18	7	10	21.06	7	10	2.00	1	1	0	0
6322739	KTI12	5	9	19.17	5	9	3.51	1	1	3.51	1	1	0	0
398366117	ELP2	4	6	6.22	4	6	5.46	3	6	1.27	1	1	0	0
6324932	CIR2	3	4	5.86	3	4	1.43	1	1	1.43	1	1	0	0
6323129	ERG27	2	2	8.36	2	2	4.61	1	1	2.59	1	1	0	0
6322737	ABF1	2	2	3.83	2	2	3.69	2	3	1.50	1	1	0	0
6321278	AIM14	0	0	0	0	0	0	0	0	1.23	1	1	0	0
398364589	DSN1	0	0	0	0	0	0	0	0	1.39	1	1	0	0
6324870	YOR296W	0	0	0	0	0	0	0	0	0.78	1	1	0	0
398365441	ISN1	0	0	0	0	0	0	0	0	2.00	1	1	0	0
6320409	RAV2	0	0	0	0	0	0	0	0	3.99	1	1	0	0
398366501	MRPS2	2	3	11.89	2	3	0	0	0	3.15	1	2	0	0
398365469	RPL40B	0	0	0	0	0	0	0	0	13.46	1	2	0	0
398364921	TFC1	1	1	2.00	1	1	0	0	0	1.23	1	2	0	0
6320130	YDL073W	0	0	0	0	0	0	0	0	0.81	1	2	0	0
6321209	SHE10	4	4	7.11	4	4	2.08	1	1	3.81	2	3	0	0
6323662	FMS1	1	3	2.76	1	3	0	0	0	1.97	1	3	0	0
398365025	KCS1	1	1	0.76	1	1	0	0	0	0.86	1	3	0	0
398365489	LST8	0	0	0	0	0	0	0	0	3.63	1	3	0	0
6322593	XPT1	1	4	5.26	1	4	0	0	0	5.26	1	4	0	0
398366545	ARO10	9	14	15.28	9	14	4.88	3	5	2.52	2	5	0	0
398366635	PHO8	1	1	1.77	1	1	0	0	0	4.06	2	8	0	0
6321005	COG3	0	0	0	0	0	0	0	0	1.37	1	1	0	0
398364553	NBP35	1	1	3.05	1	1	0	0	0	0	0	0	0	0
6323569	COG8	0	0	0	0	0	2.97	1	1	0	0	0	0	0
6322229	GTT1	6	13	31.62	6	13	24.36	6	16	0	0	0	8.97	2
398366485	ASP1	5	8	14.96	5	8	7.87	2	3	0	0	0	0	0
398365871	MGE1	4	6	17.54	4	6	10.53	3	5	0	0	0	0	0
6324690	RPO31	3	3	3.08	3	3	1.99	2	3	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6324946	MRS6	1	1	2.32	1	1	0	0	0	0	0	0	0	0
6325052	HRR25	1	2	1.82	1	2	0	0	0	0	0	0	0	0
6324631	SGT1	1	1	2.78	1	1	3.04	1	1	0	0	0	0	0
6323651	ADI1	1	2	7.26	1	2	0	0	0	0	0	0	0	0
398365629	DBF2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320725	GRH1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324056	TOF1	1	1	1.05	1	1	0	0	0	0	0	0	0	0
6321823	PIH1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383767	KCC4	0	0	0	0	0	0	0	0	0	0	0	0	0
6319556	UBC4	0	0	0	0	0	0	0	0	0	0	0	0	0
6325007	GYP5	0	0	0	0	0	0	0	0	0	0	0	0	0
6322121	SEC6	0	0	0	0	0	0	0	0	0	0	0	0	0
6323292	TMA7	0	0	0	0	0	0	0	0	0	0	0	0	0
6325227	TRM44	4	5	8.11	4	5	2.29	1	1	3.17	1	1	0	0
6322526	TOR1	0	0	0	0	0	0	0	0	0.81	1	1	0	0
6324085	SUI1	6	28	70.37	6	28	62.04	6	30	0	0	0	0	0
398366199	IVY1	3	3	6.62	3	3	0	0	0	0	0	0	0	0
6322839	ARC19	2	4	14.04	2	4	9.36	1	1	0	0	0	0	0
6321914	CIA2	1	2	5.19	1	2	0	0	0	0	0	0	0	0
6319737	RGD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322419	MHP1	1	1	0.50	1	1	0	0	0	0	0	0	0	0
398366245	LDB19	0	0	0	0	0	0	0	0	0	0	0	0	0
6325011	YPL245W	0	0	0	0	0	0	0	0	0	0	0	0	0
398364697	LSM4	2	2	10.70	2	2	14.44	3	6	8.56	2	3	0	0
6320840	FMP52	4	6	18.61	4	6	12.55	2	3	0	0	0	30.74	5
6324794	RCN2	1	1	4.53	1	1	13.21	2	3	0	0	0	0	0
6321800	VPS29	2	2	8.16	2	2	0	0	0	0	0	0	0	0
398364967	SOK1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320413	UME6	0	0	0	0	0	0	0	0	0	0	0	0	0
6322379	IML2	1	1	1.23	1	1	0	0	0	0	0	0	0	0
6325362	COG4	0	0	0	0	0	0	0	0	2.90	2	2	0	0
6322156	BCY1	5	6	11.06	5	6	16.83	6	10	0	0	0	0	0
6322424	SNX4	1	1	3.78	1	1	0	0	0	0	0	0	0	0
398365471	MCM7	1	1	1.54	1	1	1.18	1	1	0	0	0	0	0
398365269	SMY2	0	0	0	0	0	2.43	1	1	3.24	1	1	0	0
6323616	YML6	1	1	2.45	1	1	0	0	0	0	0	0	0	0
6325343	SUA7	2	2	8.70	2	2	4.93	1	1	2.90	1	1	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
10383809	FIG2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324293	YNL035C	0	0	0	0	0	0	0	0	0	0	0	0	0
398366365	EFG1	3	4	19.74	3	4	6.44	1	3	6.87	1	2	0	0
6321235	KEX1	0	0	0	0	0	0	0	0	1.37	1	6	0	0
6323606	AMD1	6	8	8.40	6	8	7.04	3	3	0	0	0	0	0
398366153	MBF1	2	5	17.88	2	5	7.95	1	2	0	0	0	9.93	1
398364537	VPS45	3	4	6.41	3	4	0	0	0	3.64	2	5	0	0
6325326	SPE3	1	1	4.44	1	1	7.17	2	2	0	0	0	4.44	1
37362654	RRP46	0	0	0	0	0	0	0	0	0	0	0	0	0
6324194	FPR1	5	20	46.49	5	20	56.14	5	21	0	0	0	0	0
46562125	VPS5	1	1	1.63	1	1	0	0	0	1.33	1	5	0	0
6322759	YKL091C	3	7	10.32	3	7	5.81	2	19	0	0	0	0	0
37362671	MIA40	2	4	9.68	2	4	6.95	2	10	0	0	0	3.97	1
6320662	GUK1	2	6	12.30	2	6	25.67	4	12	0	0	0	0	0
6324363	ARC35	3	3	10.23	3	3	5.85	2	4	0	0	0	0	0
398366563	HPT1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366275	LSC2	2	2	6.09	2	2	0	0	0	0	0	0	0	0
6323739	AIP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325215	SSN3	0	0	0	0	0	0	0	0	0	0	0	0	0
6324378	LYS9	0	0	0	0	0	3.36	1	1	3.36	1	1	0	0
6324475	WRS1	4	8	11.57	4	8	15.28	5	8	2.31	1	2	4.17	1
6323279	SSP120	0	0	0	0	0	0	0	0	5.13	1	4	0	0
6322992	PRP19	5	5	12.52	5	5	4.37	2	5	0	0	0	0	0
330443401	MAP2	2	2	6.89	2	2	3.09	1	1	0	0	0	0	0
6324564	RCL1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324379	BRE5	0	0	0	0	0	0	0	0	0	0	0	0	0
6320849	PRE10	0	0	0	0	0	0	0	0	0	0	0	0	0
6325173	BRO1	1	2	1.30	1	2	4.38	3	3	1.54	1	1	0	0
398366247	PHO81	0	0	0	0	0	0	0	0	0.85	1	4	0	0
6323007	SPA2	6	7	4.64	6	7	6.68	6	16	0	0	0	0	0
6325447	SKI3	1	1	0.77	1	1	0	0	0	0	0	0	0	0
6323332	HRI1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324574	RRP6	2	2	3.27	2	2	0	0	0	0	0	0	0	0
6324031	CLA4	0	0	0	0	0	0	0	0	0	0	0	0	0
398365831	PRE9	4	4	17.83	4	4	0	0	0	3.10	1	1	0	0
6323142	HOG1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324615	CUE5	4	4	11.19	4	4	3.89	1	3	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6322153	CST6	2	3	3.41	2	3	0	0	0	1.36	1	1	5.28	1
14318524	NIC96	2	3	2.26	2	3	2.38	2	2	1.31	1	1	0	0
6324402	AIF1	3	4	9.26	3	4	4.50	1	2	2.38	1	1	0	0
398364507	DOT5	2	2	11.16	2	2	4.19	1	2	6.05	1	1	0	0
398365867	ENT5	1	1	2.92	1	1	2.43	1	1	2.92	1	1	0	0
6324130	GCR2	1	1	1.50	1	1	0	0	0	2.25	1	1	5.24	1
6323148	SRN2	2	2	10.80	2	2	0	0	0	6.57	1	1	0	0
6319738	TAE1	1	1	4.74	1	1	0	0	0	8.62	1	1	0	0
6324656	WHI5	0	0	0	0	0	0	0	0	3.73	1	1	0	0
6324781	RET1	0	0	0	0	0	0	0	0	0.78	1	1	0	0
37362668	YJL016W	0	0	0	0	0	0	0	0	1.43	1	1	0	0
6321609	PSD2	0	0	0	0	0	0	0	0	2.11	1	1	0	0
6321481	RME1	0	0	0	0	0	0	0	0	4.00	1	1	0	0
6325431	VPS4	2	2	5.49	2	2	0	0	0	1.83	1	2	0	0
6324854	FSH3	1	1	4.14	1	1	0	0	0	4.14	1	2	0	0
330443376	MAK16	1	1	4.58	1	1	0	0	0	2.61	1	2	0	0
6322690	RCN1	0	0	0	0	0	0	0	0	6.16	1	3	0	0
6322431	VPS53	0	0	0	0	0	0	0	0	1.22	1	3	0	0
6322631	SRY1	3	6	14.11	3	6	4.60	1	1	3.37	1	5	0	0
6320580	VPS74	6	30	23.48	6	30	26.96	7	44	0	0	0	0	0
9755336	DID4	2	7	9.91	2	7	3.88	1	1	0	0	0	0	0
6323881	YML44	1	6	11.22	1	6	16.33	1	4	0	0	0	0	0
398366509	MRP1	4	5	16.20	4	5	4.36	1	1	0	0	0	0	0
6320035	FAP7	3	4	18.27	3	4	12.18	2	3	0	0	0	0	0
6323354	SFH1	1	1	3.05	1	1	3.05	1	1	0	0	0	0	0
6323531	SEC65	2	3	8.06	2	3	8.79	2	2	0	0	0	0	0
6321433	COG7	3	3	11.47	3	3	7.89	2	2	0	0	0	0	0
398364865	PIB2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320673	RMT2	2	2	4.37	2	2	2.18	1	1	0	0	0	0	0
6323417	SWC7	2	2	19.70	2	2	37.88	4	5	0	0	0	0	0
6324423	DCP1	2	2	11.69	2	2	7.79	1	1	0	0	0	0	0
398365921	TAF10	1	1	3.88	1	1	18.45	2	2	0	0	0	9.22	1
6323712	UBX4	1	1	3.13	1	1	0	0	0	0	0	0	0	0
6325166	GLR1	1	2	2.48	1	2	2.48	1	1	0	0	0	0	0
6320994	LSM5	2	2	26.88	2	2	17.20	1	3	0	0	0	0	0
398366515	TRP4	2	2	4.47	2	2	0	0	0	0	0	0	0	0
6324551	TSR4	1	1	4.17	1	1	8.33	2	3	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6324068	ORC5	0	0	0	0	0	0	0	0	0	0	0	0	0
6325213	NOP4	2	2	2.48	2	2	2.48	1	1	0	0	0	0	0
6324041	CAF40	0	0	0	0	0	0	0	0	0	0	0	0	0
6320086	CYK3	0	0	0	0	0	0	0	0	0	0	0	0	0
6321762	SNF6	0	0	0	0	0	0	0	0	0	0	0	0	0
398365087	GRX4	0	0	0	0	0	6.97	1	1	0	0	0	0	0
6323218	ATG26	0	0	0	0	0	0	0	0	0	0	0	0	0
41629693	PRP18	0	0	0	0	0	0	0	0	0	0	0	0	0
330443528	PTP3	0	0	0	0	0	0	0	0	0	0	0	0	0
398364733	YIR035C	1	1	3.54	1	1	0	0	0	0	0	0	0	0
6321766	AIM17	0	0	0	0	0	0	0	0	0	0	0	0	0
14318483	MOB2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324527	PSK2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365421	PRP16	0	0	0	0	0	0.84	1	1	0	0	0	0	0
41629685	RSC30	0	0	0	0	0	0	0	0	0	0	0	0	0
6320907	VHR2	1	1	2.97	1	1	0	0	0	0	0	0	0	0
398366405	GCN2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364549	GYP6	0	0	0	0	0	0	0	0	0	0	0	0	0
330443696	YNL193W	0	0	0	0	0	0	0	0	0	0	0	0	0
6325243	YPL014W	0	0	0	0	0	0	0	0	0	0	0	0	0
6325327	MED1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320160	MTF2	0	0	0	0	0	0	0	0	0	0	0	0	0
6319287	PRP45	0	0	0	0	0	0	0	0	0	0	0	0	0
398365919	SEC5	0	0	0	0	0	0	0	0	0	0	0	0	0
6320433	SIR4	0	0	0	0	0	0.59	1	1	0	0	0	0	0
6321725	BIO2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324177	INN1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319534	ORC2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320887	SAP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321204	SEC15	0	0	0	0	0	0	0	0	0	0	0	0	0
6323111	SRL2	0	0	0	0	0	0	0	0	0	0	0	0	0
398366601	SSN2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323541	VPS9	0	0	0	0	0	0	0	0	0	0	0	0	0
6321955	YAP1801	0	0	0	0	0	0	0	0	1.41	1	1	0	0
6323215	EMG1	3	4	15.08	3	4	17.46	3	4	5.16	1	2	0	0
6319868	HTL1	0	0	0	0	0	14.10	1	2	11.54	1	2	0	0
6321537	MDR1	1	1	0.84	1	1	0	0	0	1.16	1	2	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6320533	YSP2	1	1	0.70	1	1	0	0	0	0	0	0	0	0
6323749	YMR102C	1	1	0.96	1	1	0	0	0	0	0	0	0	0
6324651	RTS2	1	1	4.31	1	1	0	0	0	0	0	0	0	0
6323005	POM33	0	0	2.51	1	1	0	0	0	0	0	0	0	0
6322036	GUT2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365235	POP7	0	0	0	0	0	0	0	0	0	0	0	0	0
6321629	HIP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323062	SMF3	0	0	0	0	0	0	0	0	0	0	0	0	0
6320619	DFM1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366299	YGR250C	0	0	0	0	0	0	0	0	0	0	0	0	0
73858744	YIL156W-	0	0	0	0	0	0	0	0	0	0	0	0	0
6322672	LST4	0	0	0	0	0	0	0	0	0	0	0	0	0
6323224	NMT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325414	TPO3	0	0	0	0	0	0	0	0	0	0	0	0	0
9755339	TAD3	0	0	0	0	0	0	0	0	0	0	0	0	0
6320031	UGA3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323713	AVO2	0	0	0	0	0	0	0	0	0	0	0	0	0
269970308	SSY5	0	0	0	0	0	1.43	1	1	0	0	0	1.43	1
6323127	CHA4	0	0	0	0	0	0	0	0	0	0	0	2.16	1
6319498	SCO2	0	0	0	0	0	0	0	0	0	0	0	4.65	1
398364781	VFA1	0	0	0	0	0	0	0	0	0	0	0	7.39	1
269970305	ADH4	0	0	0	0	0	4.45	1	2	0	0	0	0	0
269970311	ASK1	0	0	0	0	0	6.51	1	2	0	0	0	0	0
6321833	DOG2	0	0	0	0	0	8.13	1	2	0	0	0	0	0
42759857	NBL1	0	0	0	0	0	23.29	2	2	0	0	0	0	0
398365807	REI1	0	0	0	0	0	3.82	1	2	0	0	0	0	0
398366469	SRB7	0	0	0	0	0	12.14	1	2	0	0	0	0	0
6323627	TRM9	0	0	0	0	0	5.38	1	2	0	0	0	0	0
6319933	YCRO90C	0	0	0	0	0	4.40	1	1	0	0	0	0	0
398364703	YNL134C	0	0	0	0	0	3.19	1	1	0	0	0	0	0
6321870	IRE1	0	0	0	0	0	1.17	1	1	0	0	0	0	0
398365167	ARO3	0	0	0	0	0	4.32	1	1	0	0	0	0	0
6322656	ACP1	0	0	0	0	0	12.80	1	1	0	0	0	0	0
398364449	CEG1	0	0	0	0	0	4.14	1	1	0	0	0	0	0
21450619	DAD3	0	0	0	0	0	9.57	1	1	0	0	0	0	0
398364901	MED2	0	0	0	0	0	2.32	1	1	0	0	0	0	0
6323220	PEX13	0	0	0	0	0	3.63	1	1	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1_ _WT_1	TAP_Rtr1_ _WT_1
398364729	LYS1	0	0	0	0	0	0	0	0	0	0	0	4.56	1
6322800	SFK1	0	0	0	0	0	0	0	0	0	0	0	3.40	1
6320860	SRB4	0	0	0	0	0	0	0	0	0	0	0	2.47	1
398365591	TDP1	0	0	0	0	0	0	0	0	0	0	0	2.57	1
6321769	YHL018W	0	0	0	0	0	9.17	1	1	0	0	0	0	0
6321830	BCD1	0	0	0	0	0	4.10	1	1	0	0	0	0	0
398366341	CIN1	0	0	0	0	0	0.79	1	1	0	0	0	0	0
6323894	DFG5	0	0	0	0	0	2.84	1	1	0	0	0	0	0
398366533	DXO1	0	0	0	0	0	1.81	1	1	0	0	0	0	0
398366665	EMI1	0	0	0	0	0	9.09	1	1	0	0	0	0	0
6320903	FCY22	0	0	0	0	0	1.70	1	1	0	0	0	0	0
6324540	MSE1	0	0	0	0	0	4.29	1	1	0	0	0	0	0
6323006	HIF1	0	0	0	0	0	5.19	1	1	0	0	0	0	0
6319899	YCR051W	0	0	0	0	0	5.86	1	1	0	0	0	0	0
6320512	YDR306C	0	0	0	0	0	3.56	1	1	0	0	0	0	0
6321402	YGL036W	0	0	0	0	0	1.43	1	1	0	0	0	0	0
6322314	YJL147C	0	0	0	0	0	1.83	1	1	0	0	0	0	0
6324499	YOL073C	0	0	0	0	0	3.11	1	1	0	0	0	0	0
330443728	YOL075C	0	0	0	0	0	0.93	1	1	0	0	0	0	0
6323268	LIP2	0	0	0	0	0	5.79	1	1	0	0	0	0	0
6322349	MDV1	0	0	0	0	0	2.24	1	1	0	0	0	0	0
398365753	RSM19	0	0	0	0	0	10.99	1	1	0	0	0	0	0
6322678	YML38	0	0	0	0	0	11.59	1	1	0	0	0	0	0
6323715	MOT3	0	0	0	0	0	2.24	1	1	0	0	0	0	0
330443507	MPS1	0	0	0	0	0	3.40	1	1	0	0	0	0	0
6323816	MSS11	0	0	0	0	0	1.45	1	1	0	0	0	0	0
6323714	NAT4	0	0	0	0	0	4.56	1	1	0	0	0	0	0
6323913	PET111	0	0	0	0	0	1.00	1	1	0	0	0	0	0
6321514	PEX8	0	0	0	0	0	2.21	1	1	0	0	0	0	0
330443601	PFD1	0	0	0	0	0	13.76	1	1	0	0	0	0	0
14318559	PHO4	0	0	0	0	0	3.21	1	1	0	0	0	0	0
6325444	PZF1	0	0	0	0	0	3.26	1	1	0	0	0	0	0
6324200	NRK1	0	0	0	0	0	6.25	1	1	0	0	0	0	0
6323914	ROY1	0	0	0	0	0	1.99	1	1	0	0	0	0	0
6320250	RPC11	0	0	0	0	0	16.36	1	1	0	0	0	0	0
398366011	MDM3	0	0	0	0	0	2.51	1	1	0	0	0	0	0
6320089	YDL114W	0	0	0	0	0	6.17	1	1	0	0	0	0	0

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Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1_ _WT_1	TAP_Rtr1_ _WT_1
398365719	CLB6	0	0	0	0	0	0	0	0	0	0	0	0	0
6324674	CRC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321317	GPG1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324832	HNT3	0	0	0	0	0	0	0	0	0	0	0	0	0
6321177	YGL260W	0	0	0	0	0	0	0	0	0	0	0	0	0
6323877	YMR221C	0	0	0	0	0	0	0	0	0	0	0	0	0
6319408	KIP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324872	MUM3	0	0	0	0	0	0	0	0	0	0	0	0	0
6323136	REX3	0	0	0	0	0	0	0	0	0	0	0	0	0
398366181	SLI1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323658	SOK2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320871	ZRG8	0	0	0	0	0	0	0	0	0	0	0	0	0
6324461	MDY2	3	5	16.51	3	5	5.19	1	3	5.19	1	1	0	0
6324739	SEY1	1	1	1.68	1	1	0	0	0	1.29	1	1	0	0
398365665	YNR021W	1	1	4.21	1	1	4.21	1	2	1.98	1	1	0	0
6322465	POL31	1	1	2.26	1	1	2.26	1	1	4.11	1	1	0	0
6323610	NDC1	0	0	0	0	0	0	0	0	1.83	1	1	0	0
6323736	YTA12	1	1	2.06	1	1	0	0	0	2.30	1	1	0	0
6323671	FAR8	1	1	1.34	1	1	2.68	1	1	1.72	1	1	0	0
6319503	CDS1	1	1	1.97	1	1	0	0	0	2.63	1	1	0	0
398366089	MPD1	1	1	3.46	1	1	0	0	0	2.52	1	1	0	0
6321373	ALG2	1	1	2.58	1	1	0	0	0	2.58	1	1	0	0
6325397	LOA1	1	1	5.33	1	1	0	0	0	3.00	1	1	0	0
116006497	MRPL15	0	0	0	0	0	3.56	1	1	4.74	1	1	0	0
330443583	ERG7	1	1	2.05	1	1	0	0	0	1.64	1	1	0	0
6324069	LTO1	1	1	5.05	1	1	0	0	0	6.57	1	1	0	0
6324930	MSC6	1	1	1.59	1	1	0	0	0	1.88	1	1	0	0
6320901	PCL6	1	1	1.90	1	1	0	0	0	2.62	1	1	0	0
398365081	SPC34	1	1	4.75	1	1	0	0	0	5.08	1	1	0	0
6321878	NAM8	0	0	0	0	0	0	0	0	3.63	1	1	0	0
398364895	RAV1	0	0	0	0	0	0	0	0	0.96	1	1	0	0
42759860	TOR2	0	0	0	0	0	0	0	0	0.36	1	1	0	0
14270687	AST1	0	0	0	0	0	2.33	1	1	1.63	1	1	0	0
398364431	AGE2	0	0	0	0	0	0	0	0	2.68	1	1	0	0
398365583	GAC1	0	0	0	0	0	0	0	0	1.77	1	1	0	0
6324345	RCF2	0	0	0	0	0	0	0	0	4.91	1	1	0	0
6322351	GZF3	0	0	0	0	0	0	0	0	2.18	1	1	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
330443362	BDH1	0	0	0	0	0	0	0	0	2.36	1	1	0	0
6325377	CLB5	0	0	0	0	0	0	0	0	1.84	1	1	0	0
6324301	CRZ1	0	0	0	0	0	0	0	0	1.33	1	1	0	0
6321426	ERG4	0	0	0	0	0	0	0	0	1.69	1	1	0	0
6322803	ELM1	0	0	0	0	0	0	0	0	1.41	1	1	0	0
398366135	FLD1	0	0	0	0	0	0	0	0	5.26	1	1	0	0
6321381	GEP7	0	0	0	0	0	0	0	0	2.44	1	1	0	0
6325242	HST2	0	0	0	0	0	0	0	0	3.64	1	1	0	0
330443455	YBR138C	0	0	0	0	0	0	0	0	1.72	1	1	0	0
6320002	YDL199C	0	0	0	0	0	0	0	0	4.66	1	1	0	0
6320294	YDR089W	0	0	0	0	0	0	0	0	1.38	1	1	0	0
6322972	YLL056C	0	0	0	0	0	0	0	0	2.68	1	1	0	0
6323358	YLR326W	0	0	0	0	0	0	0	0	6.25	1	1	0	0
6325189	YPL068C	0	0	0	0	0	0	0	0	7.85	1	1	0	0
398366259	KEL2	0	0	0	0	0	0	0	0	0.91	1	1	0	0
6324755	LAS17	0	0	0	0	0	0	0	0	1.42	1	1	0	0
6323686	MCM1	0	0	0	0	0	0	0	0	3.85	1	1	0	0
6321441	PEX31	0	0	0	0	0	0	0	0	3.46	1	1	0	0
6324436	PFK27	0	0	0	0	0	0	0	0	2.52	1	1	0	0
398366391	PIP2	0	0	0	0	0	0	0	0	0.90	1	1	0	0
6323238	PNP1	0	0	0	0	0	0	0	0	2.25	1	1	0	0
6319818	RRP7	0	0	0	0	0	0	0	0	3.03	1	1	0	0
6323655	SEC59	0	0	0	0	0	0	0	0	1.54	1	1	0	0
6324650	SKI7	0	0	0	0	0	0	0	0	2.01	1	1	0	0
6324847	TPO4	0	0	0	0	0	0	0	0	1.21	1	1	0	0
398366289	RMD5	0	0	0	0	0	0	0	0	2.85	1	1	0	0
398364689	YVH1	2	2	6.32	2	2	18.41	4	10	7.14	2	2	0	0
6324543	GAS5	3	4	8.88	3	4	8.26	3	3	2.48	1	2	0	0
6322303	CIS3	2	2	7.05	2	2	3.52	1	1	3.52	1	2	0	0
6325237	ULP1	2	3	3.70	2	3	0	0	0	1.29	1	2	0	0
116006493	HSP150	2	2	10.90	2	2	1.94	1	1	1.94	1	2	0	0
6226534	VAR1	1	2	2.76	1	2	0	0	0	2.51	1	2	0	0
6321942	IMP3	1	1	8.20	1	1	0	0	0	4.37	1	2	0	0
6320857	SBH2	1	1	14.77	1	1	0	0	0	14.77	1	2	0	0
6321909	TOM71	0	0	0	0	0	0	0	0	3.13	2	2	0	0
6321566	YGR127W	1	1	2.56	1	1	0	0	0	2.56	1	2	0	0
6323942	YML33	1	1	16.28	1	1	0	0	0	13.95	1	2	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP	# PSM	%SC	UP
		BY4741_F LAG_J1	BY4741_F LAG_J1	BY4741_FL AG_J2	BY4741_F LAG_J2	BY4741_F LAG_J2	BY4741_FL AG_J3	BY4741_F LAG_J3	BY4741_F LAG_J3	V5_Rtr1_ WT	V5_Rtr1_ WT	V5_Rtr1_ WT	TAP_Rtr1 _WT_1	TAP_Rtr1 _WT_1
6325075	CTI6	0	0	0	0	0	0	0	0	2.57	1	2	0	0
6323612	USA1	0	0	0	0	0	0	0	0	1.55	1	2	0	0
6324845	FSF1	0	0	0	0	0	0	0	0	3.98	1	2	0	0
398365295	YDR056C	0	0	0	0	0	0	0	0	6.83	1	2	0	0
6324510	APM4	0	0	0	0	0	0	0	0	2.04	1	2	0	0
6323263	TOP3	0	0	0	0	0	0	0	0	1.52	1	2	0	0
41629680	FMP21	0	0	0	0	0	0	0	0	5.80	1	2	0	0
6319813	GFD2	0	0	0	0	0	0	0	0	1.59	1	2	0	0
6324214	YNL115C	0	0	0	0	0	0	0	0	2.02	1	2	0	0
398364237	RML2	0	0	0	0	0	0	0	0	3.82	2	2	0	0
398365189	OCA2	0	0	0	0	0	0	0	0	5.08	1	2	0	0
398366517	SPC110	0	0	0	0	0	0	0	0	1.17	1	2	0	0
6323897	YHM2	0	0	0	0	0	0	0	0	2.55	1	3	0	0
6323573	ERV41	0	0	0	0	0	0	0	0	2.27	1	3	0	0
6320503	SUR2	0	0	0	0	0	0	0	0	5.44	2	3	0	0
6319643	TYR1	0	0	0	0	0	0	0	0	2.21	1	3	0	0
398364575	SBH1	0	0	0	0	0	0	0	0	13.41	1	3	0	0
6320189	TSC13	0	0	0	0	0	0	0	0	3.55	2	4	0	0
6324066	YIF1	0	0	0	0	0	0	0	0	4.46	1	4	0	0
6321596	CHO2	0	0	0.81	1	1	0	0	0	2.19	2	5	0	0
6321331	RMD9	0	0	0	0	0	0	0	0	4.02	2	5	0	0
6323745	ATP25	0	0	0	0	0	0	0	0	2.29	1	5	0	0
6321861	TRM5	0	0	0	0	0	0	0	0	9.82	4	7	0	0
6321963	SBP8	3	3	6.50	3	3	0	0	0	3.02	1	9	0	0
6325448	RPC82	3	3	5.50	3	3	2.45	1	1	5.35	3	10	0	0
6325370	PIS1	0	0	0	0	0	0	0	0	5.00	1	10	0	0
6319804	EMC1	0	0	0	0	0	0	0	0	3.03	2	10	0	0
6322762	CAB3	0	0	0	0	0	0	0	0	2.10	1	12	0	0
99030930	ZIM17	2	12	13.22	2	12	24.71	2	3	13.22	2	13	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ _WT_2	# PSM TAP_Rtr1_ _WT_2	%SC TAP_Rtr1_ _WT_3	UP TAP_Rtr1_ _WT_3	# PSM TAP_Rtr1_ _WT_3	%SC TAP_Rtr1_ _WT_4	UP TAP_Rtr1_ _WT_4	# PSM TAP_Rtr1_ _WT_4	%SC TAP_Rtr1_ _ctk1_del etion	UP TAP_Rtr1_ _ctk1_del etion	# PSM TAP_Rtr1_ _ctk1_del etion
398364847	RTR1	3529	75.66	14	623	79.65	21	2834	53.54	15	2499	46.46	12	4513
6320061	RPB1	845	33.47	37	254	37.05	45	464	19.85	34	456	4.15	6	20
398365417	RPB2	537	31.94	29	125	35.38	33	292	26.39	28	133	5.23	6	16
398364469	RPB3	116	52.20	10	53	57.86	11	108	13.52	3	16	0	0	0
6322321	RPB4	8	35.75	4	7	0	0	0	0	0	0	0	0	0
6319630	RPB5	108	37.67	7	67	36.74	6	99	23.72	4	38	6.98	1	3
398366567	RPB7	0	0	0	0	0	0	0	0	0	0	0	0	0
398365775	RPB10	10	52.86	2	14	15.71	1	29	0	0	0	0	0	0
6325445	RPB6	35	26.45	2	15	26.45	2	10	0	0	0	0	0	0
6321368	RPB9	9	0	0	0	12.30	1	5	0	0	0	0	0	0
6324798	RPB8	21	10.96	1	19	31.51	4	24	10.96	1	1	0	0	0
6324569	RPB11	39	25.83	3	3	47.50	5	19	0	0	0	0	0	0
6323632	SPT5	0	2.73	2	4	7.90	6	18	1.22	1	1	0	0	0
6321625	TFG1	0	2.59	1	1	0	0	0	0	0	0	0	0	0
6321937	RPB12	67	28.57	1	3	40.00	2	35	28.57	1	3	0	0	0
398365043	TFG2	0	0	0	0	0	0	0	3.00	1	1	0	0	0
154199603	IWR1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365457	SPT4	0	0	0	0	0	0	0	0	0	0	0	0	0
6325128	TAF14	0	0	0	0	6.97	1	3	0	0	0	0	0	0
398365155	NPA3	40	6.23	2	8	20.00	7	28	2.60	1	1	0	0	0
6323272	GPN3	24	10.66	2	4	10.66	2	7	0	0	0	0	0	0
6322163	IRR1	0	1.04	1	1	1.22	1	3	0	0	0	0	0	0
398366099	RRP41	0	0	0	0	0	0	0	0	0	0	0	0	0
398366555	UBA2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323670	TAP42	0	0	0	0	17.76	4	8	0	0	0	0	0	0
6323004	SSA2	191	19.56	7	31	59.78	26	311	26.92	13	49	6.42	3	11
6319505	RPL4A	29	22.10	5	32	30.94	8	46	2.76	1	1	0	0	0
398364991	RPL4B	26	22.10	5	32	30.94	8	46	2.76	1	1	0	0	0
398365303	RPL43B	320	29.35	2	19	29.35	3	12	0	0	0	0	0	0
398364349	RPL2A	32	24.41	3	3	27.17	4	8	4.33	1	6	0	0	0
6325337	TEF1	71	28.17	7	37	55.68	18	1023	12.66	6	196	11.79	6	206
144228166	SSA1	206	16.20	5	20	62.15	28	308	28.35	14	51	8.26	4	21
6319559	RPL19A	25	20.11	3	5	14.29	2	5	0	0	0	0	0	0
6320065	RPL35A	170	0	0	0	11.67	1	3	0	0	0	0	0	0
398365603	RPS4A	28	3.83	1	1	24.14	4	21	6.51	2	2	3.83	1	5
398365321	RPS6B	163	0	0	0	13.56	2	10	15.25	3	4	11.44	2	13
398364655	SSA4	75	6.23	3	14	16.20	9	93	9.66	5	26	3.89	2	10

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ _WT_2	# PSM TAP_Rtr1_ _WT_2	%SC TAP_Rtr1_ _WT_3	UP TAP_Rtr1_ _WT_3	# PSM TAP_Rtr1_ _WT_3	%SC TAP_Rtr1_ _WT_4	UP TAP_Rtr1_ _WT_4	# PSM TAP_Rtr1_ _WT_4	%SC TAP_Rtr1_ _ctk1_del etion	UP TAP_Rtr1_ _ctk1_del etion	# PSM TAP_Rtr1_ _ctk1_del etion
398366603	RPS17B	350	26.47	3	9	33.82	4	12	8.82	1	3	11.76	1	1
6324533	RPS15	332	30.28	3	13	22.54	2	12	7.75	1	1	0	0	0
6323104	RPL10	7	4.98	1	5	18.10	2	11	0	0	0	0	0	0
6319396	SSA3	62	6.16	3	6	14.79	9	90	11.86	6	16	2.00	1	1
6321786	RPL14B	25	21.74	2	17	28.26	4	65	13.04	2	2	0	0	0
6322847	RPL14A	25	21.74	2	17	28.26	4	65	13.04	2	2	0	0	0
6324670	RPS7A	92	44.21	6	31	41.05	6	28	5.26	1	1	0	0	0
6322668	RPL17A	107	15.76	2	18	15.76	2	4	0	0	0	0	0	0
398365285	CAX4	0	0	0	0	0	0	0	0	0	0	0	0	0
6322284	RPL17B	91	15.76	2	18	15.76	2	4	0	0	0	0	0	0
398364883	RPS7B	142	21.58	4	7	30.53	5	23	5.26	1	1	0	0	0
6323791	RPL13B	52	13.07	1	2	15.08	3	13	15.58	3	5	5.53	1	2
6320121	RPL13A	49	13.07	1	2	15.08	3	13	15.58	3	5	5.53	1	2
6324120	SSB2	139	20.07	7	32	60.85	24	266	16.80	9	28	15.99	7	40
6323577	RPS1B	77	3.92	1	2	17.25	4	19	8.24	2	2	5.10	1	1
6325157	ATG21	0	0	0	0	0	0	0	0	0	0	7.66	1	1
6323474	RPS1A	62	3.92	1	2	17.25	4	19	3.14	1	1	0	0	0
398364725	RPL23B	78	25.55	2	5	35.04	5	27	10.95	1	1	0	0	0
6319972	SSB1	133	23.49	8	36	56.44	23	270	16.80	9	28	15.99	7	40
154200011	RPL20B	16	0	0	0	10.47	1	1	0	0	0	0	0	0
6322989	UBI4	24	0	0	0	32.89	2	78	32.89	2	34	21.05	1	1
9755341	RPS16A	88	18.18	2	3	39.86	5	23	0	0	0	0	0	0
6319378	RPL32	105	0	0	0	22.31	2	6	0	0	0	0	0	0
6325061	APL5	0	0	0	0	11.16	7	16	0	0	0	0	0	0
398364505	RPS3	53	0	0	0	42.08	7	15	5.42	1	1	12.50	2	13
398365351	RPS13	46	35.10	4	12	19.21	2	8	12.58	2	2	0	0	0
6324260	RPL16B	2	6.06	1	1	14.14	2	11	0	0	0	7.58	1	1
398365605	RPS30B	41	15.87	1	1	15.87	1	2	15.87	1	1	0	0	0
6323481	RPL6B	1	8.52	1	2	14.20	2	5	4.55	1	2	0	0	0
398365377	RPS9B	19	0	0	0	0	0	0	4.62	1	2	8.21	1	4
6324265	YDJ1	3	17.36	3	5	36.43	12	67	14.18	5	11	0	0	0
398364745	ATP7	14	14.37	1	4	14.37	1	1	0	0	0	0	0	0
6321464	RPS25A	150	31.48	3	7	0	0	0	0	0	0	0	0	0
6323077	RPS0B	17	5.16	1	2	37.70	6	26	5.16	1	1	0	0	0
398365071	RPS11A	9	7.05	1	2	0	0	0	5.13	1	1	0	0	0
6321653	RPS0A	17	5.16	1	2	37.70	6	26	5.16	1	1	0	0	0
6323465	CNA1	56	5.79	2	7	2.53	1	2	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6321335	RPL28	14	7.38	2	2	10.07	1	2	0	0	0	0	0	0
6322505	SSC1	32	2.75	1	1	31.80	13	64	0	0	0	0	0	0
6323567	RPL6A	1	8.52	1	2	8.52	1	4	4.55	1	2	0	0	0
6321754	RPL8A	45	4.30	1	2	34.77	7	20	12.11	3	8	4.30	1	1
6322984	RPL8B	47	4.30	1	2	34.77	7	20	12.11	3	8	4.30	1	1
6323615	RPS18B	18	8.22	1	1	27.40	4	6	15.07	2	3	7.53	1	1
398366307	RPA190	0	0	0	0	14.30	14	33	0	0	0	0	0	0
6324637	RPL3	0	0	0	0	2.07	1	3	2.07	1	1	0	0	0
6320128	RPL31A	13	28.32	3	29	14.16	1	11	0	0	0	0	0	0
398366143	RPL31B	13	20.35	2	28	14.16	1	11	0	0	0	0	0	0
398365761	RPS23A	2	7.59	1	3	7.59	1	2	0	0	0	0	0	0
6322137	RPL34B	1	14.05	1	2	0	0	0	0	0	0	0	0	0
6321315	RPS2	19	0	0	0	19.29	3	11	0	0	0	0	0	0
6320949	RPS8B	13	6.50	1	1	35.50	5	15	0	0	0	0	0	0
6321798	RPL27A	2	17.65	2	5	25.74	3	12	10.29	1	1	16.91	2	8
6325176	RPS9A	18	0	0	0	0	0	0	4.57	1	2	0	0	0
86558907	RHR2	46	0	0	0	52.00	10	129	0	0	0	0	0	0
6322058	RPL16A	0	0	0	0	6.03	1	1	0	0	0	0	0	0
6324027	RPS19B	291	18.06	2	13	31.94	3	44	6.25	1	4	6.25	1	3
6323090	RPL22A	48	18.18	1	6	18.18	1	23	0	0	0	0	0	0
6320142	RPS29B	9	0	0	0	14.29	1	2	0	0	0	0	0	0
6324807	KIN4	0	0	0	0	0	0	0	0	0	0	0	0	0
6321522	RPL11B	22	0	0	0	7.47	1	2	0	0	0	0	0	0
6322583	RPS5	82	6.67	1	9	20.89	4	56	4.89	1	2	4.89	1	1
6325126	RPL5	4	5.72	1	1	8.75	2	6	3.03	1	1	0	0	0
6323073	PDC1	42	19.72	6	16	36.41	12	62	1.78	1	1	0	0	0
398366277	VMA4	44	6.44	1	2	41.20	8	51	20.17	4	6	0	0	0
6323824	HOT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325066	NAB3	0	0	0	0	0	0	0	0	0	0	0	0	0
398364517	RPS24A	47	18.52	2	8	24.44	3	6	6.67	1	1	0	0	0
6321362	RPL7A	17	13.11	2	11	27.05	5	14	0	0	0	0	0	0
6325058	RPL7B	17	13.11	2	11	27.05	5	14	0	0	0	0	0	0
37362649	GUS1	1	0	0	0	23.31	12	45	13.28	9	37	1.55	1	1
398365909	RPL24B	2	9.03	1	1	0	0	0	0	0	0	0	0	0
6324707	EFT1	32	10.81	5	9	53.92	31	257	2.85	2	2	1.66	1	2
6323376	RPL26A	68	18.90	2	7	6.30	1	1	7.09	1	1	6.30	1	14
116006491	RPL26B	77	18.90	2	7	6.30	1	1	7.09	1	1	6.30	1	14

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6325006	RPL36B	116	0	0	0	19.00	2	5	0	0	0	0	0	0
6323848	RPL36A	116	0	0	0	19.00	2	5	0	0	0	0	0	0
398366579	RPL12B	14	0	0	0	28.48	3	7	9.09	1	2	0	0	0
6321540	NOP7	0	0	0	0	7.60	3	6	1.65	1	1	0	0	0
398365599	PDC6	15	3.37	1	2	3.37	1	3	0	0	0	0	0	0
6323163	PDC5	15	3.37	1	2	5.15	2	4	0	0	0	0	0	0
6323583	CMP2	23	3.81	1	2	0	0	0	1.82	1	1	0	0	0
398365991	CLC1	28	6.87	1	3	6.87	1	1	0	0	0	0	0	0
37362619	RXT2	0	0	0	0	3.49	1	2	0	0	0	0	0	0
10383793	RPS14A	59	8.03	1	3	51.82	6	25	0	0	0	9.49	1	9
6322271	RPS22A	13	12.31	1	6	19.23	2	15	17.69	2	3	0	0	0
6321407	RPL24A	3	9.03	1	1	0	0	0	0	0	0	0	0	0
6323357	RPL38	48	33.33	2	6	19.23	1	23	0	0	0	0	0	0
398366183	HTB1	5	0	0	0	18.32	2	10	0	0	0	11.45	1	3
398364287	GCN1	0	0	0	0	18.90	39	116	0	0	0	0	0	0
6320016	VMA1	3	0	0	0	23.81	18	89	0	0	0	1.77	2	30
14318481	TUB2	6	0	0	0	42.67	13	62	0	0	0	0	0	0
398364585	KAR2	19	0	0	0	15.84	7	27	1.76	1	13	1.76	1	9
6323196	RPS31	0	0	0	0	0	0	0	0	0	0	0	0	0
6321013	PAB1	132	31.02	10	57	33.10	10	42	12.82	6	19	6.76	3	23
6324993	FUM1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366075	RPS29A	13	0	0	0	0	0	0	0	0	0	0	0	0
6322755	YJU2	0	0	0	0	3.96	1	2	0	0	0	0	0	0
398366059	PLP2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320718	SMT3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365031	YBR139W	1	0	0	0	13.39	4	14	0	0	0	0	0	0
6321850	FYV4	0	0	0	0	0	0	0	0	0	0	0	0	0
398366083	TDH3	286	28.61	5	15	66.27	14	354	14.16	5	9	4.52	1	28
6319279	CDC19	107	3.60	1	1	65.40	26	504	21.80	9	14	32.80	12	206
6322468	TDH2	257	15.06	3	7	46.99	12	245	14.16	5	9	4.52	1	28
6321249	RPS26A	23	5.88	1	1	12.61	1	14	0	0	0	12.61	1	3
6321772	RPS20	1	0	0	0	22.31	3	11	0	0	0	29.75	3	27
14318474	SEC53	1	10.24	1	1	14.17	3	5	0	0	0	0	0	0
6320589	YRA1	33	0	0	0	5.75	1	4	18.58	4	9	0	0	0
6320805	ECM10	3	2.80	1	1	3.73	2	6	0	0	0	0	0	0
6323776	ECM16	0	0	0	0	3.55	3	4	0	0	0	0	0	0
6320596	RVS167	0	4.15	1	1	32.99	10	87	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
398366359	SLH1	0	0	0	0	1.63	2	3	0	0	0	0	0	0
6324321	SIS1	10	0	0	0	12.50	3	9	9.38	3	5	7.39	2	7
6319447	NCL1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323112	EMP70	0	0	0	0	3.90	1	1	0	0	0	0	0	0
6324595	SFM1	0	0	0	0	8.92	1	6	0	0	0	0	0	0
330443411	UBP13	0	0	0	0	6.83	3	12	0	0	0	0	0	0
6323868	EFR3	0	0	0	0	0	0	0	0	0	0	0	0	0
330443676	CDC34	0	0	0	0	17.12	5	36	0	0	0	0	0	0
6320615	TRS120	0	0	0	0	0	0	0	0	0	0	0	0	0
6323487	FMP27	0	0	0	0	0	0	0	0	0	0	0	0	0
6320231	REG1	0	0	0	0	0.89	1	1	0	0	0	0	0	0
6324585	AUS1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325048	RKM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323732	SEG1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364779	YGL041W-	6	13.64	1	2	13.64	1	2	0	0	0	0	0	0
398365827	CAF130	0	0	0	0	1.07	1	1	0	0	0	0	0	0
6325063	RSA1	0	0	0	0	3.15	1	1	0	0	0	0	0	0
398365559	MSH6	0	0	0	0	1.05	1	1	0	0	0	0	0	0
398365727	YNR034W-	0	0	0	0	0	0	0	0	0	0	0	0	0
6323683	YET2	0	0	0	0	0	0	0	0	0	0	0	0	0
330443366	GPB2	0	0	0	0	0	0	0	0	0	0	0	0	0
6325012	HUT1	0	0	0	0	0	0	0	0	0	0	0	0	0
99030927	KKQ8	0	0	0	0	0	0	0	0	0	0	0	0	0
398364567	MAD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320763	RMD6	0	0	0	0	0	0	0	0	0	0	0	0	0
398365009	MEC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364407	RPL9A	18	0	0	0	23.04	3	16	6.28	1	1	0	0	0
6324298	HHF2	22	0	0	0	33.01	3	11	29.13	3	9	0	0	0
6324262	RPL9B	7	0	0	0	23.04	3	8	6.28	1	1	0	0	0
398364523	TDH1	86	4.22	1	1	48.19	11	182	16.27	5	12	4.52	1	28
398364951	VMA2	31	10.06	3	9	66.54	26	515	5.80	3	4	0	0	0
6321333	ARC1	0	0	0	0	25.00	6	40	8.24	3	10	0	0	0
6321408	RPL30	31	16.19	1	1	45.71	3	19	0	0	0	0	0	0
6324445	RPL25	12	0	0	0	19.01	2	3	9.15	1	14	0	0	0
6325114	RPL33A	0	0	0	0	24.30	2	5	0	0	0	0	0	0
398365881	RPL33B	0	0	0	0	24.30	2	4	0	0	0	0	0	0
398364339	THS1	12	0	0	0	5.31	3	6	6.27	4	7	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6320637	TIF35	0	0	0	0	13.50	2	4	0	0	0	0	0	0
398364223	NPR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6325390	SPN1	1	0	0	0	16.83	5	25	0	0	0	0	0	0
6324578	DNL4	0	0	0	0	0	0	0	0	0	0	0	0	0
6319291	FRT2	0	0	0	0	0	0	0	0	0	0	0	0	0
6322242	HXT9	0	0	0	0	0	0	0	0	0	0	0	0	0
6322272	RPL39	0	0	0	0	0	0	0	0	0	0	0	0	0
14318518	VTC2	0	0	0	0	1.93	1	1	0	0	0	0	0	0
330443752	THI22	0	0	0	0	0	0	0	0	0	0	0	0	0
6325175	MOT1	0	0	0	0	1.50	2	3	0	0	0	0	0	0
6322796	DEF1	3	0	0	0	0	0	0	0	0	0	4.34	1	1
398365059	PTK2	0	0	0	0	0	0	0	0.98	1	1	0	0	0
6320345	DOP1	0	0	0	0	1.06	2	3	0	0	0	0	0	0
6323863	HFA1	0	0	0	0	1.04	2	7	0	0	0	0	0	0
6322948	MLP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322999	FRA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324491	IRA2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324215	RPC19	2	0	0	0	8.45	1	1	0	0	0	0	0	0
6320419	UPC2	0	0	0	0	1.64	1	1	0	0	0	0	0	0
398364627	RDH54	0	0	0	0	2.09	2	2	0	0	0	0	0	0
6321752	VMR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321793	TIM10	0	0	0	0	0	0	0	0	0	0	0	0	0
6322032	BNR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325074	RTT10	0	0	0	0	0	0	0	0	0	0	0	0	0
6325235	RAD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323869	CEF1	0	0	0	0	3.39	1	1	0	0	0	1.53	1	1
6323917	TPS3	0	0	0	0	0	0	0	0	0	0	0	0	0
6321552	SPT6	0	0	0	0	4.27	5	15	0	0	0	0	0	0
6320853	BIM1	0	0	0	0	6.98	2	8	0	0	0	0	0	0
398364243	YEL043W	0	0	0	0	0	0	0	0	0	0	0	0	0
6320027	DLD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323356	PEX30	0	0	0	0	0	0	0	0	0	0	0	0	0
6320799	ANP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324393	YNR065C	0	0	0	0	0	0	0	0	0	0	0	0	0
6323266	THI7	0	2.51	1	1	0	0	0	0	0	0	0	0	0
6323445	YLR413W	0	0	0	0	0	0	0	0	0	0	0	0	0
398364667	THO2	0	0	0	0	0.50	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
37362637	JIP4	0	0	0	0	0	0	0	0	0	0	0	0	0
398364707	AAC3	2	0	0	0	3.91	1	1	0	0	0	0	0	0
6322570	YMR1	0	0	0	0	3.78	2	2	0	0	0	0	0	0
6324879	ISW2	0	0	0	0	5.09	5	14	0.80	1	1	0	0	0
10383772	LEU2	1	0	0	0	6.32	1	1	0	0	0	0	0	0
6322079	HOS4	0	0	0	0	1.57	1	1	0	0	0	0	0	0
6321210	VID30	0	0	0	0	2.40	1	1	0	0	0	0	0	0
398365701	YNR029C	0	0	0	0	5.83	2	6	0	0	0	0	0	0
6320692	VPS52	0	0	0	0	4.21	2	5	0	0	0	0	0	0
398365525	SMD1	0	0	0	0	17.12	2	3	0	0	0	0	0	0
6322086	SLM1	2	0	0	0	6.85	3	3	0	0	0	0	0	0
398364897	CBP6	5	0	0	0	0	0	0	0	0	0	0	0	0
6323726	NAM7	0	0	0	0	0	0	0	0	0	0	0	0	0
6324982	SAM4	0	0	0	0	0	0	0	0	0	0	0	0	0
6320324	VBA4	0	0	0	0	0	0	0	0	0	0	0	0	0
6324612	HIR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324981	SAM3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325379	AXL1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321020	BRR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324694	GCY1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322312	DAS1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364625	YKL033W-	0	0	0	0	4.66	1	2	0	0	0	0	0	0
6320034	NRP1	0	0	0	0	15.86	5	6	0	0	0	0	0	0
6325374	YPR117W	0	0	0	0	0	0	0	0	0	0	0	0	0
6324546	MDM38	2	0	0	0	0	0	0	0	0	0	0	0	0
6323289	LCB5	0	0	0	0	0	0	0	0	0	0	0	0	0
6321205	TAN1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319351	PHO11	0	0	0	0	0	0	0	0	0	0	0	0	0
6319523	REB1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320921	YER077C	0	0	0	0	0	0	0	2.76	1	1	0	0	0
6320826	GLC3	0	0	0	0	0	0	0	3.41	1	1	0	0	0
398365141	INP53	0	0	0	0	1.26	1	3	0	0	0	0	0	0
398366583	SIP1	0	0	0	0	1.84	1	1	0	0	0	0	0	0
398364669	MAD3	0	0	0	0	3.88	1	2	0	0	0	0	0	0
6322700	DBR1	0	0	0	0	2.72	1	1	0	0	0	0	0	0
6323317	YLR287C	0	0	0	0	3.66	1	1	0	0	0	0	0	0
398365819	SAN1	0	0	0	0	3.28	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
398365335	YNL033W	0	0	0	0	0	0	0	0	0	0	0	0	0
33438893	YPL119C-	0	0	0	0	0	0	0	0	0	0	0	0	0
6322579	JHD2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324141	KAR1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383756	LRE1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364917	NHP10	0	0	0	0	0	0	0	0	0	0	0	0	0
6323035	NSE1	0	0	0	0	0	0	0	0	0	0	0	0	0
50593425	OMA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321910	ORC6	0	0	0	0	0	0	0	0	0	0	0	0	0
6322494	PET191	0	0	0	0	0	0	0	0	0	0	0	0	0
398365057	SPO14	0	0	0	0	0	0	0	0	0	0	0	0	0
398364971	TRP1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365003	PSF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321697	RAD2	0	0	0	0	0	0	0	0	0	0	0	0	0
6319525	REG2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320271	RTR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324162	SKO1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366633	SNM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322778	STB6	0	0	0	0	0	0	0	0	0	0	0	0	0
6325081	TRE1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324447	TRM13	0	0	0	0	0	0	0	0	0	0	0	0	0
6323236	HRD3	0	0	0	0	0	0	0	0	0	0	0	0	0
6325184	UBP16	0	0	0	0	0	0	0	0	0	0	0	0	0
398364885	UBP5	0	0	0	0	0	0	0	0	0	0	0	0	0
6325071	UIP4	0	0	0	0	0	0	0	0	0	0	0	0	0
398365127	UIP5	0	0	0	0	0	0	0	0	0	0	0	0	0
6319904	YIH1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325266	SUT2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324449	HRP1	2	4.31	1	1	0	0	0	0	0	0	0	0	0
6323278	YEF3	20	3.07	2	3	16.00	11	30	5.36	5	30	0.77	1	1
6322084	PFK26	0	0	0	0	0	0	0	0	0	0	0	0	0
6323057	RPL15A	2	0	0	0	13.73	2	4	0	0	0	0	0	0
6323769	RPL15B	2	0	0	0	13.73	2	4	0	0	0	0	0	0
6319282	FUN12	0	0	0	0	30.04	18	110	5.79	6	8	0	0	0
6320190	NOP1	10	0	0	0	37.92	10	33	18.96	6	20	0	0	0
6323654	CLU1	0	0	0	0	22.87	21	89	9.55	12	19	0	0	0
6324886	NOP58	0	6.85	2	2	34.44	11	70	10.18	4	4	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6321807	DED81	6	0	0	0	6.86	3	5	1.99	1	1	9.39	4	48
398364767	SUI2	0	0	0	0	16.78	3	9	3.62	1	2	0	0	0
6323011	DPS1	4	0	0	0	6.82	3	3	7.18	4	5	5.75	2	21
6323885	RRP5	1	0	0	0	9.66	11	30	1.27	2	2	0	0	0
398366547	RPP2B	16	0	0	0	53.64	4	8	10.00	1	1	0	0	0
6323288	HSP60	50	2.80	1	1	37.76	18	112	0	0	0	0	0	0
6321599	NSR1	0	0	0	0	16.91	5	11	9.90	4	4	0	0	0
6320863	GCD11	0	0	0	0	26.76	10	38	10.06	5	10	3.04	1	2
398366591	NPL3	19	19.32	4	14	11.84	3	13	11.84	3	53	3.38	1	1
6319483	IPP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325221	PMA2	1	0	0	0	6.65	4	9	0	0	0	0	0	0
6322745	GFA1	3	0	0	0	30.82	14	92	0	0	0	0	0	0
398364525	LSG1	0	0	0	0	0	0	0	1.72	1	4	0	0	0
6325038	SAR1	0	15.79	2	3	86.32	13	152	0	0	0	0	0	0
6323840	HSC82	44	7.94	3	3	37.45	24	300	4.96	3	3	1.99	1	1
6321881	GAR1	0	0	0	0	32.68	4	22	0	0	0	0	0	0
6320164	NAT1	0	0	0	0	0	0	0	3.63	3	5	0	0	0
398366403	RPS12	1	0	0	0	39.86	4	7	0	0	0	0	0	0
6322830	CDC16	0	0	0	0	0	0	0	0	0	0	0	0	0
330443397	ATP1	14	0	0	0	42.57	17	114	0	0	0	0	0	0
6324709	IDH2	3	0	0	0	44.17	9	76	0	0	0	0	0	0
6325178	RPL21B	4	0	0	0	5.63	1	1	0	0	0	0	0	0
6321903	UBA4	0	0	0	0	4.09	2	3	0	0	0	0	0	0
6325138	DBP1	12	1.78	1	1	6.16	3	7	3.24	2	10	0	0	0
6325248	TAE2	0	0	0	0	4.14	3	8	0	0	0	0	0	0
330443505	DBP10	0	0	0	0	8.64	6	7	0	0	0	0	0	0
6320148	PSA1	4	4.99	1	2	62.88	15	983	0	0	0	0	0	0
6325016	HSP82	40	4.09	2	2	36.39	22	248	4.94	3	3	1.97	1	1
6320874	ARB1	0	0	0	0	26.39	11	49	6.07	3	3	0	0	0
6319673	PGI1	2	0	0	0	8.12	3	12	0	0	0	0	0	0
398365251	GSC2	0	0	0	0	0	0	0	0.53	1	1	0	0	0
6325018	SUI3	3	0	0	0	18.25	4	14	9.47	3	5	0	0	0
6323374	FKS1	0	0	0	0	0	0	0	0.53	1	1	0	0	0
14318479	ACT1	19	7.73	2	4	24.80	5	27	2.67	1	1	0	0	0
6323554	TUB1	4	0	0	0	25.73	8	66	0	0	0	0	0	0
6321724	ZUO1	2	0	0	0	3.70	1	1	2.77	1	1	2.31	1	1
6320842	NUG1	0	0	0	0	27.31	10	32	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6322652	YKT6	2	0	0	0	56.00	9	79	0	0	0	0	0	0
6321706	FOL2	0	0	0	0	3.70	1	1	0	0	0	0	0	0
6319546	HSP26	13	0	0	0	30.84	5	11	5.14	2	3	22.43	3	112
398364355	RPL29	2	32.20	2	2	13.56	1	1	0	0	0	0	0	0
398365973	TIF4631	6	0	0	0	18.91	10	35	1.16	1	1	0	0	0
6321929	ARO9	0	0	0	0	6.63	2	4	0	0	0	0	0	0
6321808	YHR020W	8	2.76	1	1	41.13	18	64	2.33	1	2	0	0	0
6323729	ADH3	0	0	0	0	0	0	0	0	0	0	0	0	0
6319395	ILS1	2	0	0	0	4.20	3	3	0	0	0	1.59	1	1
6323969	NIP1	4	0	0	0	5.30	3	6	1.23	1	1	0	0	0
6322666	FAS1	8	0.93	1	1	12.14	16	48	0	0	0	0	0	0
6323568	TCB3	0	0	0	0	1.04	1	1	0	0	0	0	0	0
398366505	HXT7	0	0	0	0	0	0	0	0	0	0	0	0	0
6322289	CPS1	0	0	0	0	5.21	3	4	0	0	0	0	0	0
398366285	ALA1	4	0	0	0	10.86	8	19	0	0	0	0	0	0
6319553	RPG1	15	0	0	0	7.78	5	9	1.24	1	1	0	0	0
6325298	TIF5	0	0	0	0	3.21	1	2	2.72	1	1	0	0	0
398366229	FAA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323613	TSA1	11	0	0	0	33.16	6	23	4.08	1	1	0	0	0
6323183	RNH203	0	0	0	0	0	0	0	0	0	0	0	0	0
6320359	CPR1	0	0	0	0	20.99	2	5	0	0	0	0	0	0
6321236	ARO8	0	0	0	0	8.80	3	9	0	0	0	0	0	0
6323861	PFK2	42	2.50	1	1	16.58	9	42	0	0	0	5.94	4	11
398365253	TIF1	21	4.05	1	1	52.66	15	215	5.32	2	2	0	0	0
6323511	TUB3	3	0	0	0	18.20	5	54	0	0	0	0	0	0
37362625	SRO9	10	25.58	5	11	28.34	7	57	0	0	0	0	0	0
398365115	BMH1	9	0	0	0	38.95	8	37	0	0	0	0	0	0
398366063	TYS1	0	0	0	0	0	0	0	2.03	1	2	0	0	0
6323002	HSP104	2	0	0	0	14.98	10	24	3.85	3	3	0	0	0
6324902	MYO2	0	0	0	0	15.76	17	50	0	0	0	0	0	0
6323947	HAS1	0	0	0	0	12.08	5	11	4.36	2	2	0	0	0
6320376	HSP42	0	0	0	0	2.93	1	1	0	0	0	0	0	0
398364255	HYP2	0	0	0	0	11.46	1	1	0	0	0	0	0	0
6320809	SNU13	0	19.05	1	2	36.51	2	13	0	0	0	0	0	0
6323021	DRS1	0	0	0	0	16.49	8	11	0	0	0	0	0	0
398366329	PYK2	4	0	0	0	1.98	1	17	1.98	1	1	1.98	1	22
398365741	NOC2	0	0	0	0	7.46	4	9	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6321027	DMC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364483	HOR2	27	0	0	0	19.20	4	8	0	0	0	0	0	0
398365429	PDR5	1	0	0	0	3.64	3	4	0	0	0	0	0	0
6325245	RRP12	0	0	0	0	6.03	6	22	0	0	0	0	0	0
398364937	TOP2	0	0	0	0	0.91	1	1	0.77	1	1	1	0	0
6324660	TCB1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320056	COP1	0	0	0	0	23.23	18	55	0.67	1	1	0	0	0
398365597	BMH2	9	0	0	0	29.30	6	32	0	0	0	0	0	0
6325030	NEW1	1	0	0	0	6.19	5	10	0	0	0	0	0	0
6321809	RPS27B	0	0	0	0	0	0	0	0	0	0	0	0	0
6321012	CHD1	0	0	0	0	8.45	9	24	0.68	1	1	0	0	0
6320499	SSD1	0	0	0	0	1.60	1	1	1.28	2	3	0	0	0
6323072	TRX1	0	0	0	0	41.75	4	11	0	0	0	0	0	0
6325238	VTC3	1	0	0	0	0	0	0	0	0	0	0	0	0
398365319	MAK21	0	0	0	0	7.41	6	10	0	0	0	0	0	0
398364327	LSB3	0	0	0	0	40.09	11	86	1.74	1	1	3.92	1	14
6319513	ATP3	30	0	0	0	33.12	7	36	13.50	4	4	0	0	0
6324140	SRP1	0	0	0	0	16.42	5	15	0	0	0	0	0	0
6324297	HHT2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323306	DBP9	0	0	0	0	15.32	7	20	0	0	0	0	0	0
6324039	RFC3	0	0	0	0	12.94	3	7	2.65	1	1	0	0	0
6321193	RTF1	0	0	0	0	5.73	2	3	0	0	0	0	0	0
398366149	TRX2	0	0	0	0	67.31	7	22	0	0	0	0	0	0
6322452	CCT8	2	0	0	0	51.23	20	82	0	0	0	0	0	0
6321679	PFK1	7	0	0	0	17.02	11	34	0	0	0	5.37	3	3
6324219	NOP15	0	0	0	0	5.91	1	4	0	0	0	0	0	0
330443717	CTR9	0	0	0	0	4.36	3	4	2.32	1	1	0	0	0
6323755	ILV2	4	3.06	1	1	31.00	13	140	6.55	2	2	0	0	0
6320276	PAA1	0	0	0	0	18.85	3	12	0	0	0	0	0	0
6681849	RPL42A	0	0	0	0	0	0	0	0	0	0	0	0	0
14318517	SEC4	0	0	0	0	13.02	2	9	0	0	0	0	0	0
6320400	MSS116	0	0	0	0	0	0	0	0	0	0	0	0	0
6321389	TIF4632	4	0	0	0	2.08	2	8	0	0	0	0	0	0
6324081	RPA49	0	0	0	0	28.67	7	22	0	0	0	0	0	0
398366347	MES1	0	0	0	0	19.17	10	25	0	0	0	0	0	0
6320418	TCP1	6	0	0	0	49.19	18	126	0	0	0	0	0	0
398365595	PIL1	16	5.31	1	2	20.94	5	16	2.95	1	1	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
398364417	SAH1	1	0	0	0	17.59	7	24	2.45	1	1	0	0	0
6320396	RVB1	0	0	0	0	44.28	13	88	0	0	0	0	0	0
6320077	CDC48	6	0	0	0	27.54	15	84	0	0	0	0	0	0
6325025	FAS2	8	0	0	0	8.69	9	18	0	0	0	0	0	0
6324517	THI20	0	0	0	0	3.09	1	1	0	0	0	0	0	0
6323030	NOC3	0	0	0	0	3.77	2	3	0	0	0	0	0	0
6320143	TSR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321318	PRP43	0	0	0	0	14.86	8	18	3.91	3	4	0	0	0
6319618	MAK5	0	0	0	0	3.36	2	7	1.42	1	1	0	0	0
398364755	URB1	0	0	0	0	1.42	2	3	0	0	0	0	0	0
6320780	POL5	0	0	0	0	0	0	0	0	0	0	0	0	0
6325021	RVB2	2	0	0	0	32.70	11	48	0	0	0	0	0	0
6322350	CCT7	1	0	0	0	25.09	10	59	0	0	0	0	0	0
14270688	UTP20	0	0	0	0	1.72	3	5	0	0	0	0	0	0
6324197	KRE33	0	0	0	0	10.89	7	11	0.85	1	2	0	0	0
6321816	RPN1	0	0	0	0	8.86	6	25	0	0	0	0	0	0
6322901	NAP1	3	0	0	0	20.38	5	12	0	0	0	0	0	0
6320030	GLT1	3	0	0	0	14.73	21	65	0	0	0	0.56	1	2
6323470	CAR2	2	0	0	0	11.32	3	5	0	0	0	0	0	0
6320956	KAP123	1	0	0	0	44.38	29	205	0	0	0	0	0	0
398365941	SEC7	0	0	0	0	13.64	16	49	0	0	0	0.40	1	1
6323324	GSP1	0	0	0	0	20.55	4	14	4.57	1	1	0	0	0
6322697	GPM1	24	4.86	1	2	45.75	10	136	14.17	3	5	12.15	2	10
6325013	SRP68	0	0	0	0	9.68	4	6	0	0	0	0	0	0
6322103	AVT7	0	0	0	0	0	0	0	0	0	0	0	0	0
6322049	CCT2	5	6.45	2	3	59.58	20	111	0	0	0	0	0	0
330443694	SLA2	0	0	0	0	23.66	13	39	2.07	2	2	0	0	0
6321804	YSC84	0	0	0	0	25.00	7	42	1.71	1	1	0	0	0
398366021	CCT6	1	0	0	0	42.86	14	63	0	0	0	0	0	0
6324042	SEC21	0	0	0	0	16.79	9	22	0	0	0	0	0	0
398364741	SCS2	0	0	0	0	12.30	2	3	0	0	0	0	0	0
6325253	LSP1	26	0	0	0	18.48	4	17	3.52	1	1	0	0	0
6321390	RPT6	1	0	0	0	6.42	2	9	0	0	0	0	0	0
398364807	PNC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325164	NOG1	0	0	0	0	9.27	4	9	0	0	0	1.85	1	1
6319814	GRX1	0	0	0	0	31.82	3	6	0	0	0	0	0	0
6322823	MAE1	0	2.39	1	2	29.00	11	83	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6320255	TPI1	4	0	0	0	8.06	1	3	0	0	0	6.05	1	2
6320983	GDI1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365205	RPT5	0	5.99	2	2	33.64	10	39	0	0	0	0	0	0
6324992	KEL3	0	0	0	0	0	0	0	1.54	1	1	0	0	0
6323028	DNM1	0	0	0	0	20.48	10	28	0	0	0	0	0	0
398365977	RPT4	1	0	0	0	16.93	5	13	0	0	0	0	0	0
6320058	CCT4	1	0	0	0	21.59	7	48	1.52	1	1	3.03	1	2
6320379	HMO1	0	0	0	0	4.88	1	7	0	0	0	0	0	0
6322508	CYC1	0	0	0	0	7.34	1	1	0	0	0	0	0	0
6325137	VPS30	0	0	0	0	0	0	0	0	0	0	0	0	0
6321301	SEC27	0	0	0	0	16.20	10	39	0	0	0	1.80	1	2
6321673	YHB1	0	0	0	0	42.86	10	72	0	0	0	0	0	0
27469359	SYN8	0	0	0	0	0	0	0	0	0	0	0	0	0
6324433	CDC33	0	0	0	0	30.52	6	27	7.51	1	1	0	0	0
6323571	POB3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365157	NOP2	0	0	0	0	8.90	4	7	0	0	0	0	0	0
6321228	YPT32	0	0	0	0	0	0	0	0	0	0	0	0	0
398366559	RPT3	1	0	0	0	23.36	6	37	0	0	0	0	0	0
6322704	RPT1	2	0	0	0	13.92	5	12	0	0	0	0	0	0
398366379	PRT1	1	0	0	0	6.29	4	7	0	0	0	0	0	0
398365125	PDA1	3	0	0	0	8.57	3	11	0	0	0	0	0	0
398364913	GND1	23	2.86	1	3	12.07	4	11	0	0	0	14.31	5	13
6322637	SAC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365933	CYS4	17	4.14	1	1	46.15	15	97	0	0	0	0	0	0
6323903	FAA4	0	0	0	0	12.39	6	22	0	0	0	0	0	0
6319806	PDI1	10	0	0	0	4.21	1	3	0	0	0	0	0	0
6323512	PHO84	0	0	0	0	0	0	0	0	0	0	0	0	0
6325367	RPC40	0	0	0	0	25.07	5	18	0	0	0	0	0	0
6319835	GBP2	0	0	0	0	0	0	0	7.26	3	4	0	0	0
6319290	MYO4	0	0	0	0	8.16	9	15	0	0	0	0	0	0
14318531	GCN20	0	0	0	0	0	0	0	0	0	0	0	0	0
6320869	YPT31	0	0	0	0	0	0	0	0	0	0	0	0	0
398365617	UTP22	0	0	0	0	7.03	6	13	0	0	0	0	0	0
398366209	RTN1	5	0	0	0	5.76	1	2	4.07	1	1	4.07	1	1
6325097	CDC60	0	0	0	0	1.38	1	1	1.01	1	1	0	0	0
6323294	RPS28B	1	0	0	0	31.34	2	4	13.43	1	1	13.43	1	2
6325151	SSE1	29	0	0	0	38.96	18	117	0	0	0	1.59	1	1

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
398364441	NOT3	0	0	0	0	9.21	5	9	0	0	0	0	0	0
398366191	RPN13	0	0	0	0	28.21	3	12	0	0	0	0	0	0
6321453	YGR016W	0	0	0	0	5.26	1	1	0	0	0	0	0	0
6320930	ILV1	1	0	0	0	48.96	17	97	0	0	0	0	0	0
6323016	PUF3	0	0	0	0	3.19	2	4	0	0	0	0	0	0
37362700	SRP72	0	0	0	0	21.88	9	21	0	0	0	0	0	0
6321326	TAF6	0	3.49	1	1	12.98	4	9	0	0	0	0	0	0
6322565	ADO1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319722	ISW1	0	0	0	0	5.58	6	16	0.80	1	1	0	0	0
330443605	SET2	0	0	0	0	3.00	2	3	0	0	0	0	0	0
398365101	ARP3	0	0	0	0	17.59	5	8	0	0	0	1.78	1	1
6321780	STE20	0	0	0	0	9.80	5	8	0	0	0	0	0	0
6321776	PRS3	0	0	0	0	37.50	9	34	0	0	0	0	0	0
6323214	RPL37A	8	0	0	0	0	0	0	0	0	0	0	0	0
6321757	ECM29	0	0	0	0	8.57	12	19	0	0	0	0	0	0
398365069	ADH5	1	0	0	0	2.28	1	5	0	0	0	0	0	0
398365695	CPR8	0	0	0	0	0	0	0	0	0	0	0	0	0
6319854	RVS161	0	7.92	1	1	56.98	11	106	4.15	1	2	0	0	0
6324620	DBP5	0	0	0	0	49.79	15	66	0	0	0	0	0	0
6321399	YGL039W	0	0	0	0	16.95	3	6	0	0	0	0	0	0
398365061	SUP45	1	0	0	0	44.62	14	69	0	0	0	0	0	0
6324673	KTR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323387	ILV5	0	5.32	1	1	5.32	1	4	0	0	0	0	0	0
6323795	TIF34	0	5.48	1	1	36.31	8	19	2.31	1	1	0	0	0
6322667	PRS1	2	0	0	0	18.27	6	21	0	0	0	0	0	0
6320910	RGI1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365575	HOM6	0	6.69	1	1	0	0	0	0	0	0	0	0	0
6323916	TIF11	0	0	0	0	7.84	1	1	0	0	0	0	0	0
6321612	RBG2	10	0	0	0	21.74	5	20	0	0	0	0	0	0
6323225	PWP1	0	0	0	0	2.26	1	1	3.99	2	2	0	0	0
6324737	SSP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323450	CDC73	0	0	0	0	2.54	1	1	0	0	0	3.56	1	1
398365461	SRP40	0	0	0	0	0	0	0	0	0	0	0	0	0
6320708	RPL37B	0	0	0	0	0	0	0	0	0	0	0	0	0
398364957	MKT1	0	0	0	0	7.71	4	7	0	0	0	0	0	0
6320079	YDL124W	0	0	0	0	0	0	0	0	0	0	0	0	0
398364371	RSC8	0	0	0	0	6.46	3	8	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6321024	ECM32	0	0	0	0	0	0	0	0	0	0	0	0	0
37362660	FUR1	0	8.80	1	2	33.33	4	17	0	0	0	0	0	0
398364665	SEC18	2	0	0	0	24.93	14	43	0	0	0	0	0	0
6323391	ADE13	0	0	0	0	0	0	0	0	0	0	0	0	0
6323699	STV1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321412	TRP5	8	0	0	0	56.58	26	212	0	0	0	0	0	0
6321973	OYE2	0	0	0	0	0	0	0	0	0	0	0	0	0
6321801	ARD1	0	0	0	0	6.30	1	1	0	0	0	0	0	0
10383761	HIS4	0	0	0	0	2.13	1	1	0	0	0	0	0	0
398365759	SAC6	0	0	0	0	16.20	6	11	0	0	0	0	0	0
6321721	BGL2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364979	CCZ1	0	0	0	0	0	0	0	0	0	0	0	0	0
42742256	VTC4	0	0	0	0	0	0	0	0	0	0	0	0	0
6321814	THR1	0	0	0	0	21.57	5	20	0	0	0	3.08	1	1
10383798	PHO87	0	0	0	0	0	0	0	0	0	0	0	0	0
6319847	CDC10	0	0	0	0	11.49	2	6	0	0	0	0	0	0
398365363	URA8	0	0	0	0	11.25	3	21	0	0	0	0	0	0
6323138	AHP1	0	0	0	0	11.36	1	7	0	0	0	0	0	0
6320657	UTP6	0	0	0	0	0	0	0	0	0	0	0	0	0
6321919	YHR127W	0	0	0	0	0	0	0	0	0	0	0	0	0
41629688	PSP2	0	0	0	0	7.08	3	22	11.80	5	10	0	0	0
398364633	APS3	0	0	0	0	11.86	2	2	0	0	0	0	0	0
6322843	MRT4	0	0	0	0	24.15	4	13	0	0	0	0	0	0
330443486	PAT1	1	0	0	0	9.30	4	5	0	0	0	0	0	0
37362659	RRP3	0	0	0	0	31.34	11	42	2.20	1	1	0	0	0
6320845	SEC3	0	0	0	0	7.49	6	10	0	0	0	0	0	0
6324791	RFC1	0	0	0	0	7.32	5	10	0	0	0	0	0	0
6325369	MRD1	0	0	0	0	6.76	5	9	0	0	0	0	0	0
6322635	YRA2	0	0	0	0	0	0	0	9.36	1	1	0	0	0
6320368	SEC1	0	0	0	0	21.69	9	31	0	0	0	0	0	0
398364935	UBP3	0	2.30	2	2	4.93	3	10	0.99	1	1	0	0	0
6320789	GLY1	0	0	0	0	4.65	1	1	0	0	0	0	0	0
6325172	SEC16	0	0	0	0	8.75	11	25	0	0	0	0	0	0
398364311	TMA20	0	0	0	0	6.08	1	1	0	0	0	0	0	0
6320926	UTP7	0	0	0	0	3.25	1	4	0	0	0	0	0	0
330443568	ENP2	0	0	0	0	2.12	1	2	0	0	0	0	0	0
6324328	CIT1	0	0	0	0	2.51	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6322845	CAP1	1	0	0	0	13.06	2	5	0	0	0	0	0	0
6323625	PPZ1	0	0	0	0	3.90	2	3	0	0	0	0	0	0
6321267	ROK1	0	0	0	0	9.04	3	10	0	0	0	0	0	0
6321422	KAP122	0	0	0	0	6.66	4	6	0	0	0	0	0	0
6323717	ABF2	9	0	0	0	0	0	0	0	0	0	0	0	0
27808701	IST2	0	0	0	0	0	0	0	0	0	0	0	0	0
6322942	TGL4	0	0	0	0	0	0	0	0	0	0	0	0	0
6319307	CYS3	2	0	0	0	7.87	2	7	0	0	0	0	0	0
6322719	SHE2	0	0	0	0	10.98	2	2	0	0	0	0	0	0
6323837	SSO2	0	0	0	0	0	0	0	0	0	0	0	0	0
6319923	SRB8	0	0	0	0	0	0	0	0	0	0	0	0	0
6321290	ARO2	0	0	0	0	6.65	2	2	0	0	0	0	0	0
6325072	MRN1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324291	IDH1	0	0	0	0	14.72	4	5	0	0	0	0	0	0
6325024	SSO1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366609	PPN1	0	0	0	0	2.82	1	2	0	0	0	0	0	0
6323639	GLO1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321832	NCP1	3	0	0	0	0	0	0	0	0	0	0	0	0
398364239	FRD1	0	0	0	0	3.40	1	1	0	0	0	0	0	0
6320235	PST2	5	0	0	0	19.70	2	8	0	0	0	0	0	0
6322770	VMA5	0	4.34	1	2	38.27	11	105	0	0	0	0	0	0
6325209	CAM1	1	0	0	0	0	0	0	5.06	2	2	0	0	0
6320041	DHH1	4	0	0	0	29.25	10	53	6.32	3	4	0	0	0
6323635	ERG6	0	0	0	0	4.70	1	1	0	0	0	0	0	0
6319698	PDB1	0	0	0	0	17.21	4	11	0	0	0	0	0	0
398365225	IMD2	0	0	0	0	4.40	2	3	0	0	0	2.87	1	1
6324990	DIP5	0	0	0	0	0	0	0	0	0	0	0	0	0
6323585	IMD4	2	3.05	1	1	5.92	2	3	0	0	0	2.86	1	1
6323386	TAL1	1	0	0	0	0	0	0	0	0	0	0	0	0
398366255	IMD3	2	3.06	1	1	5.93	2	3	0	0	0	2.87	1	1
6319426	COR1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364269	SEC24	0	0	0	0	3.13	2	3	0	0	0	0	0	0
6320695	RIB3	0	0	0	0	0	0	0	0	0	0	0	0	0
398364515	URM1	0	0	0	0	13.13	1	1	0	0	0	0	0	0
398364803	GLC7	0	0	0	0	19.55	4	11	0	0	0	0	0	0
398365877	HOM2	1	0	0	0	20.82	5	19	0	0	0	0	0	0
6324428	NOP8	0	0	0	0	2.69	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6324550	IFM1	0	0	0	0	1.92	1	1	0	0	0	0	0	0
6324950	ALD4	1	0	0	0	6.55	2	2	0	0	0	0	0	0
6324635	CKA2	0	0	0	0	17.11	4	7	0	0	0	0	0	0
6319849	YCP4	0	0	0	0	0	0	0	0	0	0	0	0	0
6325202	LGE1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366027	SUR4	0	0	0	0	0	0	0	0	0	0	0	0	0
6321641	PCT1	0	0	0	0	4.72	1	2	0	0	0	0	0	0
6325280	EAF3	0	0	0	0	6.98	2	2	0	0	0	0	0	0
6320803	MCM3	0	0	0	0	14.52	10	24	0	0	0	0	0	0
6323872	SKY1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325140	IDI1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320049	SAS10	0	0	0	0	5.74	2	2	0	0	0	0	0	0
6324989	DIM1	0	0	0	0	9.12	3	4	3.46	1	1	0	0	0
398366039	PBP1	14	7.62	3	7	16.20	7	26	3.32	2	2	0	0	0
6323088	REX2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324899	PRO2	0	0	0	0	4.17	1	2	0	0	0	0	0	0
398365467	YNL010W	0	0	0	0	0	0	0	0	0	0	0	0	0
41629671	SEA4	0	0	0	0	0	0	0	0	0	0	0	0	0
6320704	PUF6	0	0	0	0	1.68	1	1	0	0	0	0	0	0
6323650	PLB1	0	0	0	0	6.33	1	1	0	0	0	0	0	0
6323158	DIP2	0	0	0	0	1.27	1	1	0	0	0	0	0	0
6322727	SRP21	0	0	0	0	4.79	1	2	0	0	0	0	0	0
14318504	MDJ1	0	0	0	0	10.76	3	5	0	0	0	0	0	0
6322878	GCN3	0	0	0	0	16.39	3	11	0	0	0	0	0	0
330443380	PSK1	0	0	0	0	5.16	4	5	0	0	0	0	0	0
398364789	UFD4	0	0	0	0	8.56	8	13	0	0	0	0	0	0
6319903	PWP2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324613	CKB2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324568	TOP1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365393	MDM32	0	0	0	0	0	0	0	0	0	0	0	0	0
6322731	VPH2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323653	HXT2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320236	MRH1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364559	NSP1	2	1.82	1	1	2.07	1	2	0	0	0	0	0	0
6321498	ADE6	0	0	0	0	6.55	6	16	0	0	0	0	0	0
6319326	ADE1	0	0	0	0	3.92	1	1	0	0	0	0	0	0
6321812	MYO1	14	0	0	0	3.37	5	8	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
398365005	RIX1	0	0	0	0	3.01	2	5	0	0	0	4.59	1	1
6323975	YMR315	1	0	0	0	4.01	1	4	0	0	0	0	0	0
6323740	UTP15	0	0	0	0	3.70	1	2	0	0	0	0	0	0
6322743	AAT1	0	0	0	0	2.22	1	2	0	0	0	0	0	0
6323033	SSL1	0	0	0	0	3.90	1	2	0	0	0	0	0	0
398365521	SWT1	0	0	0	0	2.84	1	2	0	0	0	0	0	0
398365403	NUP133	0	0	0	0	1.12	1	1	0	0	0	0	0	0
6319276	CDC24	0	0	0	0	1.76	1	1	0	0	0	0	0	0
6323251	UTP13	0	0	0	0	2.45	1	2	0	0	0	0	0	0
6321913	LSM12	7	16.04	2	7	25.13	3	20	0	0	0	0	0	0
6322257	RCY1	0	0	0	0	7.14	3	5	0	0	0	0	0	0
6319900	RSC6	0	0	0	0	8.90	3	5	0	0	0	0	0	0
398364743	IRC2	0	0	0	0	15.97	3	4	0	0	0	0	0	0
6320070	CDC53	2	0	0	0	5.40	3	5	0	0	0	0	0	0
398364601	NFU1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323229	YKE2	1	0	0	0	0	0	0	0	0	0	0	0	0
6322154	CKA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323687	IOC4	0	0	0	0	0	0	0	0	0	0	0	0	0
6325372	RGC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325234	MET12	0	0	0	0	0	0	0	0	0	0	0	0	0
10383790	RHB1	0	0	0	0	0	0	0	0	0	0	0	0	0
10383792	SYP1	8	0	0	0	14.94	7	14	1.72	1	1	0	0	0
398365079	SEC9	0	0	0	0	2.46	1	1	0	0	0	0	0	0
398365907	APM3	0	0	0	0	6.00	2	7	0	0	0	0	0	0
6322783	YNK1	0	0	0	0	18.95	2	3	0	0	0	0	0	0
6324470	TPT1	0	0	0	0	22.17	2	10	0	0	0	0	0	0
6321843	CIC1	0	0	0	0	12.77	3	11	0	0	0	0	0	0
6319298	CCR4	0	0	0	0	9.44	6	17	0	0	0	0	0	0
6321257	GTS1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321341	SRM1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365245	PFY1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324375	FPK1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364253	AYR1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365397	YGR054W	0	0	0	0	0	0	0	0	0	0	0	0	0
398365527	RLP7	0	0	0	0	10.56	2	8	0	0	0	0	0	0
6319976	SHS1	0	0	0	0	8.53	3	11	0	0	0	0	0	0
398364769	RSP5	0	0	0	0	8.78	5	8	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6320843	PAC2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365529	SIR1	0	0	0	0	0	0	0	0	0	0	0	0	0
14318435	TIP20	0	0	0	0	0	0	0	0	0	0	0	0	0
6323645	AIM34	0	0	0	0	0	0	0	0	0	0	0	0	0
330443386	BUD14	0	0	0	0	0	0	0	0	0	0	0	0	0
6319751	CHK1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319452	APN2	0	0	0	0	0	0	0	0	0	0	0	0	0
10383783	YCR015C	0	0	0	0	0	0	0	0	0	0	0	0	0
33438820	YJL133C-A	0	0	0	0	0	0	0	0	0	0	0	0	0
398365435	ERG25	0	0	0	0	0	0	0	0	0	0	0	0	0
6321363	MPS2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323180	PCD1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364799	PMD1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322709	TGL1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325305	TAH18	0	0	0	0	0	0	0	0	0	0	0	0	0
6322698	YKL151C	0	0	0	0	0	0	0	0	0	0	0	0	0
6322341	RPE1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323336	STT4	0	0	0	0	0	0	0	0	0	0	0	0	0
6320542	MSN5	0	0	0	0	0	0	0	0	0	0	0	0	0
6325171	ELP3	0	0	0	0	0	0	0	0	0	0	0	0	0
6319274	ERV46	0	0	0	0	0	0	0	0	0	0	0	0	0
398366207	CRM1	0	0	0	0	9.13	7	18	0	0	0	0	0	0
6321198	CSE1	0	0	0	0	3.54	3	5	0	0	0	0	0	0
398366491	SKP1	0	0	0	0	8.76	1	5	0	0	0	0	0	0
6323819	MLH1	0	0	0	0	2.21	1	1	0	0	0	0	0	0
398365033	IRA1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366427	SRP101	0	0	0	0	0	0	0	0	0	0	0	0	0
6320008	NUS1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324098	PDR16	0	0	0	0	33.33	9	35	3.42	1	1	0	0	0
6324093	SIN4	0	0	0	0	7.29	4	11	0	0	0	0	0	0
6322063	MET18	0	0	0	0	8.53	6	17	0	0	0	0	0	0
6319901	THR4	0	0	0	0	0	0	0	0	0	0	0	0	0
6321899	CDC12	0	0	0	0	0	0	0	0	0	0	0	0	0
398364305	YND1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325365	RPN7	2	0	0	0	16.32	6	13	2.10	1	1	2.10	1	3
6322288	RFA3	0	0	0	0	27.05	2	12	0	0	0	0	0	0
6323832	YMR178	0	0	0	0	6.20	1	1	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6321327	NSA1	0	0	0	0	2.59	1	1	0	0	0	0	0	0
398366645	PKH1	0	0	0	0	1.17	1	1	0	0	0	0	0	0
6323507	MSC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366107	YPP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323586	SPC2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324079	RAD50	0	0	0	0	0	0	0	0	0	0	0	0	0
6320865	GAL83	0	0	0	0	0	0	0	0	0	0	0	0	0
14318465	YFL054C	0	0	0	0	0	0	0	0	0	0	0	0	0
6325349	NVJ2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323962	UBP15	0	0	0	0	0	0	0	0	0	0	0	0	0
6323891	RNA1	0	0	0	0	35.63	11	35	0	0	0	0	0	0
6323273	MAP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320935	TRP2	2	0	0	0	31.56	11	25	0	0	0	0	0	0
6324994	YPL260W	0	0	0	0	2.36	1	9	0	0	0	0	0	0
27469358	PMT2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365969	CDC31	0	0	0	0	31.68	4	14	0	0	0	0	0	0
6320214	SNQ2	0	0	0	0	0	0	0	0	0	0	0	0	0
6322263	PHO90	0	0	0	0	0	0	0	0	0	0	0	0	0
42742305	YNL208W	2	0	0	0	28.64	3	37	0	0	0	0	0	0
6321432	PMC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323221	HCR1	0	0	0	0	0	0	0	4.15	1	2	0	0	0
6322722	PGM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323089	FRS1	0	0	0	0	2.18	1	7	0	0	0	0	0	0
6321271	PMR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322097	LYS12	22	0	0	0	43.94	15	68	0	0	0	0	0	0
6320151	SLC1	1	0	0	0	0	0	0	0	0	0	0	0	0
6319312	ERP2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365779	YCF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324082	YNL247W	0	0	0	0	0	0	0	0	0	0	0	0	0
6324625	ETT1	0	0	0	0	21.60	6	41	0	0	0	0	0	0
6321492	MUP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322434	RNR2	0	0	0	0	6.77	2	9	0	0	0	0	0	0
6320245	ENA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323853	VTI1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319837	ILV6	0	0	0	0	21.68	4	12	0	0	0	0	0	0
6319441	PET9	7	3.14	1	1	7.55	2	2	0	0	0	0	0	0
398365855	KGD2	29	0	0	0	1.94	1	1	0	0	0	3.46	1	2

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6322980	YBT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320717	GNP1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365985	RPN8	0	0	0	0	26.63	6	36	0	0	0	0	0	0
398364411	MMF1	3	0	0	0	9.66	1	2	0	0	0	0	0	0
398366385	RNH202	0	0	0	0	0	0	0	0	0	0	0	0	0
6320332	ARO1	2	0	0	0	11.52	13	39	0	0	0	0	0	0
330443439	ECM33	14	0	0	0	0	0	0	0	0	0	0	0	0
6322640	STE6	0	0	0	0	0	0	0	0	0	0	0	0	0
398365037	DNF1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366015	VPH1	0	0	0	0	1.79	1	1	0	0	0	0	0	0
398364221	SIT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323965	FKS3	0	0	0	0	0	0	0	0.56	1	1	0	0	0
6321346	NUP145	0	0	0	0	0	0	0	0.68	1	1	0	0	0
6324067	POL2	0	0	0	0	0.54	1	1	0	0	0	0	0	0
6325441	DPM1	0	0	0	0	4.87	1	1	0	0	0	0	0	0
6320150	PBP4	8	12.43	1	1	20.00	2	5	0	0	0	0	0	0
6319704	MCX1	0	0	0	0	7.31	3	5	0	0	0	0	0	0
6323235	ENT2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365451	PXL1	0	0	0	0	0	0	0	0	0	0	0	0	0
37362705	YPR089W	0	0	0	0	5.63	4	5	0	0	0	0	0	0
116006501	MGM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324111	MGS1	0	0	0	0	2.21	1	1	0	0	0	0	0	0
6321880	RPF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325407	NCE102	0	0	0	0	0	0	0	0	0	0	0	0	0
398366221	YML7	0	0	0	0	0	0	0	0	0	0	0	0	0
398364731	MPP10	0	0	0	0	3.04	1	2	0	0	0	0	0	0
6319449	PIM1	0	0	0	0	12.71	9	27	0	0	0	0	0	0
6319448	MCM2	0	0	0	0	7.95	5	9	0	0	0	1.73	1	1
6321851	VMA22	0	0	0	0	9.39	1	1	0	0	0	0	0	0
398365621	YJR149W	0	0	0	0	0	0	0	0	0	0	0	0	0
6319646	SSE2	13	0	0	0	3.75	2	4	0	0	0	0	0	0
6321332	MLC1	2	14.77	1	1	44.97	5	57	0	0	0	0	0	0
398365795	YGR126W	0	0	0	0	3.04	1	4	0	0	0	0	0	0
6324350	SNF12	0	0	0	0	29.86	11	27	0	0	0	2.47	1	1
398366497	RQC1	0	0	0	0	0	0	0	1.24	1	1	0	0	0
6322676	EBP2	0	0	0	0	3.75	1	2	0	0	0	0	0	0
6323944	HSH155	0	0	0	0	1.75	1	3	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6322531	LIA1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364353	PRO3	0	0	0	0	10.14	2	4	0	0	0	0	0	0
6322645	EAP1	1	0	0	0	0	0	0	1.74	1	1	1	0	0
6323237	SEC13	0	0	0	0	0	0	0	0	0	0	0	0	0
398365809	RUD3	0	0	0	0	0	0	0	0	0	0	0	0	0
6319930	YCR087C-	0	0	0	0	0	0	0	7.19	1	1	1	0	0
398366045	CSR1	2	0	0	0	28.92	9	25	0	0	0	0	0	0
6323938	AEP2	0	0	0	0	0	0	0	0	0	0	0	0	0
6322167	TIM44	23	0	0	0	35.03	11	60	2.09	1	1	0	0	0
6320000	TRM8	0	0	0	0	0	0	0	0	0	0	0	0	0
6319321	RFA1	0	0	0	0	0	0	0	0	0	0	1.93	1	1
6323934	PGM3	0	0	0	0	0	0	0	0	0	0	1.13	1	2
6324629	VHS3	0	0	0	0	0	0	0	0	0	0	2.37	1	1
398366511	ATP22	4	0	0	0	2.49	1	1	2.49	1	4	2.49	1	4
6323706	RNA14	0	0	0	0	0.89	1	1	0.89	1	1	0	0	0
6321395	DST1	2	0	0	0	10.36	2	3	9.71	3	3	0	0	0
398364447	CAP2	0	0	0	0	4.53	1	2	0	0	0	0	0	0
398365841	RIF1	0	0	0	0	1.10	1	1	0	0	0	0	0	0
398365073	RAD3	0	0	0	0	1.93	1	1	0	0	0	0	0	0
6319622	MRPS9	0	0	0	0	3.60	1	1	0	0	0	0	0	0
6323895	RNT1	0	0	0	0	4.03	1	2	0	0	0	0	0	0
6319317	TFC3	0	0	0	0	1.29	1	1	0	0	0	0	0	0
6323216	SKG3	0	0	0	0	1.46	1	2	0	0	0	0	0	0
6323341	CDC25	0	0	0	0	0.94	1	2	0	0	0	0	0	0
6320087	NUP84	0	0	0	0	2.34	1	2	0	0	0	0	0	0
6322188	CFD1	0	0	0	0	5.12	1	2	0	0	0	0	0	0
6324424	SPT20	0	0	0	0	2.65	1	2	0	0	0	0	0	0
6323034	SSK1	0	0	0	0	1.97	1	2	0	0	0	0	0	0
398365691	SRB5	0	0	0	0	5.86	1	1	0	0	0	0	0	0
398364427	PIG2	0	0	0	0	2.60	1	1	0	0	0	0	0	0
398364977	ALY1	0	0	0	0	1.53	1	1	0	0	0	0	0	0
6320315	FOB1	0	0	0	0	3.36	1	1	0	0	0	0	0	0
6323621	YML020W	0	0	0	0	3.77	1	1	0	0	0	0	0	0
14318535	IOC3	0	0	0	0	1.91	1	1	0	0	0	0	0	0
10383754	MRC1	0	0	0	0	1.92	1	1	0	0	0	0	0	0
6319988	NOP6	0	0	0	0	4.89	1	1	0	0	0	0	0	0
6323394	STE11	0	0	0	0	2.79	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
330443700	UBP10	0	0	0	0	2.02	1	1	0	0	0	0	0	0
330443395	PKC1	0	0	0	0	0.96	1	1	0	0	0	0	0	0
6321782	MRP4	0	0	0	0	2.79	1	1	0	0	0	0	0	0
6323846	GYL1	0	0	0	0	1.39	1	2	0	0	0	0	0	0
398365475	VPS41	0	0	0	0	1.11	1	1	0	0	0	0	0	0
6323526	COQ5	0	0	0	0	10.10	2	2	0	0	0	0	0	0
6325226	PHO85	0	0	0	0	5.57	2	3	0	0	0	0	0	0
398364611	PUP3	0	0	0	0	16.59	2	13	0	0	0	0	0	0
6323915	TRM732	0	0	0	0	1.83	2	6	0	0	0	0	0	0
6320438	HEM1	0	0	0	0	5.29	2	6	0	0	0	0	0	0
6325102	KIP2	0	0	0	0	4.25	2	4	0	0	0	0	0	0
6323575	ORC1	0	0	0	0	2.52	2	3	0	0	0	0	0	0
398365487	RRP8	0	0	0	0	8.93	2	2	0	0	0	0	0	0
6324338	PRP2	0	0	0	0	2.40	2	2	0	0	0	0	0	0
6325104	RAD53	0	0	0	0	3.90	2	2	0	0	0	0	0	0
6324107	SSU72	0	0	0	0	20.87	2	2	0	0	0	0	0	0
6319526	RFS1	0	0	0	0	15.71	2	3	0	0	0	0	0	0
6320952	NUP157	0	0	0	0	2.30	2	2	0	0	0	0	0	0
6324147	IPI3	0	0	0	0	8.83	3	6	0	0	0	0	0	0
330443409	PRS4	0	0	0	0	11.66	3	5	0	0	0	0	0	0
6322806	PRI2	0	0	0	0	10.04	3	12	0	0	0	0	0	0
330443500	UFD2	0	0	0	0	5.52	3	6	0	0	0	0	0	0
6324062	PIK1	0	0	0	0	4.60	3	3	0	0	0	0	0	0
6324801	HER1	0	1.77	1	2	3.69	3	5	0	0	0	0	0	0
398366237	PRP28	0	0	0	0	8.16	3	5	0	0	0	0	0	0
6325096	BEM4	0	0	0	0	4.90	3	5	0	0	0	0	0	0
6325402	NOC4	0	0	0	0	8.15	3	5	0	0	0	0	0	0
6324770	LIP5	6	0	0	0	13.77	3	8	0	0	0	0	0	0
398364793	DMA2	0	0	0	0	10.15	3	7	0	0	0	0	0	0
398364875	YML13	0	0	0	0	24.24	4	8	0	0	0	0	0	0
6322491	GEA1	0	0	0	0	4.47	4	9	0	0	0	0	0	0
6323061	RSC58	0	0	0	0	13.15	4	7	0	0	0	0	0	0
6323842	MRPS17	0	0	0	0	21.10	4	6	0	0	0	0	0	0
6325121	GIP3	0	0	0	0	4.23	4	10	0	0	0	0	0	0
16198342	RGD2	0	0	0	0	10.22	4	12	0	0	0	0	0	0
6323045	MEU1	0	0	0	0	21.66	5	17	0	0	0	0	0	0
6322286	SWI3	0	0	0	0	10.91	6	20	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6323630	ERV25	0	0	0	0	0	0	0	0	0	0	0	0	0
6323716	TVP18	0	0	0	0	0	0	0	0	0	0	0	0	0
6319289	SNC1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364281	EMP24	0	0	0	0	0	0	0	0	0	0	0	0	0
6320298	DNF2	0	0	0	0	0	0	0	0	0	0	0	0	0
330443597	MNN5	0	0	0	0	3.07	1	1	0	0	0	0	0	0
6320897	FCY2	0	0	0	0	0	0	0	0	0	0	0	0	0
330443542	SAP155	0	0	0	0	2.00	1	2	0	0	0	0	0	0
14318521	SPB4	0	0	0	0	1.82	1	2	0	0	0	0	0	0
6323367	NUP2	0	0	0	0	4.58	2	5	0	0	0	0	0	0
6319303	TPD3	0	0	0	0	8.19	4	11	0	0	0	0	0	0
6324166	RIA1	0	0	0	0	4.95	4	9	0	0	0	0	0	0
398364557	AIM21	0	0	0	0	0	0	0	0	0	0	0	0	0
6322369	SRS2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324088	ZWF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319890	TAF2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320033	SFA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6325429	BSP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323022	YLL007C	0	0	0	0	0	0	0	0	0	0	0	0	0
6324918	YOR342C	0	0	0	0	0	0	0	0	0	0	0	0	0
6320091	TRM3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322925	SIS2	0	0	0	0	3.91	1	2	0	0	0	0	0	0
6321999	SCH9	0	0	0	0	1.82	1	1	0	0	0	0	0	0
6320106	RPN6	0	0	0	0	19.59	6	19	0	0	0	0	0	0
6320934	PTC2	0	0	0	0	0	0	0	0	0	0	0	0	0
6322545	STE18	0	0	0	0	0	0	0	0	0	0	0	0	0
27808714	LAP3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322410	IRC8	0	0	0	0	0	0	0	0	0	0	0	0	0
14318532	UBP6	0	0	0	0	6.61	2	3	0	0	0	0	0	0
6320129	BRE1	0	0	0	0	9.43	5	17	0	0	0	0	0	0
6324287	COG6	0	0	0	0	9.89	5	14	0	0	0	0	0	0
6320846	NTF2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324508	MET22	0	0	0	0	0	0	0	0	0	0	0	0	0
6319975	GCS1	0	0	0	0	9.66	2	6	0	0	0	0	0	0
63535562	CHA1	0	0	0	0	16.39	3	9	0	0	0	0	0	0
6322141	NEO1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365175	CDC28	0	0	0	0	6.38	2	3	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6323323	SEC72	0	0	0	0	0	0	0	0	0	0	0	0	0
6321649	YGR210C	0	0	0	0	0	0	0	0	0	0	0	0	0
6323423	CCW14	0	0	0	0	0	0	0	0	0	0	0	0	0
398366295	HSP78	0	0	0	0	11.10	6	17	0	0	0	0	0	0
6324169	YGP1	0	0	0	0	3.39	1	2	0	0	0	0	0	0
6323085	ERG3	0	0	0	0	0	0	0	0	0	0	0	0	0
6319997	RTN2	0	0	0	0	0	0	0	0	0	0	0	0	0
14318526	RPN11	0	0	0	0	8.50	2	2	0	0	0	0	0	0
6320913	ARG5,6	0	0	0	0	1.51	1	1	0	0	0	0	0	0
398365815	PHB1	2	13.94	1	1	6.97	1	3	0	0	0	0	0	0
14318497	FRS2	0	0	0	0	0	0	0	0	0	0	0	0	0
6320291	SSS1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319367	ECM21	1	0	0	0	1.16	1	1	0	0	0	0.54	1	1
6319714	PRP5	0	0	0	0	0	0	0	1.06	1	1	1.06	1	1
6323841	YMR187C	0	0	0	0	0	0	0	0	0	0	1.39	1	1
6321813	MAS2	0	0	0	0	4.98	2	2	0	0	0	1.66	1	1
6322607	HMS2	0	3.35	1	1	0	0	0	0	0	0	2.23	1	1
6319750	UBX7	0	0	0	0	4.59	1	1	0	0	0	2.29	1	1
6322917	OAF3	0	0	0	0	0	0	0	0	0	0	3.82	1	1
37362687	FOL1	0	0	0	0	1.21	1	3	1.21	1	1	0	0	0
37362652	PGD1	0	0	0	0	8.31	2	3	2.52	1	1	0	0	0
37362689	GPI15	0	0	0	0	0	0	0	3.93	1	3	0	0	0
398364551	CBK1	0	0	0	0	0	0	0	0.79	1	1	0	0	0
6325386	SCD6	0	0	0	0	4.58	1	1	6.59	2	3	0	0	0
6320372	CDC37	0	0	0	0	3.75	1	1	0	0	0	0	0	0
6321266	NUP49	0	0	0	0	1.48	1	1	0	0	0	0	0	0
6325406	YPR148C	0	0	0	0	3.22	1	2	0	0	0	0	0	0
6321873	LRP1	0	0	0	0	7.07	1	3	0	0	0	0	0	0
6325152	SYH1	0	0	0	0	1.65	1	1	0	0	0	0	0	0
6320036	CDC36	0	0	0	0	7.33	1	1	0	0	0	0	0	0
6322810	VPS24	0	0	0	0	7.14	1	1	0	0	0	0	0	0
6323125	KIN2	0	0	0	0	0.78	1	2	0	0	0	0	0	0
6319555	SPT7	0	0	0	0	0.90	1	1	0	0	0	0	0	0
6322523	RPA12	0	0	0	0	12.00	1	2	0	0	0	0	0	0
6324622	RAT1	0	0	0	0	1.29	1	1	0	0	0	0	0	0
6322788	MSN4	0	0	0	0	2.70	1	5	0	0	0	0	0	0
6321889	YHR097C	0	0	0	0	4.64	1	2	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6324330	RPC34	0	0	0	0	4.10	1	1	0	0	0	0	0	0
6321796	SOD2	0	0	0	0	6.01	1	5	0	0	0	0	0	0
6320862	YAT2	4	0	0	0	1.52	1	1	0	0	0	0	0	0
6320360	RPA14	0	0	0	0	14.60	1	4	0	0	0	0	0	0
6324174	YNL155W	0	0	0	0	4.38	1	2	0	0	0	0	0	0
6323305	SMD2	0	0	0	0	7.27	1	2	0	0	0	0	0	0
6322296	HAL5	0	0	0	0	2.11	1	2	0	0	0	0	0	0
6323559	TDA9	0	0	0	0	1.04	1	1	0	0	0	0	0	0
6324374	TRM112	0	0	0	0	12.59	1	1	0	0	0	0	0	0
6325225	SVL3	0	0	0	0	1.94	1	1	0	0	0	0	0	0
6324874	BUD7	0	0	0	0	2.28	1	3	0	0	0	0	0	0
6322261	ACO2	0	0	0	0	1.01	1	1	0	0	0	0	0	0
398365029	BCH2	0	0	0	0	1.57	1	1	0	0	0	0	0	0
6324371	MVD1	0	0	0	0	4.80	1	1	0	0	0	0	0	0
6321623	UBR1	0	0	0	0	1.08	1	1	0	0	0	0	0	0
6321425	PDR1	0	0	0	0	1.31	1	1	0	0	0	0	0	0
6324203	SPC98	0	0	0	0	1.89	1	1	0	0	0	0	0	0
6325129	TBF1	0	0	0	0	2.31	1	1	0	0	0	0	0	0
6322644	LOS1	0	0	0	0	1.73	1	1	0	0	0	0	0	0
6322363	SAP185	0	0	0	0	1.04	1	1	0	0	0	0	0	0
6323081	IES3	0	0	0	0	8.40	1	2	0	0	0	0	0	0
6321378	YBP2	0	0	0	0	1.72	1	2	0	0	0	0	0	0
398364843	POL1	0	0	0	0	1.29	1	1	0	0	0	0	0	0
6321764	NPR3	0	0	0	0	2.53	1	1	0	0	0	0	0	0
398366625	SDC1	0	0	0	0	6.86	1	1	0	0	0	0	0	0
6322540	AIM24	0	0	0	0	2.79	1	1	0	0	0	0	0	0
6320026	AIR2	0	0	0	0	4.94	1	1	0	0	0	0	0	0
6323114	ARP6	0	0	0	0	2.51	1	1	0	0	0	0	0	0
6324096	BNI4	0	0	0	0	1.79	1	1	0	0	0	0	0	0
6322876	DBP7	0	0	0	0	2.29	1	1	0	0	0	0	0	0
6321458	YGR021W	0	0	0	0	3.45	1	1	0	0	0	0	0	0
398365777	YGR122W	0	0	0	0	3.73	1	1	0	0	0	0	0	0
398365293	LAS1	0	0	0	0	2.79	1	1	0	0	0	0	0	0
398364695	MAK11	0	0	0	0	3.85	1	1	0	0	0	0	0	0
6324805	MKK1	0	0	0	0	3.35	1	1	0	0	0	0	0	0
6321398	HEM2	0	0	0	0	5.56	1	1	0	0	0	0	0	0
6319365	RTG3	0	0	0	0	2.06	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6322549	GRR1	0	0	0	0	0.96	1	1	0	0	0	0	0	0
6323551	UFO1	0	0	0	0	1.80	1	1	0	0	0	0	0	0
6323108	SIC1	0	0	0	0	6.69	1	1	0	0	0	0	0	0
6324100	URE2	0	0	0	0	3.39	1	1	0	0	0	0	0	0
6321534	ASK10	0	0	0	0	1.48	1	1	0	0	0	0	0	0
6322907	DYN1	0	0	0	0	0.49	1	1	0	0	0	0	0	0
6319529	PRP6	0	0	0	0	1.67	1	3	0	0	0	0	0	0
6321907	DMA1	0	0	0	0	5.77	1	1	0	0	0	0	0	0
398364783	FES1	0	0	0	0	10.69	2	7	0	0	0	0	0	0
6324188	AAH1	0	0	0	0	9.22	2	2	0	0	0	0	0	0
398364975	SHE3	0	0	0	0	7.29	2	4	0	0	0	0	0	0
398364251	RAD23	0	0	0	0	7.54	2	4	0	0	0	0	0	0
6320565	GGA1	0	0	0	0	5.21	2	2	0	0	0	0	0	0
398365259	UBP2	0	0	0	0	2.28	2	4	0	0	0	0	0	0
398364701	CYR1	0	0	0	0	1.33	2	2	0	0	0	0	0	0
398365573	GRX3	0	0	0	0	8.42	2	5	0	0	0	0	0	0
6321217	NIF3	0	0	0	0	16.32	2	2	0	0	0	0	0	0
6323525	BUL2	0	0	0	0	3.15	2	3	0	0	0	0	0	0
6324490	ATG19	0	0	0	0	7.23	2	4	0	0	0	0	0	0
14318523	LOC1	0	0	0	0	11.76	2	3	0	0	0	0	0	0
37362641	CIN8	0	0	0	0	2.50	2	2	0	0	0	0	0	0
6324033	MON2	0	0	0	0	1.22	2	2	0	0	0	0	0	0
6321948	RTT107	0	0	0	0	2.15	2	2	0	0	0	0	0	0
14318438	MET5	0	0	0	0	1.87	2	3	0	0	0	0	0	0
6320454	YDR248C	0	0	0	0	17.10	2	6	0	0	0	0	0	0
398366363	SNX3	0	0	0	0	10.49	2	4	0	0	0	0	0	0
398365949	NAT5	0	0	0	0	17.61	2	3	0	0	0	0	0	0
6324279	SFB2	0	0	0	0	3.31	2	4	0	0	0	0	0	0
6322400	NUP82	0	0	0	0	5.19	2	8	0	0	0	0	0	0
398366177	ZPR1	0	0	0	0	7.41	3	5	0	0	0	0	0	0
6323054	SED5	0	0	0	0	12.94	3	4	0	0	0	0	0	0
6324123	RTT106	0	0	0	0	9.45	3	8	0	0	0	0	0	0
6319977	WHI4	0	0	0	0	6.63	3	6	0	0	0	0	0	0
398365083	SKN7	0	0	0	0	7.88	3	8	0	0	0	0	0	0
14318486	RIM15	0	0	0	0	2.82	3	5	0	0	0	0	0	0
398366197	PCF11	0	0	0	0	5.27	3	5	0	0	0	0	0	0
6681848	MOB1	0	0	0	0	16.24	3	4	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6322378	TAX4	0	0	0	0	0	0	0	0	0	0	0	0	0
6320848	TIR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320093	TMA17	0	0	0	0	0	0	0	0	0	0	0	0	0
6322332	TRK1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320738	QCR7	0	0	0	0	0	0	0	0	0	0	0	0	0
6321543	VOA1	0	0	0	0	0	0	0	0	0	0	0	0	0
41629668	VPS8	0	0	0	0	0	0	0	0	0	0	0	0	0
6323150	YPS3	0	0	0	0	0	0	0	0	0	0	0	0	0
398365307	EHT1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322739	KTI12	0	0	0	0	0	0	0	0	0	0	0	0	0
398366117	ELP2	0	0	0	0	0	0	0	0	0	0	0	0	0
6324932	CIR2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323129	ERG27	0	0	0	0	0	0	0	0	0	0	0	0	0
6322737	ABF1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321278	AIM14	0	0	0	0	0	0	0	0	0	0	0	0	0
398364589	DSN1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324870	YOR296W	0	0	0	0	0	0	0	0	0	0	0	0	0
398365441	ISN1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320409	RAV2	0	0	0	0	0	0	0	0	0	0	0	0	0
398366501	MRPS2	0	0	0	0	0	0	0	0	0	0	0	0	0
398365469	RPL40B	0	0	0	0	0	0	0	0	0	0	0	0	0
398364921	TFC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320130	YDL073W	0	0	0	0	0	0	0	0	0	0	0	0	0
6321209	SHE10	0	0	0	0	0	0	0	0	0	0	0	0	0
6323662	FMS1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365025	KCS1	0	0	0	0	0	0	0	0	0	0	0	0	0
398365489	LST8	0	0	0	0	0	0	0	0	0	0	0	0	0
6322593	XPT1	0	0	0	0	0	0	0	0	0	0	0	0	0
398366545	ARO10	0	0	0	0	0	0	0	0	0	0	0	0	0
398366635	PHO8	0	0	0	0	0	0	0	0	0	0	0	0	0
6321005	COG3	0	0	0	0	4.99	3	3	0	0	0	2.00	1	1
398364553	NBP35	0	0	0	0	6.71	1	1	0	0	0	2.44	1	1
6323569	COG8	0	0	0	0	4.78	2	4	0	0	0	2.97	1	3
6322229	GTT1	5	0	0	0	8.97	2	2	3.85	1	1	0	0	0
398366485	ASP1	0	0	0	0	3.67	1	1	0	0	0	0	0	0
398365871	MGE1	0	0	0	0	4.39	1	1	0	0	0	0	0	0
6324690	RPO31	0	0	0	0	1.30	1	3	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6324946	MRS6	0	0	0	0	2.32	1	4	0	0	0	0	0	0
6325052	HRR25	0	0	0	0	4.66	1	1	0	0	0	0	0	0
6324631	SGT1	0	0	0	0	1.77	1	2	0	0	0	0	0	0
6323651	ADI1	0	0	0	0	7.26	1	2	0	0	0	0	0	0
398365629	DBF2	0	0	0	0	2.27	1	2	0	0	0	0	0	0
6320725	GRH1	0	0	0	0	3.49	1	2	0	0	0	0	0	0
6324056	TOF1	0	0	0	0	1.21	1	1	0	0	0	0	0	0
6321823	PIH1	0	0	0	0	6.98	1	2	0	0	0	0	0	0
10383767	KCC4	0	0	0	0	1.74	1	1	0	0	0	0	0	0
6319556	UBC4	0	0	0	0	18.24	1	1	0	0	0	0	0	0
6325007	GYP5	0	0	0	0	2.13	1	1	0	0	0	0	0	0
6322121	SEC6	0	0	0	0	1.74	1	1	0	0	0	0	0	0
6323292	TMA7	0	0	0	0	17.19	1	1	0	0	0	0	0	0
6325227	TRM44	0	0	0	0	2.29	1	2	0	0	0	0	0	0
6322526	TOR1	0	0	0	0	0.57	1	1	0	0	0	0	0	0
6324085	SUI1	0	0	0	0	25.93	2	4	0	0	0	0	0	0
398366199	IVY1	0	4.42	1	1	6.84	2	6	0	0	0	0	0	0
6322839	ARC19	0	0	0	0	14.04	2	2	0	0	0	0	0	0
6321914	CIA2	0	0	0	0	11.69	2	6	0	0	0	0	0	0
6319737	RGD1	0	0	0	0	3.45	2	5	0	0	0	0	0	0
6322419	MHP1	0	0	0	0	2.43	2	3	0	0	0	0	0	0
398366245	LDB19	0	0	0	0	2.93	2	4	0	0	0	0	0	0
6325011	YPL245W	0	0	0	0	5.95	2	2	0	0	0	0	0	0
398364697	LSM4	0	0	0	0	9.63	2	3	0	0	0	0	0	0
6320840	FMP52	26	0	0	0	14.72	3	5	0	0	0	0	0	0
6324794	RCN2	0	0	0	0	14.72	3	7	0	0	0	0	0	0
6321800	VPS29	0	0	0	0	13.12	3	7	0	0	0	0	0	0
398364967	SOK1	0	0	0	0	6.99	3	6	0	0	0	0	0	0
6320413	UME6	0	0	0	0	4.43	3	5	0	0	0	0	0	0
6322379	IML2	0	0	0	0	4.51	3	3	0	0	0	0	0	0
6325362	COG4	0	0	0	0	5.92	3	7	0	0	0	0	0	0
6322156	BCY1	0	0	0	0	13.46	4	6	0	0	0	0	0	0
6322424	SNX4	0	0	0	0	15.84	4	7	0	0	0	0	0	0
398365471	MCM7	0	0	0	0	7.93	4	4	0	0	0	0	0	0
398365269	SMY2	0	0	0	0	7.43	4	12	0	0	0	0	0	0
6323616	YML6	0	0	0	0	21.33	5	5	0	0	0	0	0	0
6325343	SUA7	0	0	0	0	24.93	5	23	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6320490	DPP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321749	CBP2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323230	COQ9	0	0	0	0	0	0	0	0	0	0	0	0	0
6323020	COX17	0	0	0	0	0	0	0	0	0	0	0	0	0
6325319	FCY1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323523	DAT1	0	0	0	0	0	0	0	0	0	0	0	0	0
330443382	DEP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322633	URA1	0	0	0	0	0	0	0	0	0	0	0	0	0
6322657	DPH2	0	0	0	0	0	0	0	0	0	0	0	0	0
6321200	HAP2	0	0	0	0	0	0	0	0	0	0	0	0	0
9755326	YBR085C-	0	0	0	0	0	0	0	0	0	0	0	0	0
42759851	YCL012C	0	0	0	0	0	0	0	0	0	0	0	0	0
6323669	YMR027	0	0	0	0	0	0	0	0	0	0	0	0	0
398365291	ACB1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323941	NGL2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323177	PEP3	0	0	0	0	0	0	0	0	0	0	0	0	0
6324678	PIN2	0	0	0	0	0	0	0	0	0	0	0	0	0
14318491	CAF16	0	0	0	0	0	0	0	0	0	0	0	0	0
398365407	RCM1	0	0	0	0	0	0	0	0	0	0	0	0	0
398364993	RPN14	0	0	0	0	0	0	0	0	0	0	0	0	0
6324072	SIP3	0	0	0	0	0	0	0	0	0	0	0	0	0
6324655	TGL5	0	0	0	0	0	0	0	0	0	0	0	0	0
398366623	TLG1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320100	QRI1	0	0	0	0	0	0	0	0	0	0	0	0	0
6321226	VAM7	0	0	0	0	0	0	0	0	0	0	0	0	0
6322459	PRE3	0	0	0	0	0	0	0	0	0	0	0	0	0
6322083	YIL108W	0	0	0	0	0	0	0	0	0	0	0	0	0
398366489	UTP4	0	0	0	0	0	0	0	0	0	0	0	0	0
6325194	TIM50	0	0	0	0	0	0	0	0	0	0	0	0	0
398365181	HOC1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320380	RSM24	0	0	0	0	0	0	0	0	0	0	0	0	0
330443750	YPR097W	0	0	0	0	4.01	2	5	1.58	1	1	0	0	0
6324984	ATP15	1	0	0	0	27.42	1	1	0	0	0	0	0	0
6323999	RPD3	0	0	0	0	3.70	1	1	0	0	0	0	0	0
6320974	LCP5	0	0	0	0	5.04	1	1	0	0	0	0	0	0
6324552	DIS3	0	0	0	0	1.60	1	3	0	0	0	0	0	0
398365639	PBP2	0	0	0	0	3.39	1	5	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6325146	CAR1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319895	BUD23	0	0	0	0	0	0	0	0	0	0	0	0	0
398365793	SHM1	2	0	0	0	2.86	1	3	0	0	0	0	0	0
398365279	ADE2	0	0	0	0	2.28	1	1	0	0	0	0	0	0
6322564	SOD1	1	0	0	0	6.49	1	1	0	0	0	0	0	0
6323931	BUL1	0	0	0	0	2.36	1	1	0	0	0	0	0	0
6322714	APL2	0	0	0	0	4.96	3	5	0	0	0	0	0	0
398365723	DPB4	0	0	0	0	0	0	0	0	0	0	0	0	0
6324250	TPM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324382	ESF2	0	0	0	0	0	0	0	0	0	0	0	0	0
398364299	NOP16	1	0	0	0	0	0	0	0	0	0	0	0	0
398364461	HMF1	0	0	0	0	0	0	0	0	0	0	0	0	0
330443644	APE2	0	0	0	0	0	0	0	0	0	0	0	0	0
6323951	JNM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6323103	BUD20	1	0	0	0	7.23	1	1	0	0	0	0	0	0
6324245	END3	2	0	0	0	29.51	6	23	0	0	0	0	0	0
398365535	GLN4	4	0	0	0	0	0	0	0	0	0	0	0	0
6321829	VMA10	0	0	0	0	9.65	1	2	0	0	0	0	0	0
6323340	IMH1	0	0	0	0	16.25	11	17	0	0	0	0	0	0
330443364	ECM1	0	0	0	0	0	0	0	0	0	0	0	0	0
6324601	STI1	2	0	0	0	3.06	1	1	0	0	0	0	0	0
6321563	ASN2	2	0	0	0	0	0	0	0	0	0	0	0	0
6323779	RRB1	0	0	0	0	0	0	0	0	0	0	0	0	0
6320306	ARX1	1	0	0	0	3.04	1	2	0	0	0	5.40	1	1
398365749	MTC5	0	0	0	0	0	0	0	0	0	0	0	0	0
6320152	LHP1	0	0	0	0	0	0	0	0	0	0	0	0	0
6319926	TUP1	0	0	0	0	3.23	2	4	0	0	0	0	0	0
398364983	EGD2	8	0	0	0	20.69	2	5	0	0	0	7.47	1	1
398366421	GDH1	1	0	0	0	10.79	3	3	0	0	0	0	0	0
6325421	TIF3	0	0	0	0	0	0	0	0	0	0	0	0	0
339046261	NHP6BP	0	0	0	0	13.13	1	1	0	0	0	0	0	0
6325229	ERG10	0	0	0	0	9.80	3	3	0	0	0	0	0	0
398366043	CAF20	0	0	0	0	42.86	3	9	0	0	0	0	0	0
6325014	IQG1	0	0	0	0	1.40	1	1	0	0	0	0.54	1	2
398365281	BUD4	0	0	0	0	0.76	1	2	0	0	0	0.83	1	1
398365669	TEL2	0	0	0	0	0	0	0	0	0	0	1.02	1	1
10383780	ADP1	0	0	0	0	0	0	0	0	0	0	1.05	1	1

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6323181	YLR152C	0	0	0	0	0	0	0	0	0	0	1.22	1	1
6323217	MDL1	0	0	0	0	0	0	0	0	0	0	1.44	1	1
6321439	SWC4	0	0	0	0	3.78	1	3	0	0	0	1.68	1	1
6325116	FRK1	0	0	0	0	0	0	0	0	0	0	1.97	1	1
6321223	CLG1	0	0	0	0	0	0	0	0	0	0	2.21	1	1
398364759	PRP40	0	0	0	0	0	0	0	0	0	0	2.23	1	1
330443403	TEL1	0	0	0	0	0	0	0	0	0	0	2.26	2	2
6323724	CTF18	0	0	0	0	0	0	0	0	0	0	2.43	1	1
6324482	MSH2	0	0	0	0	0	0	0	0	0	0	2.90	1	1
6322471	YJR011C	0	0	0	0	0	0	0	0	0	0	3.07	1	17
6322075	HIS5	1	0	0	0	12.21	4	7	0	0	0	3.64	1	1
6323435	SFP1	0	0	0	0	10.10	3	6	0	0	0	5.27	1	1
330443407	PET112	0	0	0	0	0	0	0	0	0	0	5.91	1	1
6325210	SGF11	0	0	0	0	0	0	0	0	0	0	7.07	1	1
6320137	IDP1	0	0	0	0	2.57	1	1	0	0	0	8.41	3	3
6323640	YML003W	0	0	0	0	0	0	0	0	0	0	10.69	1	1
6321215	COG1	0	0	0	0	2.88	1	1	0	0	0	11.51	2	2
6323826	DDR48	0	0	0	0	0	0	0	0	0	0	22.33	3	4
398365503	RSF2	0	0	0	0	0.80	1	2	0.43	1	1	0	0	0
398366065	VAC14	0	0	0	0	2.27	1	1	2.16	1	1	0	0	0
398365047	DAS2	0	0	0	0	13.79	2	2	8.19	1	1	0	0	0
6324235	APP1	0	0	0	0	9.54	3	3	1.02	1	1	0	0	0
6321557	NUP57	1	0	0	0	7.58	4	6	1.85	1	1	0	0	0
6320775	PRB1	0	0	0	0	0	0	0	2.05	1	1	0	0	0
6325212	VPS16	0	0	0	0	0	0	0	1.75	1	1	0	0	0
6321418	GET1	0	0	0	0	0	0	0	2.98	1	1	0	0	0
6323517	NGL3	0	0	0	0	0	0	0	1.98	1	3	0	0	0
6325228	SUV3	0	0	0	0	0	0	0	1.22	1	1	0	0	0
398364413	PCL7	0	0	0	0	0	0	0	4.21	1	2	0	0	0
6322556	YJR096W	0	0	0	0	0	0	0	11.35	1	1	0	0	0
6322804	ANR2	0	0	0	0	0	0	0	1.94	1	1	0	0	0
6323271	ARV1	0	0	0	0	0	0	0	10.90	1	1	0	0	0
6324506	RIB2	0	0	0	0	0	0	0	3.05	1	1	0	0	0
6320046	CLB3	0	0	0	0	0	0	0	3.51	1	1	0	0	0
82795250	YGL194C-	0	0	0	0	0	0	0	26.25	1	1	0	0	0
398365389	YGR053C	0	0	0	0	0	0	0	3.18	1	1	0	0	0
33438858	YMR230	0	0	0	0	0	0	0	20.97	1	1	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6325279	YPRO22C	0	0	0	0	0	0	0	0.79	1	1	0	0	0
6321028	ISC10	0	0	0	0	0	0	0	3.75	1	1	0	0	0
6321274	YRB30	0	0	0	0	0	0	0	1.59	1	1	0	0	0
6324519	AIM39	0	0	0	0	0	0	0	2.78	1	1	0	0	0
6322658	CNB1	3	0	0	0	10.29	1	2	0	0	0	0	0	0
10383760	LSB5	0	0	0	0	6.50	1	1	0	0	0	0	0	0
398364999	CKS1	0	0	0	0	10.00	1	1	0	0	0	0	0	0
398366627	TRS31	0	0	0	0	8.83	1	1	0	0	0	0	0	0
6324672	NUP1	11	0	0	0	1.21	1	1	0	0	0	0	0	0
6323031	CMS1	0	0	0	0	5.15	1	6	0	0	0	0	0	0
6323287	GSY2	0	0	0	0	2.70	1	1	0	0	0	0	0	0
6323425	ATP10	0	0	0	0	7.53	1	3	0	0	0	0	0	0
398365917	RSR1	0	0	0	0	4.04	1	2	0	0	0	0	0	0
6321849	MED6	0	0	0	0	5.42	1	7	0	0	0	0	0	0
6322281	ATP12	0	0	0	0	3.69	1	5	0	0	0	0	0	0
6319867	MAK31	0	0	0	0	10.23	1	4	0	0	0	0	0	0
6321785	HSE1	0	0	0	0	3.54	1	4	0	0	0	0	0	0
6325087	DAP1	0	0	0	0	8.55	1	2	0	0	0	0	0	0
6324502	NBA1	0	0	0	0	3.99	1	1	0	0	0	0	0	0
6323019	PSR1	0	0	0	0	3.04	1	7	0	0	0	0	0	0
398364635	IXR1	7	0	0	0	1.84	1	2	0	0	0	0	0	0
6322996	YLL032C	0	0	0	0	1.70	1	4	0	0	0	0	0	0
6323533	NUP188	0	0	0	0	0.91	1	3	0	0	0	0	0	0
6322337	LSM1	0	0	0	0	9.30	1	2	0	0	0	0	0	0
6321906	BZZ1	0	0	0	0	2.84	1	2	0	0	0	0	0	0
6320163	SIR2	0	0	0	0	3.38	1	5	0	0	0	0	0	0
398364637	PRS2	0	0	0	0	5.03	1	2	0	0	0	0	0	0
398364293	FMC1	4	0	0	0	6.45	1	1	0	0	0	0	0	0
6325113	POC4	0	0	0	0	10.14	1	1	0	0	0	0	0	0
6323162	CKI1	0	0	0	0	1.89	1	1	0	0	0	0	0	0
6320127	RXT3	0	0	0	0	3.40	1	1	0	0	0	0	0	0
330443546	MDS3	0	0	0	0	1.08	1	4	0	0	0	0	0	0
398366201	ATG17	0	0	0	0	2.88	1	2	0	0	0	0	0	0
6324934	HAP5	0	0	0	0	4.96	1	2	0	0	0	0	0	0
398364577	DOT6	0	0	0	0	1.79	1	1	0	0	0	0	0	0
6324023	YMS18	0	0	0	0	5.07	1	1	0	0	0	0	0	0
37362669	ESS1	0	0	0	0	8.82	1	1	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6320537	UBX5	0	0	0	0	2.40	1	1	0	0	0	0	0	0
398366051	REF2	0	0	0	0	2.63	1	1	0	0	0	0	0	0
93117371	PEX19	0	0	0	0	4.68	1	1	0	0	0	0	0	0
6323146	CLF1	0	0	0	0	2.04	1	5	0	0	0	0	0	0
6322990	ENT4	0	0	0	0	5.67	1	5	0	0	0	0	0	0
6320990	MAG1	0	0	0	0	6.42	1	5	0	0	0	0	0	0
398365849	EKI1	0	0	0	0	1.69	1	4	0	0	0	0	0	0
6322662	MTR2	0	0	0	0	9.24	1	3	0	0	0	0	0	0
6320507	CFT1	0	0	0	0	1.33	1	3	0	0	0	0	0	0
6325155	ELP4	1	0	0	0	2.41	1	2	0	0	0	0	0	0
6319894	IMG1	0	0	0	0	9.47	1	2	0	0	0	0	0	0
6321705	YGR266W	0	0	0	0	1.43	1	2	0	0	0	0	0	0
14318554	SMC2	0	0	0	0	1.45	1	2	0	0	0	0	0	0
398366369	VTS1	0	0	0	0	4.02	1	1	0	0	0	0	0	0
6325307	MAK3	0	0	0	0	9.09	1	1	0	0	0	0	0	0
398364521	NAS2	0	0	0	0	4.55	1	1	0	0	0	0	0	0
6323222	UPS1	0	0	0	0	5.14	1	1	0	0	0	0	0	0
6321924	ECM14	0	0	0	0	3.26	1	4	0	0	0	0	0	0
398365805	MSO1	0	0	0	0	5.71	1	4	0	0	0	0	0	0
6322100	UTP25	0	1.94	1	1	2.08	1	2	0	0	0	0	0	0
398365089	FKH2	0	0	0	0	3.25	1	2	0	0	0	0	0	0
398364225	IMP2'P	0	0	0	0	4.62	1	2	0	0	0	0	0	0
6320643	PPM1	0	0	0	0	4.88	1	2	0	0	0	0	0	0
6319436	POL12	0	0	0	0	2.70	1	1	0	0	0	0	0	0
6322334	SPT10	0	0	0	0	4.06	1	1	0	0	0	0	0	0
6319631	CNS1	0	0	0	0	4.68	1	1	0	0	0	0	0	0
6320050	RPC53	0	0	0	0	2.13	1	1	0	0	0	0	0	0
6321713	TAF1	0	0	0	0	1.13	1	1	0	0	0	0	0	0
6325118	UME1	0	0	0	0	3.70	1	1	0	0	0	0	0	0
6321493	RSC1	0	0	0	0	2.48	1	1	0	0	0	0	0	0
6323274	CDD1	0	0	0	0	8.45	1	3	0	0	0	0	0	0
398364583	YGL082W	0	0	0	0	5.51	1	3	0	0	0	0	0	0
398364535	YNL165W	0	0	0	0	2.71	1	3	0	0	0	0	0	0
398365015	AIM46	1	0	0	0	4.52	1	2	0	0	0	0	0	0
6325183	YTA6	0	0	0	0	1.33	1	2	0	0	0	0	0	0
6324128	PSY2	0	0	0	0	1.28	1	2	0	0	0	0	0	0
6320259	CDC34	0	0	0	0	5.42	1	2	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
398364911	SPT15	0	0	0	0	2.92	1	2	0	0	0	0	0	0
330443461	COS111	1	0.87	1	1	1.52	1	1	0	0	0	0	0	0
6320652	YDR444W	0	0	0	0	2.18	1	1	0	0	0	0	0	0
398365541	UBC13	0	0	0	0	12.42	1	1	0	0	0	0	0	0
398365707	GPX2	0	0	0	0	8.64	1	1	0	0	0	0	0	0
398365979	GTR2	0	0	0	0	2.64	1	1	0	0	0	0	0	0
6323528	YML108W	0	0	0	0	9.52	1	1	0	0	0	0	0	0
6321894	KIC1	0	0	0	0	1.11	1	1	0	0	0	0	0	0
398364893	PTC1	0	0	0	0	6.05	1	1	0	0	0	0	0	0
398366573	SIZ1	0	0	0	0	1.55	1	1	0	0	0	0	0	0
398366291	TEA1	0	0	0	0	3.16	1	1	0	0	0	0	0	0
398366471	TFB1	0	0	0	0	1.40	1	1	0	0	0	0	0	0
6325426	NUT2	0	0	0	0	5.10	1	1	0	0	0	0	0	0
154199620	CGI121	0	0	0	0	4.97	1	1	0	0	0	0	0	0
6320156	SIT4	0	0	0	0	6.43	1	1	0	0	0	0	0	0
398366599	APT2	0	0	0	0	6.08	1	2	0	0	0	0	0	0
6324668	ARF3	0	0	0	0	8.74	1	2	0	0	0	0	0	0
6324087	ATG2	0	0	0	0	1.13	1	2	0	0	0	0	0	0
6323143	AVL9	0	0	0	0	2.23	1	2	0	0	0	0	0	0
398364821	BYE1	0	0	0	0	2.19	1	2	0	0	0	0	0	0
6322088	DPH1	0	0	0	0	2.59	1	2	0	0	0	0	0	0
6323972	ELP6	0	0	0	0	4.03	1	2	0	0	0	0	0	0
398364641	FLO11	0	0	0	0	1.32	1	2	0	0	0	0	0	0
6322659	HYM1	0	0	0	0	3.01	1	2	0	0	0	0	0	0
6322040	YIL151C	0	0	0	0	0.98	1	2	0	0	0	0	0	0
398365375	YKR078W	0	0	0	0	1.88	1	2	0	0	0	0	0	0
6323758	YMR111C	0	0	0	0	6.49	1	2	0	0	0	0	0	0
6323793	YMR144	0	0	0	0	4.09	1	2	0	0	0	0	0	0
6320208	MAF1	0	0	0	0	5.32	1	2	0	0	0	0	0	0
6324437	MED7	0	0	0	0	7.21	1	2	0	0	0	0	0	0
6323880	MRE11	0	0	0	0	1.45	1	2	0	0	0	0	0	0
6323680	MSN2	0	0	0	0	2.27	1	2	0	0	0	0	0	0
6324503	NUF2	0	0	0	0	3.10	1	2	0	0	0	0	0	0
6323025	ORC3	0	0	0	0	2.11	1	2	0	0	0	0	0	0
6323147	YLR118C	0	0	0	0	6.61	1	2	0	0	0	0	0	0
6322733	PRR1	0	0	0	0	2.51	1	2	0	0	0	0	0	0
6319504	RKM3	0	0	0	0	2.17	1	2	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6321876	STE12	0	0	0	0	2.47	1	2	0	0	0	0	0	0
6324836	GPN2	0	6.34	1	1	5.48	1	1	0	0	0	0	0	0
6321605	TRS65	1	0	0	0	2.86	1	1	0	0	0	0	0	0
6324138	DUG3	0	0	0	0	4.48	1	1	0	0	0	0	0	0
6319940	HMRA1	0	0	0	0	6.35	1	1	0	0	0	0	0	0
330443533	YFL034W	0	0	0	0	0.84	1	1	0	0	0	0	0	0
6323815	INP2	0	0	0	0	2.13	1	1	0	0	0	0	0	0
330443515	PSP1	0	0	0	0	1.78	1	1	0	0	0	0	0	0
6323711	SOV1	0	0	0	0	1.78	1	1	0	0	0	0	0	0
330443539	FAB1	0	0	0	0	0.83	1	1	0	0	0	0	0	0
398365149	ICS2	0	0	0	0	7.06	1	1	0	0	0	0	0	0
6323764	SPC24	0	0	0	0	5.63	1	1	0	0	0	0	0	0
6325028	CET1	0	0	0	0	3.46	1	1	0	0	0	0	0	0
6322081	HPM1	0	0	0	0	4.77	1	1	0	0	0	0	0	0
6322195	IST3	0	0	0	0	9.46	1	1	0	0	0	0	0	0
398366647	RSM28	0	0	0	0	3.88	1	1	0	0	0	0	0	0
6320113	RAM1	0	0	0	0	3.71	1	1	0	0	0	0	0	0
6323522	TAF8	0	0	0	0	3.92	1	1	0	0	0	0	0	0
6323874	TRS130	0	0	0	0	0.73	1	1	0	0	0	0	0	0
6323456	SPP382	0	0	0	0	1.69	1	1	0	0	0	0	0	0
6324386	BIO3	0	0	0	0	2.92	1	1	0	0	0	0	0	0
6324228	AVT4	0	0	0	0	4.77	1	1	0	0	0	0	0	0
398364711	RAM2	0	0	0	0	4.11	1	1	0	0	0	0	0	0
14318494	BST1	0	0	0	0	0.97	1	1	0	0	0	0	0	0
6320579	CTS2	0	0	0	0	2.15	1	1	0	0	0	0	0	0
6322760	CUE2	0	0	0	0	3.84	1	1	0	0	0	0	0	0
6320023	DLD2	0	0	0	0	5.28	1	1	0	0	0	0	0	0
6323060	RAD5	0	0	0	0	1.03	1	1	0	0	0	0	0	0
6319754	DPB3	0	0	0	0	3.98	1	1	0	0	0	0	0	0
6319994	GLE1	0	0	0	0	2.42	1	1	0	0	0	0	0	0
6323950	HER2	0	0	0	0	3.66	1	1	0	0	0	0	0	0
398366409	GPB1	0	0	0	0	0.89	1	1	0	0	0	0	0	0
6319390	YBL081W	0	0	0	0	3.80	1	1	0	0	0	0	0	0
398364947	YDR003W	0	0	0	0	32.50	1	1	0	0	0	0	0	0
6323384	YLR352W	0	0	0	0	2.35	1	1	0	0	0	0	0	0
6324536	YOL036W	0	0	0	0	1.84	1	1	0	0	0	0	0	0
6325180	YPL077C	0	0	0	0	4.17	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6320328	INO2	0	0	0	0	3.95	1	1	0	0	0	0	0	0
6324064	IST1	0	0	0	0	3.36	1	1	0	0	0	0	0	0
6321196	KAP114	0	0	0	0	1.49	1	1	0	0	0	0	0	0
398366685	KRE28	0	0	0	0	3.64	1	1	0	0	0	0	0	0
398364391	KRE29	0	0	0	0	3.88	1	1	0	0	0	0	0	0
6320055	LDB17	0	0	0	0	3.67	1	1	0	0	0	0	0	0
6321950	LIN1	0	0	0	0	4.71	1	1	0	0	0	0	0	0
6325219	MET31	0	0	0	0	9.60	1	1	0	0	0	0	0	0
6320158	YMS-T	0	0	0	0	9.47	1	1	0	0	0	0	0	0
6323472	YML4	0	0	0	0	4.08	1	1	0	0	0	0	0	0
6324640	MSA1	0	0	0	0	3.18	1	1	0	0	0	0	0	0
6319938	OCA4	0	0	0	0	3.04	1	1	0	0	0	0	0	0
14318475	OTU1	0	0	0	0	6.64	1	1	0	0	0	0	0	0
6324012	PFS2	0	0	0	0	1.94	1	1	0	0	0	0	0	0
398366005	PLP1	0	0	0	0	6.96	1	1	0	0	0	0	0	0
6321904	YHR112C	0	0	0	0	7.41	1	1	0	0	0	0	0	0
6323631	RAD33	0	0	0	0	5.65	1	1	0	0	0	0	0	0
6323930	RCE1	0	0	0	0	8.57	1	1	0	0	0	0	0	0
6324701	RGA1	0	0	0	0	1.69	1	1	0	0	0	0	0	0
6323919	SAP30	0	0	0	0	4.98	1	1	0	0	0	0	0	0
6322536	CDC11	0	0	0	0	3.13	1	1	0	0	0	0	0	0
6319410	SKT5	0	0	0	0	1.29	1	1	0	0	0	0	0	0
6321311	SOH1	0	0	0	0	12.60	1	1	0	0	0	0	0	0
398364839	SLX5	0	0	0	0	1.94	1	1	0	0	0	0	0	0
330443427	TCM62	0	0	0	0	2.45	1	1	0	0	0	0	0	0
6324684	TFC7	0	0	0	0	3.91	1	1	0	0	0	0	0	0
398364235	TID3	0	0	0	0	2.17	1	1	0	0	0	0	0	0
6325297	TIP41	0	0	0	0	3.65	1	1	0	0	0	0	0	0
6320112	UBX3	0	0	0	0	3.30	1	1	0	0	0	0	0	0
6324852	HEM4	0	0	0	0	5.45	1	1	0	0	0	0	0	0
398366151	UTP21	0	0	0	0	1.17	1	1	0	0	0	0	0	0
6323405	VID22	0	0	0	0	2.11	1	1	0	0	0	0	0	0
398366111	VPS33	0	0	0	0	1.74	1	1	0	0	0	0	0	0
398366641	VPS60	0	0	0	0	8.73	1	1	0	0	0	0	0	0
398364823	YAF9	0	0	0	0	5.31	1	1	0	0	0	0	0	0
27469361	YCG1	0	0	0	0	1.35	1	1	0	0	0	0	0	0
14318529	YFH7	0	0	0	0	3.40	1	1	0	0	0	0	0	0

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Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6322153	CST6	11	0	0	0	3.92	1	3	0	0	0	0	0	0
14318524	NIC96	0	0	0	0	1.07	1	1	0	0	0	0	0	0
6324402	AIF1	0	0	0	0	5.03	1	3	0	0	0	0	0	0
398364507	DOT5	0	0	0	0	4.19	1	2	0	0	0	0	0	0
398365867	ENT5	0	0	0	0	4.14	1	2	0	0	0	0	0	0
6324130	GCR2	1	0	0	0	2.62	1	4	0	0	0	0	0	0
6323148	SRN2	0	0	0	0	6.10	1	1	0	0	0	0	0	0
6319738	TAE1	0	0	0	0	3.88	1	1	0	0	0	0	0	0
6324656	WHI5	0	0	0	0	2.71	1	4	0	0	0	0	0	0
6324781	RET1	0	0	0	0	1.04	1	2	0	0	0	0	0	0
37362668	YJL016W	0	0	0	0	2.14	1	1	0	0	0	0	0	0
6321609	PSD2	0	0	0	0	2.20	1	1	0	0	0	0	0	0
6321481	RME1	0	0	0	0	3.67	1	1	0	0	0	0	0	0
6325431	VPS4	0	0	0	0	3.89	1	1	0	0	0	0	0	0
6324854	FSH3	0	0	0	0	6.39	1	1	0	0	0	0	0	0
330443376	MAK16	0	0	0	0	4.58	1	1	0	0	0	0	0	0
6322690	RCN1	0	0	0	0	7.11	1	5	0	0	0	0	0	0
6322431	VPS53	0	0	0	0	1.70	1	1	0	0	0	0	0	0
6322631	SRY1	0	0	0	0	4.91	1	1	0	0	0	0	0	0
6320580	VPS74	0	0	0	0	9.57	2	3	0	0	0	0	0	0
9755336	DID4	0	0	0	0	9.91	2	2	0	0	0	0	0	0
6323881	YML44	0	0	0	0	27.55	2	4	0	0	0	0	0	0
398366509	MRP1	0	0	0	0	9.66	2	3	0	0	0	0	0	0
6320035	FAP7	0	0	0	0	12.69	2	5	0	0	0	0	0	0
6323354	SFH1	0	0	0	0	7.28	2	13	0	0	0	0	0	0
6323531	SEC65	0	0	0	0	9.16	2	6	0	0	0	0	0	0
6321433	COG7	0	0	0	0	9.32	2	4	0	0	0	0	0	0
398364865	PIB2	0	2.68	1	1	5.67	2	12	0	0	0	0	0	0
6320673	RMT2	0	0	0	0	5.10	2	7	0	0	0	0	0	0
6323417	SWC7	0	0	0	0	19.70	2	2	0	0	0	0	0	0
6324423	DCP1	0	0	0	0	9.09	2	6	0	0	0	0	0	0
398365921	TAF10	1	0	0	0	20.87	2	4	0	0	0	0	0	0
6323712	UBX4	0	0	0	0	5.77	2	9	0	0	0	0	0	0
6325166	GLR1	0	0	0	0	6.83	2	3	0	0	0	0	0	0
6320994	LSM5	0	0	0	0	26.88	2	2	0	0	0	0	0	0
398366515	TRP4	0	0	0	0	7.89	2	4	0	0	0	0	0	0
6324551	TSR4	0	0	0	0	6.62	2	2	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6324068	ORC5	0	0	0	0	5.22	2	5	0	0	0	0	0	0
6325213	NOP4	0	0	0	0	5.11	2	2	0	0	0	0	0	0
6324041	CAF40	0	0	0	0	6.43	2	6	0	0	0	0	0	0
6320086	CYK3	0	0	0	0	3.62	2	6	0	0	0	0	0	0
6321762	SNF6	0	12.95	1	1	10.84	2	5	0	0	0	0	0	0
398365087	GRX4	0	0	0	0	8.20	2	5	0	0	0	0	0	0
6323218	ATG26	0	0	0	0	2.59	2	5	0	0	0	0	0	0
41629693	PRP18	0	0	0	0	7.97	2	5	0	0	0	0	0	0
330443528	PTP3	0	0	0	0	3.66	2	5	0	0	0	0	0	0
398364733	YIR035C	0	0	0	0	10.24	2	2	0	0	0	0	0	0
6321766	AIM17	0	0	0	0	6.67	2	4	0	0	0	0	0	0
14318483	MOB2	0	0	0	0	10.10	2	4	0	0	0	0	0	0
6324527	PSK2	0	0	0	0	2.72	2	4	0	0	0	0	0	0
398365421	PRP16	0	0	0	0	2.89	2	3	0	0	0	0	0	0
41629685	RSC30	0	0	0	0	2.04	2	3	0	0	0	0	0	0
6320907	VHR2	0	0	0	0	4.75	2	2	0	0	0	0	0	0
398366405	GCN2	0	0	0	0	1.57	2	3	0	0	0	0	0	0
398364549	GYP6	0	0	0	0	5.02	2	3	0	0	0	0	0	0
330443696	YNL193W	0	0	0	0	5.38	2	3	0	0	0	0	0	0
6325243	YPL014W	0	0	0	0	9.97	2	3	0	0	0	0	0	0
6325327	MED1	0	0	0	0	5.12	2	3	0	0	0	0	0	0
6320160	MTF2	0	0	0	0	5.23	2	3	0	0	0	0	0	0
6319287	PRP45	0	0	0	0	6.86	2	3	0	0	0	0	0	0
398365919	SEC5	0	0	0	0	2.78	2	3	0	0	0	0	0	0
6320433	SIR4	0	0	0	0	2.36	2	2	0	0	0	0	0	0
6321725	BIO2	0	0	0	0	9.60	2	2	0	0	0	0	0	0
6324177	INN1	0	0	0	0	6.11	2	2	0	0	0	0	0	0
6319534	ORC2	0	0	0	0	4.35	2	2	0	0	0	0	0	0
6320887	SAP1	0	0	0	0	3.34	2	2	0	0	0	0	0	0
6321204	SEC15	0	0	0	0	2.75	2	2	0	0	0	0	0	0
6323111	SRL2	0	0	0	0	6.63	2	2	0	0	0	0	0	0
398366601	SSN2	0	0	0	0	2.32	2	2	0	0	0	0	0	0
6323541	VPS9	0	0	0	0	7.76	2	2	0	0	0	0	0	0
6321955	YAP1801	0	0	0	0	4.87	2	5	0	0	0	0	0	0
6323215	EMG1	0	0	0	0	13.10	2	3	0	0	0	0	0	0
6319868	HTL1	0	0	0	0	33.33	2	5	0	0	0	0	0	0
6321537	MDR1	0	0	0	0	2.84	2	3	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	# PSM TAP_Rtr1_ WT_1	%SC TAP_Rtr1_ WT_2	UP TAP_Rtr1_ WT_2	# PSM TAP_Rtr1_ WT_2	%SC TAP_Rtr1_ WT_3	UP TAP_Rtr1_ WT_3	# PSM TAP_Rtr1_ WT_3	%SC TAP_Rtr1_ WT_4	UP TAP_Rtr1_ WT_4	# PSM TAP_Rtr1_ WT_4	%SC TAP_Rtr1_ ctk1_del etion	UP TAP_Rtr1_ ctk1_del etion	# PSM TAP_Rtr1_ ctk1_del etion
6321941	YML16	0	0	0	0	10.75	2	2	0	0	0	0	0	0
398365813	POP2	0	0	0	0	4.85	2	2	0	0	0	0	0	0
6322855	OSH6	12	0	0	0	4.69	2	4	0	0	0	0	0	0
6319407	PRX1	0	0	0	0	14.94	3	7	0	0	0	0	0	0
398366397	SCP1	0	0	0	0	22.00	3	7	0	0	0	0	0	0
6325247	RET3	0	0	0	0	21.16	3	7	0	0	0	0	0	0
6322574	RSM7	1	0	0	0	15.79	3	3	0	0	0	0	0	0
6321831	SRB2	0	0	0	0	23.81	3	7	0	0	0	0	0	0
6324815	MET7	0	0	0	0	8.21	3	7	0	0	0	0	0	0
6322771	SMY1	0	0	0	0	7.16	3	8	0	0	0	0	0	0
398365395	MED8	0	0	0	0	19.73	3	11	0	0	0	0	0	0
16740527	ATG18	0	0	0	0	9.40	3	8	0	0	0	0	0	0
6320526	SWA2	0	0	0	0	7.49	3	10	0	0	0	0	0	0
6319430	PRE7	0	0	0	0	17.84	3	9	0	0	0	0	0	0
6323626	TAF11	0	0	0	0	10.69	3	4	0	0	0	0	0	0
6319558	MIS1	0	0	0	0	4.10	3	3	0	0	0	0	0	0
398366353	SOG2	0	0	0	0	5.94	3	6	0	0	0	0	0	0
6321377	DUO1	0	0	0	0	18.22	3	3	0	0	0	0	0	0
6319820	BIK1	0	0	0	0	10.00	3	6	0	0	0	0	0	0
6325187	MUK1	0	0	0	0	7.35	3	6	0	0	0	0	0	0
6320957	SWI4	0	0	0	0	4.85	3	6	0	0	0	0	0	0
398366093	EBS1	0	0	0	0	8.03	3	5	0	0	0	0	0	0
398364617	RAD51	0	0	0	0	12.75	3	3	0	0	0	0	0	0
398365445	BEM1	0	0	0	0	10.89	3	4	0	0	0	0	0	0
6324824	CLP1	0	0	0	0	10.79	3	4	0	0	0	0	0	0
6323772	YMR124	0	0	0	0	4.14	3	4	0	0	0	0	0	0
6325420	ORC4	0	0	0	0	9.07	3	4	0	0	0	0	0	0
398366267	SCD5	0	0	0	0	5.28	3	4	0	0	0	0	0	0
6324327	DOM34	0	0	0	0	10.62	3	9	0	0	0	0	0	0
6323429	AFG2	0	0	0	0	6.92	3	3	0	0	0	0	0	0
398365705	MCA1	0	0	0	0	13.89	4	21	0	0	0	0	0	0
398366069	HGH1	0	0	0	0	14.21	4	14	0	0	0	0	0	0
6325299	PUF2	0	0	0	0	6.23	4	17	0	0	0	0	0	0
6321269	SUA5	0	0	0	0	17.61	4	14	0	0	0	0	0	0
6319902	CTR86	0	0	0	0	7.10	4	9	0	0	0	0	0	0
398366309	SIR3	0	0	0	0	5.21	4	7	0	0	0	0	0	0
398366425	HRQ1	1	0	0	0	4.92	4	6	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398364847	RTR1	47.79	12	728	48.67	10	465	45.58	11	680	43.81	7	20
6320061	RPB1	21.00	33	863	13.21	18	124	19.91	27	406	65.67	128	14665
398365417	RPB2	18.63	20	360	8.33	8	54	16.34	16	167	73.20	105	14108
398364469	RPB3	29.25	6	84	9.12	2	6	35.85	7	42	74.53	26	5873
6322321	RPB4	28.51	3	5	4.98	1	2	0	0	0	82.81	18	4185
6319630	RPB5	20.00	3	30	13.02	2	7	21.86	3	30	63.26	17	3182
398366567	RPB7	7.60	1	2	0	0	0	0	0	0	87.13	13	2096
398365775	RPB10	0	0	0	0	0	0	0	0	0	68.57	6	1373
6325445	RPB6	0	0	0	14.19	1	2	14.19	1	3	87.10	12	1289
6321368	RPB9	20.49	2	5	0	0	0	0	0	0	71.31	8	1230
6324798	RPB8	25.34	3	51	0	0	0	17.12	2	7	70.55	13	944
6324569	RPB11	36.67	4	5	15.00	1	1	19.17	2	2	94.17	15	895
6323632	SPT5	10.91	8	16	0	0	0	0	0	0	61.71	52	636
6321625	TFG1	5.99	3	4	0	0	0	0	0	0	47.07	35	363
6321937	RPB12	38.57	2	21	0	0	0	28.57	1	44	52.86	4	356
398365043	TFG2	3.00	1	1	0	0	0	0	0	0	36.25	16	344
154199603	IWR1	0	0	0	0	0	0	0	0	0	43.06	12	160
398365457	SPT4	0	0	0	0	0	0	0	0	0	65.69	3	130
6325128	TAF14	6.56	1	1	0	0	0	0	0	0	35.25	6	77
398365155	NPA3	11.43	4	4	4.16	1	1	9.87	3	34	0	0	0
6323272	GPN3	0	0	0	4.78	1	1	5.15	1	1	0	0	0
6322163	IRR1	0.87	1	1	1.04	1	1	0	0	0	0	0	0
398366099	RRP41	4.07	1	1	0	0	0	0	0	0	3.66	1	1
398366555	UBA2	0	0	0	0	0	0	0	0	0	0	0	0
6323670	TAP42	23.77	6	127	6.01	2	37	15.85	5	43	0	0	0
6323004	SSA2	41.16	21	1833	45.07	19	399	44.13	22	613	38.65	17	63
6319505	RPL4A	6.63	2	66	43.09	12	417	33.98	10	416	44.20	11	82
398364991	RPL4B	6.63	2	66	38.67	11	416	33.43	9	388	39.78	9	76
398365303	RPL43B	19.57	1	49	29.35	3	168	30.43	4	100	43.48	6	67
398364349	RPL2A	4.33	1	194	32.68	6	329	27.17	4	299	26.38	5	62
6325337	TEF1	17.03	10	430	30.13	11	342	28.60	10	297	40.61	12	57
144228166	SSA1	40.97	21	1735	54.52	21	476	43.93	22	606	32.55	15	57
6319559	RPL19A	19.05	3	49	30.16	7	125	23.81	4	118	29.63	7	48
6320065	RPL35A	16.67	2	3	22.50	4	18	19.17	5	43	34.17	5	48
398365603	RPS4A	19.92	4	12	34.87	8	145	24.14	4	80	35.63	8	40
398365321	RPS6B	16.53	3	7	42.80	11	382	16.95	4	26	40.68	9	38
398364655	SSA4	8.88	5	666	9.50	5	170	6.23	4	124	12.77	7	35

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398366603	RPS17B	44.12	5	40	45.59	6	246	31.62	4	85	47.06	7	34
6324533	RPS15	16.90	2	15	30.28	3	171	30.28	3	21	30.28	3	34
6323104	RPL10	12.22	3	3	25.34	5	110	17.19	4	9	18.55	3	33
6319396	SSA3	11.09	6	613	15.56	7	57	8.47	5	234	9.55	6	33
6321786	RPL14B	17.39	4	82	22.46	3	68	13.77	2	34	33.33	5	31
6322847	RPL14A	23.91	5	83	22.46	3	68	13.77	2	34	33.33	5	31
6324670	RPS7A	30.00	6	195	37.37	5	28	13.16	2	15	46.32	7	31
6322668	RPL17A	7.07	1	1	39.67	6	236	34.78	5	136	35.33	5	27
398365285	CAX4	0	0	0	0	0	0	0	0	0	10.88	1	25
6322284	RPL17B	7.07	1	1	39.67	6	235	34.78	5	116	29.89	4	21
398364883	RPS7B	38.95	7	204	24.21	4	49	14.21	2	9	35.79	6	21
6323791	RPL13B	20.60	4	125	29.65	7	227	34.17	6	40	33.17	5	20
6320121	RPL13A	20.60	4	125	29.65	7	227	32.66	5	39	33.17	5	20
6324120	SSB2	51.71	25	2235	35.89	15	232	35.56	15	238	19.25	8	19
6323577	RPS1B	11.37	3	66	44.71	11	295	20.78	5	20	21.57	5	16
6325157	ATG21	0	0	0	0	0	0	0	0	0	3.83	1	16
6323474	RPS1A	11.37	3	32	48.24	12	237	20.78	5	20	16.47	4	15
398364725	RPL23B	5.84	1	1	16.79	2	14	10.95	1	63	31.39	3	15
6319972	SSB1	51.71	25	2462	32.46	14	230	34.26	15	231	15.82	7	15
154200011	RPL20B	20.35	4	106	34.30	5	192	30.81	5	49	29.07	4	15
6322989	UBI4	32.89	2	304	0	0	0	50.00	3	12	32.89	2	15
9755341	RPS16A	29.37	4	201	48.25	6	47	27.27	3	35	28.67	3	14
6319378	RPL32	0	0	0	30.00	4	110	17.69	2	176	17.69	2	14
6325061	APL5	6.01	4	5	0	0	0	0	0	0	1.93	1	14
398364505	RPS3	39.17	8	296	52.92	10	454	21.67	4	39	30.83	5	13
398365351	RPS13	23.84	3	45	51.66	7	277	33.77	4	41	33.11	4	13
6324260	RPL16B	17.17	3	12	19.70	4	59	19.70	4	82	18.69	5	13
398365605	RPS30B	15.87	1	6	15.87	1	8	0	0	0	28.57	3	12
6323481	RPL6B	27.27	5	89	37.50	6	88	12.50	2	55	16.48	2	11
398365377	RPS9B	32.31	7	80	29.74	7	109	12.31	2	20	23.08	6	11
6324265	YDJ1	11.25	3	53	5.87	2	2	11.00	4	24	28.36	5	11
398364745	ATP7	7.47	1	1	0	0	0	0	0	0	40.23	5	11
6321464	RPS25A	21.30	3	6	39.81	5	49	21.30	2	5	14.81	3	10
6323077	RPS0B	8.33	2	5	0	0	0	0	0	0	4.76	1	10
398365071	RPS11A	13.46	2	6	26.28	5	22	14.10	2	48	17.31	2	10
6321653	RPS0A	8.33	2	5	0	0	0	0	0	0	4.76	1	10
6323465	CNA1	2.53	1	1	0	0	0	0	0	0	17.36	6	10

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6321335	RPL28	16.11	2	6	18.79	3	155	17.45	3	22	30.87	5	9
6322505	SSC1	29.05	15	568	15.29	7	37	30.73	13	77	6.27	3	9
6323567	RPL6A	15.34	3	48	31.82	5	77	12.50	2	40	8.52	1	8
6321754	RPL8A	30.47	8	88	53.52	14	355	35.94	10	90	18.36	4	8
6322984	RPL8B	30.47	8	88	50.78	14	377	36.72	11	91	18.36	4	8
6323615	RPS18B	29.45	4	91	41.10	6	120	13.01	2	5	35.62	5	8
398366307	RPA190	20.19	28	190	0.72	1	1	0	0	0	0.54	1	8
6324637	RPL3	20.67	10	27	27.13	8	194	28.94	9	144	11.37	3	7
6320128	RPL31A	0	0	0	38.05	4	18	14.16	1	2	30.97	3	7
398366143	RPL31B	0	0	0	30.09	3	17	14.16	1	2	30.97	3	7
398365761	RPS23A	0	0	0	13.10	2	2	0	0	0	26.21	2	7
6322137	RPL34B	6.61	1	32	6.61	1	49	0	0	0	20.66	2	7
6321315	RPS2	11.02	2	27	44.49	9	159	13.39	2	48	12.60	2	6
6320949	RPS8B	14.00	2	78	34.50	6	280	20.50	3	157	21.00	3	6
6321798	RPL27A	16.91	2	122	37.50	6	97	34.56	5	68	22.06	3	6
6325176	RPS9A	32.99	7	73	21.32	6	99	4.06	1	6	23.35	4	6
86558907	RHR2	24.40	5	114	11.60	2	23	11.20	2	9	15.60	3	6
6322058	RPL16A	21.61	4	10	14.07	3	17	19.60	4	30	18.59	4	6
6324027	RPS19B	50.00	6	230	45.14	5	36	20.14	2	9	24.31	2	6
6323090	RPL22A	18.18	1	29	41.32	2	5	18.18	1	13	32.23	2	6
6320142	RPS29B	17.86	1	1	0	0	0	0	0	0	17.86	1	6
6324807	KIN4	2.88	2	2	0	0	0	0	0	0	2.38	1	6
6321522	RPL11B	12.64	2	26	27.59	4	102	8.05	1	5	13.79	2	5
6322583	RPS5	26.22	6	292	21.33	4	331	11.56	2	20	12.44	2	5
6325126	RPL5	7.74	2	9	21.89	4	34	21.89	4	29	10.77	2	5
6323073	PDC1	27.35	12	175	36.41	12	388	29.66	10	190	6.04	2	5
398366277	VMA4	40.77	8	110	4.72	1	4	12.45	3	15	21.46	3	5
6323824	HOT1	0	0	0	0	0	0	0	0	0	6.40	2	5
6325066	NAB3	0	0	0	0	0	0	1.87	1	1	7.48	4	5
398364517	RPS24A	13.33	2	62	37.78	5	114	11.11	1	38	15.56	2	4
6321362	RPL7A	21.72	4	67	30.74	7	129	27.05	6	96	9.84	2	4
6325058	RPL7B	18.03	3	63	30.74	7	98	23.36	5	92	9.84	2	4
37362649	GUS1	15.96	11	104	7.20	5	40	8.47	5	20	2.82	2	4
398365909	RPL24B	7.74	1	1	16.77	2	19	0	0	0	16.13	3	4
6324707	EFT1	26.01	20	570	13.90	10	71	17.93	11	84	8.19	4	4
6323376	RPL26A	35.43	7	49	30.71	5	17	18.11	3	48	14.17	2	4
116006491	RPL26B	35.43	7	53	30.71	5	17	18.11	3	48	14.17	2	4

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6325006	RPL36B	33.00	4	39	42.00	5	298	33.00	5	53	7.00	1	4
6323848	RPL36A	33.00	4	39	33.00	4	282	23.00	3	51	7.00	1	4
398366579	RPL12B	33.33	4	538	28.48	3	144	37.58	4	164	16.97	2	4
6321540	NOP7	7.93	3	3	0	0	0	0	0	0	2.98	1	4
398365599	PDC6	3.37	1	19	3.37	1	28	4.97	2	66	3.37	1	4
6323163	PDC5	7.28	3	35	9.59	3	35	7.82	2	67	3.37	1	4
6323583	CMP2	2.65	1	1	0	0	0	0	0	0	8.61	3	4
398365991	CLC1	4.29	1	2	6.87	1	3	11.59	2	16	12.45	2	4
37362619	RXT2	0	0	0	0	0	0	0	0	0	3.72	1	4
10383793	RPS14A	37.96	5	133	19.71	4	10	29.93	3	4	28.47	2	3
6322271	RPS22A	23.85	3	23	25.38	3	24	12.31	1	13	19.23	2	3
6321407	RPL24A	7.74	2	2	22.58	3	18	0	0	0	9.68	2	3
6323357	RPL38	32.05	2	95	28.21	2	16	19.23	1	11	19.23	1	3
398366183	HTB1	13.74	2	11	0	0	0	0	0	0	11.45	1	3
398364287	GCN1	8.57	20	78	6.29	12	47	6.96	13	38	0.60	1	3
6320016	VMA1	20.63	21	108	5.32	4	8	11.95	9	75	1.87	1	3
14318481	TUB2	13.79	6	99	8.75	3	21	6.56	2	7	15.10	3	3
398364585	KAR2	14.66	8	99	15.69	8	154	8.21	4	41	1.76	1	3
6323196	RPS31	10.53	1	2	0	0	0	0	0	0	22.37	1	3
6321013	PAB1	19.58	10	168	9.01	4	12	5.55	2	9	4.51	2	3
6324993	FUM1	9.43	4	29	0	0	0	3.07	1	2	3.28	1	3
398366075	RPS29A	14.29	1	1	14.29	1	1	0	0	0	17.86	1	3
6322755	YJU2	4.32	1	1	0	0	0	0	0	0	12.23	2	3
398366059	PLP2	3.50	1	1	0	0	0	0	0	0	3.50	1	3
6320718	SMT3	10.89	1	51	0	0	0	0	0	0	32.67	2	3
398365031	YBR139W	0	0	0	0	0	0	0	0	0	6.10	2	3
6321850	FYV4	0	0	0	0	0	0	0	0	0	14.62	1	3
398366083	TDH3	52.11	14	985	58.73	13	441	34.64	10	475	7.53	2	2
6319279	CDC19	47.00	22	1858	44.00	19	634	31.00	14	344	3.60	1	2
6322468	TDH2	44.58	12	956	35.84	9	361	27.71	8	453	7.53	2	2
6321249	RPS26A	20.17	2	121	20.17	2	153	12.61	1	3	20.17	2	2
6321772	RPS20	38.02	5	361	31.40	3	14	9.92	1	32	9.92	1	2
14318474	SEC53	25.20	6	25	32.68	5	15	11.81	2	21	5.51	1	2
6320589	YRA1	22.12	5	187	9.73	2	13	5.75	1	2	5.75	1	2
6320805	ECM10	4.50	4	64	0	0	0	5.12	2	6	1.40	1	2
6323776	ECM16	1.66	2	2	0	0	0	0	0	0	1.10	1	2
6320596	RVS167	9.75	3	5	0	0	0	14.94	4	20	5.19	2	2

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398366359	SLH1	0.51	1	1	0	0	0	0	0	0	2.80	2	2
6324321	SIS1	25.00	5	10	6.53	2	4	12.22	3	5	5.97	1	2
6319447	NCL1	17.54	10	77	1.90	1	1	0	0	0	5.56	2	2
6323112	EMP70	4.35	2	2	2.40	1	12	2.55	1	11	3.15	1	2
6324595	SFM1	3.29	1	1	0	0	0	0	0	0	12.68	1	2
330443411	UBP13	1.07	1	1	0	0	0	0	0	0	3.21	1	2
6323868	EFR3	3.58	2	4	0	0	0	0	0	0	2.05	1	2
330443676	CDC34	8.27	4	5	0	0	0	0	0	0	2.69	1	2
6320615	TRS120	0	0	0	0	0	0	0	0	0	1.16	1	2
6323487	FMP27	0	0	0	0	0	0	0	0	0	1.14	2	2
6320231	REG1	0	0	0	0	0	0	1.28	1	1	1.58	1	2
6324585	AUS1	0	0	0	1.08	1	5	3.59	3	4	1.15	1	2
6325048	RKM1	3.43	2	2	0	0	0	0	0	0	5.32	2	2
6323732	SEG1	5.52	4	6	0	0	0	0	0	0	1.56	1	2
398364779	YGL041W-	0	0	0	0	0	0	0	0	0	13.64	1	2
398365827	CAF130	0	0	0	0	0	0	0	0	0	1.07	1	2
6325063	RSA1	0	0	0	0	0	0	0	0	0	6.30	1	2
398365559	MSH6	0	0	0	0	0	0	0	0	0	1.85	2	2
398365727	YNR034W-	0	0	0	0	0	0	0	0	0	11.22	1	2
6323683	YET2	0	0	0	0	0	0	0	0	0	11.88	1	2
330443366	GPB2	0	0	0	0	0	0	0	0	0	1.14	1	2
6325012	HUT1	0	0	0	0	0	0	0	0	0	6.19	1	2
99030927	KKQ8	0	0	0	0	0	0	0	0	0	1.10	1	2
398364567	MAD1	0	0	0	0	0	0	0	0	0	2.14	2	2
6320763	RMD6	0	0	0	0	0	0	0	0	0	6.49	1	2
398365009	MEC1	24.58	54	2197	8.91	17	163	17.53	36	383	0.68	1	1
398364407	RPL9A	33.51	4	41	24.08	4	56	25.65	4	105	5.24	1	1
6324298	HHF2	21.36	2	7	33.01	3	131	40.78	4	110	11.65	1	1
6324262	RPL9B	33.51	4	30	14.14	3	43	15.71	3	100	5.24	1	1
398364523	TDH1	38.25	11	237	39.46	10	369	23.49	7	348	4.22	1	1
398364951	VMA2	36.17	14	511	22.63	8	39	24.18	9	53	2.71	1	1
6321333	ARC1	14.63	5	47	7.18	2	2	7.18	2	13	4.26	1	1
6321408	RPL30	44.76	3	108	39.05	4	89	27.62	3	36	11.43	1	1
6324445	RPL25	25.35	4	68	34.51	4	182	42.25	5	31	9.15	1	1
6325114	RPL33A	24.30	2	54	29.91	3	83	14.95	1	58	10.28	1	1
398365881	RPL33B	9.35	1	2	29.91	3	63	14.95	1	51	10.28	1	1
398364339	THS1	17.44	11	51	3.68	2	7	5.99	3	6	1.23	1	1

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
330443609	URA2	16.26	30	229	6.01	10	24	4.29	7	21	0.59	1	1
6323209	SAM1	13.61	5	108	12.04	3	22	10.99	3	24	2.36	1	1
398364415	HXK1	7.01	3	20	6.80	3	38	12.99	4	11	3.71	1	1
6321232	CHC1	8.65	13	75	7.20	10	21	6.17	8	51	0.73	1	1
6322352	UTP10	2.94	4	5	0	0	0	0.90	1	1	0.90	1	1
398366655	SAM2	11.20	4	36	9.11	2	17	8.07	2	14	2.34	1	1
6322115	RPN2	15.56	13	33	6.67	5	11	6.03	5	15	1.06	1	1
6320073	RPP1B	15.09	1	178	15.09	1	12	15.09	1	2	15.09	1	1
6320122	RPP1A	25.47	2	4	0	0	0	0	0	0	20.75	1	1
6323693	ERB1	3.97	3	9	0	0	0	0	0	0	2.60	1	1
6320784	VMA8	3.91	1	8	0	0	0	4.30	1	3	4.30	1	1
6323087	SHM2	19.83	9	69	6.18	2	14	8.74	3	32	1.92	1	1
6320119	SUB2	4.26	2	3	7.62	3	38	3.36	1	1	7.17	1	1
6320573	ESF1	7.64	4	4	0	0	0	0	0	0	2.39	1	1
6323179	STM1	13.92	2	5	0	0	0	7.33	1	11	8.79	1	1
398366241	SEN1	0.49	1	1	0	0	0	0	0	0	0.58	1	1
6323763	ASC1	20.06	6	254	3.76	1	2	4.70	1	1	4.70	1	1
6324772	BFR1	17.87	7	28	2.55	1	3	4.04	1	2	3.19	1	1
6320175	ARP2	6.91	2	9	0	0	0	11.25	3	6	5.12	1	1
398366669	EMI2	0	0	0	0	0	0	0	0	0	2.80	1	1
6323389	RSC2	0	0	0	0	0	0	0	0	0	2.02	1	1
6324663	VPS21	5.24	1	1	12.38	2	4	7.14	1	1	8.57	1	1
6319543	TAT1	2.10	1	1	0	0	0	6.46	3	7	4.36	1	1
6325417	KRE6	7.92	4	59	2.08	1	1	6.39	3	11	2.64	1	1
6324021	KRI1	6.09	3	8	0	0	0	1.69	1	1	2.20	1	1
6323454	YLR422W	0	0	0	0	0	0	0.47	1	1	0.67	1	1
6319756	PAF1	24.04	7	65	0	0	0	0	0	0	2.25	1	1
6323725	SEC14	8.55	4	10	3.29	1	1	0	0	0	7.57	1	1
6323368	SGD1	3.89	3	4	0	0	0	0	0	0	2.45	1	1
6322598	IML1	4.99	6	11	0	0	0	0	0	0	1.14	1	1
398366503	YDR341C	7.74	4	13	0	0	0	0	0	0	1.65	1	1
6320423	RAD9	0	0	0	0	0	0	0	0	0	1.45	1	1
6325147	GDE1	10.87	10	38	0	0	0	1.23	1	1	1.55	1	1
6319585	CMD1	38.10	4	19	8.84	1	2	16.33	2	6	8.84	1	1
6322946	PTR2	0	0	0	0	0	0	8.65	3	42	3.66	1	1
398366323	GND2	4.47	2	79	2.03	1	6	0	0	0	5.08	1	1
6324231	RAS2	18.32	4	4	0	0	0	0	0	0	4.66	1	1

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6320637	TIF35	15.33	3	6	0	0	0	0	0	0	2.55	1	1
398364223	NPR2	7.80	4	13	0	0	0	0	0	0	2.44	1	1
6325390	SPN1	13.66	5	14	0	0	0	0	0	0	3.41	1	1
6324578	DNL4	0	0	0	0	0	0	0	0	0	1.69	1	1
6319291	FRT2	0	0	0	0	0	0	0	0	0	2.84	1	1
6322242	HXT9	0	0	0	0	0	0	0	0	0	4.06	1	1
6322272	RPL39	0	0	0	0	0	0	0	0	0	15.69	1	1
14318518	VTC2	1.81	1	1	1.45	1	1	6.52	3	3	1.57	1	1
330443752	THI22	0	0	0	1.75	1	12	3.15	2	5	2.97	1	1
6325175	MOT1	0.54	1	1	1.12	1	1	0	0	0	0.80	1	1
6322796	DEF1	6.64	3	7	1.08	1	1	0	0	0	4.20	1	1
398365059	PTK2	1.47	1	1	0	0	0	0	0	0	2.81	1	1
6320345	DOP1	1.88	2	2	0	0	0	0	0	0	0.71	1	1
6323863	HFA1	0.52	1	2	0	0	0	0	0	0	1.08	1	1
6322948	MLP1	1.12	2	2	0	0	0	0	0	0	0.64	1	1
6322999	FRA1	2.94	2	2	0	0	0	0	0	0	2.27	1	1
6324491	IRA2	0.23	1	2	0	0	0	0	0	0	0.23	1	1
6324215	RPC19	37.32	4	11	0	0	0	0	0	0	10.56	1	1
6320419	UPC2	0	0	0	0	0	0	0	0	0	3.40	1	1
398364627	RDH54	0	0	0	0	0	0	0	0	0	2.82	1	1
6321752	VMR1	0	0	0	0	0	0	0	0	0	0.75	1	1
6321793	TIM10	0	0	0	0	0	0	0	0	0	7.53	1	1
6322032	BNR1	0	0	0	0	0	0	0	0	0	1.09	1	1
6325074	RTT10	0	0	0	0	0	0	0	0	0	0.69	1	1
6325235	RAD1	0	0	0	0	0	0	0	0	0	1.82	1	1
6323869	CEF1	3.39	1	1	0	0	0	2.20	1	1	2.20	1	1
6323917	TPS3	4.46	4	6	0	0	0	1.61	1	1	2.47	1	1
6321552	SPT6	8.34	11	17	0	0	0	1.24	1	1	0.69	1	1
6320853	BIM1	0	0	0	0	0	0	4.65	1	1	4.07	1	1
398364243	YEL043W	0	0	0	0	0	0	1.78	1	1	1.15	1	1
6320027	DLD1	0	0	0	0	0	0	2.21	1	1	3.58	1	1
6323356	PEX30	0	0	0	0	0	0	3.44	1	1	3.25	1	1
6320799	ANP1	0	0	0	0	0	0	3.60	1	2	4.40	1	1
6324393	YNR065C	3.14	3	57	0	0	0	1.25	1	7	1.70	1	1
6323266	THI7	0	0	0	0	0	0	5.02	2	8	6.02	1	1
6323445	YLR413W	2.96	2	2	0	0	0	5.78	3	14	2.52	1	1
398364667	THO2	0.63	1	1	0.75	1	1	0	0	0	1.19	1	1

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
37362637	JIP4	1.48	1	1	1.03	1	1	0	0	0	1.14	1	1
398364707	AAC3	3.58	1	1	6.19	2	2	0	0	0	8.79	1	1
6322570	YMR1	1.02	1	1	1.02	1	2	0	0	0	1.74	1	1
6324879	ISW2	1.34	1	1	0	0	0	0	0	0	2.23	1	1
10383772	LEU2	6.32	1	1	0	0	0	0	0	0	6.32	1	1
6322079	HOS4	0.92	1	1	0	0	0	0	0	0	2.49	1	1
6321210	VID30	1.36	1	1	0	0	0	0	0	0	2.19	1	1
398365701	YNR029C	2.10	1	1	0	0	0	0	0	0	4.20	1	1
6320692	VPS52	1.40	1	1	0	0	0	0	0	0	2.03	1	1
398365525	SMD1	6.16	1	1	0	0	0	0	0	0	23.29	1	1
6322086	SLM1	1.60	1	1	0	0	0	0	0	0	1.75	1	1
398364897	CBP6	6.79	1	1	0	0	0	0	0	0	9.26	1	1
6323726	NAM7	0.93	1	1	0	0	0	0	0	0	1.75	1	1
6324982	SAM4	3.08	1	1	0	0	0	0	0	0	5.85	1	1
6320324	VBA4	1.69	1	1	0	0	0	0	0	0	3.78	1	1
6324612	HIR2	1.26	1	1	0	0	0	0	0	0	2.29	1	1
6324981	SAM3	2.39	1	1	0	0	0	0	0	0	3.24	1	1
6325379	AXL1	0.99	1	1	0	0	0	0	0	0	0.91	1	1
6321020	BRR2	0.46	1	1	0	0	0	0	0	0	0.83	1	1
6324694	GCY1	3.21	1	1	0	0	0	0	0	0	6.09	1	1
6322312	DAS1	1.06	1	1	0	0	0	0	0	0	2.71	1	1
398364625	YKL033W-	3.81	1	2	0	0	0	0	0	0	6.36	1	1
6320034	NRP1	3.06	2	2	0	0	0	0	0	0	2.50	1	1
6325374	YPR117W	0.40	1	2	0	0	0	0	0	0	0.68	1	1
6324546	MDM38	7.16	2	2	0	0	0	0	0	0	3.66	1	1
6323289	LCB5	2.33	1	3	0	0	0	0	0	0	2.33	1	1
6321205	TAN1	13.84	4	4	0	0	0	0	0	0	3.11	1	1
6319351	PHO11	1.71	1	4	0	0	0	0	0	0	4.71	1	1
6319523	REB1	6.67	4	5	0	0	0	0	0	0	3.70	1	1
6320921	YER077C	0	0	0	0	0	0	0	0	0	3.05	1	1
6320826	GLC3	0	0	0	0	0	0	0	0	0	2.41	1	1
398365141	INP53	0	0	0	0	0	0	0	0	0	1.99	1	1
398366583	SIP1	0	0	0	0	0	0	0	0	0	2.70	1	1
398364669	MAD3	0	0	0	0	0	0	0	0	0	3.88	1	1
6322700	DBR1	0	0	0	0	0	0	0	0	0	3.95	1	1
6323317	YLR287C	0	0	0	0	0	0	0	0	0	6.20	1	1
398365819	SAN1	0	0	0	0	0	0	0	0	0	3.44	1	1

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6319273	PTA1	0	0	0	0	0	0	0	0	0	1.78	1	1
6323592	RSE1	0	0	0	0	0	0	0	0	0	0.88	1	1
398364677	GLE2	0	0	0	0	0	0	0	0	0	3.56	1	1
6324505	RTG1	0	0	0	0	0	0	0	0	0	10.73	1	1
6325281	YME1	0	0	0	0	0	0	0	0	0	5.09	1	1
6320159	FAD1	0	0	0	0	0	0	0	0	0	5.88	1	1
6319711	ARC40	0	0	0	0	0	0	0	0	0	3.65	1	1
6323933	FCP1	0	0	0	0	0	0	0	0	0	3.96	1	1
6323641	YML002W	0	0	0	0	0	0	0	0	0	1.90	1	1
6323762	MGR3	0	0	0	0	0	0	0	0	0	3.39	1	1
6324948	NDD1	0	0	0	0	0	0	0	0	0	3.61	1	1
398365185	AZF1	0	0	0	0	0	0	0	0	0	3.94	1	1
6325040	YPL216W	0	0	0	0	0	0	0	0	0	1.27	1	1
6322932	TRZ1	0	0	0	0	0	0	0	0	0	3.82	1	1
6322965	AYT1	0	0	0	0	0	0	0	0	0	4.85	1	1
398366037	NUP42	0	0	0	0	0	0	0	0	0	6.51	1	1
398364347	GPA2	0	0	0	0	0	0	0	0	0	3.34	1	1
6322492	CPR7	0	0	0	0	0	0	0	0	0	4.07	1	1
37362610	FUN19	0	0	0	0	0	0	0	0	0	6.30	1	1
398364541	INP51	0	0	0	0	0	0	0	0	0	2.96	1	1
6323580	OGG1	0	0	0	0	0	0	0	0	0	4.26	1	1
398366675	AGE1	0	0	0	0	0	0	0	0	0	3.32	1	1
6320060	BPL1	0	0	0	0	0	0	0	0	0	2.61	1	1
330443457	BMT2	0	0	0	0	0	0	0	0	0	6.23	1	1
6323385	BUD8	0	0	0	0	0	0	0	0	0	4.64	1	1
398364867	CTR2	0	0	0	0	0	0	0	0	0	6.88	1	1
398364539	RTT101	0	0	0	0	0	0	0	0	0	1.66	1	1
398364351	CUP2	0	0	0	0	0	0	0	0	0	10.22	1	1
6323896	CUS1	0	0	0	0	0	0	0	0	0	2.52	1	1
6323760	FOL3	0	0	0	0	0	0	0	0	0	4.45	1	1
398365821	EFM2	0	0	0	0	0	0	0	0	0	6.44	1	1
6322629	FRE2	0	0	0	0	0	0	0	0	0	4.36	1	1
6322324	GLG2	0	0	0	0	0	0	0	0	0	2.37	1	1
6321676	YGR237C	0	0	0	0	0	0	0	0	0	1.27	1	1
6322432	YJL028W	0	0	0	0	0	0	0	0	0	8.11	1	1
6322498	YJR039W	0	0	0	0	0	0	0	0	0	1.43	1	1
6323075	YLR046C	0	0	0	0	0	0	0	0	0	4.44	1	1

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398365335	YNL033W	0	0	0	0	0	0	0	0	0	4.93	1	1
33438893	YPL119C-	0	0	0	0	0	0	0	0	0	25.29	1	1
6322579	JHD2	0	0	0	0	0	0	0	0	0	3.43	1	1
6324141	KAR1	0	0	0	0	0	0	0	0	0	2.31	1	1
10383756	LRE1	0	0	0	0	0	0	0	0	0	3.77	1	1
398364917	NHP10	0	0	0	0	0	0	0	0	0	4.93	1	1
6323035	NSE1	0	0	0	0	0	0	0	0	0	6.85	1	1
50593425	OMA1	0	0	0	0	0	0	0	0	0	3.77	1	1
6321910	ORC6	0	0	0	0	0	0	0	0	0	5.75	1	1
6322494	PET191	0	0	0	0	0	0	0	0	0	23.15	1	1
398365057	SPO14	0	0	0	0	0	0	0	0	0	0.83	1	1
398364971	TRP1	0	0	0	0	0	0	0	0	0	11.16	1	1
398365003	PSF1	0	0	0	0	0	0	0	0	0	7.69	1	1
6321697	RAD2	0	0	0	0	0	0	0	0	0	0.78	1	1
6319525	REG2	0	0	0	0	0	0	0	0	0	3.85	1	1
6320271	RTR2	0	0	0	0	0	0	0	0	0	5.61	1	1
6324162	SKO1	0	0	0	0	0	0	0	0	0	3.40	1	1
398366633	SNM1	0	0	0	0	0	0	0	0	0	5.05	1	1
6322778	STB6	0	0	0	0	0	0	0	0	0	2.61	1	1
6325081	TRE1	0	0	0	0	0	0	0	0	0	2.68	1	1
6324447	TRM13	0	0	0	0	0	0	0	0	0	2.52	1	1
6323236	HRD3	0	0	0	0	0	0	0	0	0	1.80	1	1
6325184	UBP16	0	0	0	0	0	0	0	0	0	3.61	1	1
398364885	UBP5	0	0	0	0	0	0	0	0	0	1.86	1	1
6325071	UIP4	0	0	0	0	0	0	0	0	0	10.20	1	1
398365127	UIP5	0	0	0	0	0	0	0	0	0	4.74	1	1
6319904	YIH1	0	0	0	0	0	0	0	0	0	11.24	1	1
6325266	SUT2	0	0	0	0	0	0	0	0	0	7.46	1	1
6324449	HRP1	0	0	0	0	0	0	0	0	0	4.31	1	1
6323278	YEF3	19.44	17	205	10.63	9	72	12.36	12	100	0	0	0
6322084	PFK26	37.00	27	1046	48.85	32	1153	45.59	28	1064	0	0	0
6323057	RPL15A	25.49	5	82	24.02	4	76	11.27	2	4	0	0	0
6323769	RPL15B	25.49	5	75	20.59	3	58	7.84	1	3	0	0	0
6319282	FUN12	15.77	13	32	0	0	0	0	0	0	0	0	0
6320190	NOP1	19.27	5	17	0	0	0	0	0	0	0	0	0
6323654	CLU1	22.08	21	499	0	0	0	6.11	5	10	0	0	0
6324886	NOP58	17.42	7	21	0	0	0	15.85	5	11	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6323226	NOP56	18.65	7	25	0	0	0	10.52	3	10	0	0	0
37362658	SSZ1	26.39	11	119	17.29	6	51	13.75	5	14	0	0	0
6324452	RPL18A	23.66	4	161	22.04	4	311	12.37	2	89	0	0	0
398366653	LCD1	31.99	21	565	6.43	5	27	16.47	9	86	0	0	0
6322313	RPA34	44.64	7	48	32.62	6	71	26.61	4	150	0	0	0
398364437	RPL1B	6.45	2	9	28.11	5	343	24.42	5	136	0	0	0
6321968	ENO2	37.76	15	1712	50.57	14	415	46.00	16	633	0	0	0
398365729	DED1	24.83	12	51	15.40	6	40	5.13	2	2	0	0	0
6323566	FPR3	2.68	1	73	2.68	1	1	4.14	1	2	0	0	0
398366315	ENO1	38.44	14	1548	47.14	13	427	35.70	12	518	0	0	0
10383781	PGK1	40.14	16	691	39.90	13	164	37.50	14	273	0	0	0
6321874	KSP1	11.08	9	148	0	0	0	2.62	2	3	0	0	0
6321531	VAS1	23.82	23	229	5.71	5	7	5.25	4	5	0	0	0
6322769	TEF4	14.08	5	173	9.71	3	24	22.57	7	65	0	0	0
398366187	HTA1	6.82	1	15	6.82	1	4	0	0	0	0	0	0
330443451	GRS1	25.49	17	98	2.40	1	1	6.60	3	10	0	0	0
6321203	ADE5,7	15.71	9	23	4.99	3	5	9.10	5	14	0	0	0
6324486	ADH1	31.90	11	1603	43.10	10	217	46.55	13	501	0	0	0
6323371	RPP0	15.38	4	115	24.04	7	186	28.85	7	157	0	0	0
6323886	RPS10B	7.62	1	2	0	0	0	0	0	0	0	0	0
6320009	ARF1	18.78	3	11	13.81	2	25	29.28	4	54	0	0	0
6321429	LEU1	26.57	18	905	10.01	7	88	7.32	4	22	0	0	0
6323204	CBF5	3.93	1	4	0	0	0	0	0	0	0	0	0
6320064	ARF2	18.78	3	11	13.81	2	10	23.76	3	24	0	0	0
6321265	XRN1	10.67	12	49	0	0	0	3.01	3	4	0	0	0
6323904	RKR1	17.99	26	637	5.06	6	12	3.20	5	18	0	0	0
6322790	FBA1	12.81	3	602	18.11	5	59	24.51	5	160	0	0	0
6324562	HTZ1	7.46	1	1	0	0	0	7.46	1	3	0	0	0
6321231	SPT16	5.60	5	10	0	0	0	0	0	0	0	0	0
398365455	TEF3	3.07	3	71	2.59	2	2	2.01	2	19	0	0	0
398366333	FPR4	0	0	0	0	0	0	4.34	1	2	0	0	0
398364607	DBP3	20.46	9	74	0	0	0	0	0	0	0	0	0
6324534	RPP2A	61.32	4	75	37.74	2	6	50.94	3	10	0	0	0
6321430	PMA1	16.12	14	78	19.93	13	99	14.16	11	138	0	0	0
398364809	DBP2	15.57	8	18	9.16	4	8	0	0	0	0	0	0
6319668	RPL21A	21.88	4	21	21.88	4	106	16.25	3	16	0	0	0
6323981	FET4	11.23	5	18	1.99	1	24	8.51	4	52	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6321807	DED81	13.18	6	32	5.96	2	5	10.11	4	20	0	0	0
398364767	SUI2	25.33	6	20	3.62	1	1	9.21	2	5	0	0	0
6323011	DPS1	20.65	10	56	1.62	1	2	3.23	1	3	0	0	0
6323885	RRP5	4.80	8	10	0	0	0	0	0	0	0	0	0
398366547	RPP2B	50.00	4	138	40.00	2	2	61.82	5	33	0	0	0
6323288	HSP60	35.49	16	256	33.39	14	215	31.82	13	167	0	0	0
6321599	NSR1	11.11	3	16	0	0	0	0	0	0	0	0	0
6320863	GCD11	22.39	10	73	3.23	1	3	9.11	3	9	0	0	0
398366591	NPL3	14.73	4	20	15.70	4	215	10.87	3	13	0	0	0
6319483	IPP1	19.86	5	89	17.07	3	30	30.66	5	28	0	0	0
6325221	PMA2	12.35	11	75	14.99	11	86	10.77	9	133	0	0	0
6322745	GFA1	16.32	7	75	9.90	5	26	18.41	8	30	0	0	0
398364525	LSG1	14.22	8	10	1.72	1	2	0	0	0	0	0	0
6325038	SAR1	25.79	4	33	20.53	3	53	6.84	1	1	0	0	0
6323840	HSC82	34.04	23	914	17.02	10	109	21.13	12	62	0	0	0
6321881	GAR1	0	0	0	0	0	0	6.34	1	1	0	0	0
6320164	NAT1	5.62	4	7	0	0	0	0	0	0	0	0	0
398366403	RPS12	22.38	3	21	7.69	2	4	26.57	4	65	0	0	0
6322830	CDC16	21.19	17	255	15.48	11	86	17.74	11	83	0	0	0
330443397	ATP1	10.46	5	43	19.45	9	89	21.10	8	47	0	0	0
6324709	IDH2	23.85	7	188	15.99	4	19	15.99	4	19	0	0	0
6325178	RPL21B	21.88	4	21	21.88	4	106	16.25	3	16	0	0	0
6321903	UBA4	28.86	13	382	23.18	11	343	15.00	6	159	0	0	0
6325138	DBP1	6.32	4	36	1.78	1	2	0	0	0	0	0	0
6325248	TAE2	1.16	1	1	0	0	0	0	0	0	0	0	0
330443505	DBP10	4.52	4	6	0	0	0	0	0	0	0	0	0
6320148	PSA1	10.25	3	4	21.61	5	36	16.07	5	57	0	0	0
6325016	HSP82	31.17	21	887	19.32	11	110	21.44	12	63	0	0	0
6320874	ARB1	20.82	9	113	8.52	4	9	9.67	4	8	0	0	0
6319673	PGI1	12.09	6	72	0	0	0	2.53	1	1	0	0	0
398365251	GSC2	3.43	6	9	6.60	9	130	16.89	21	465	0	0	0
6325018	SUI3	35.44	8	73	0	0	0	4.56	1	1	0	0	0
6323374	FKS1	9.49	14	41	8.80	12	156	19.03	26	633	0	0	0
14318479	ACT1	20.53	7	65	14.93	5	49	23.20	6	55	0	0	0
6323554	TUB1	10.96	5	90	5.37	2	2	8.05	3	3	0	0	0
6321724	ZUO1	20.09	7	55	21.94	5	21	3.70	1	4	0	0	0
6320842	NUG1	10.58	4	4	2.50	1	3	0	0	0	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6321027	DMC1	0	0	0	0	0	0	0	0	0	0	0	0
398364483	HOR2	17.20	4	100	6.00	1	22	6.00	1	8	0	0	0
398365429	PDR5	10.13	12	19	0	0	0	17.94	19	375	0	0	0
6325245	RRP12	5.54	6	8	0	0	0	0	0	0	0	0	0
398364937	TOP2	0.77	1	1	0	0	0	0	0	0	0	0	0
6324660	TCB1	1.01	1	1	1.26	1	1	1.60	1	2	0	0	0
6320056	COP1	11.32	11	20	0.67	1	1	6.74	5	12	0	0	0
398365597	BMH2	24.54	7	105	8.42	1	11	15.38	3	12	0	0	0
6325030	NEW1	8.53	9	24	2.84	3	3	0.84	1	2	0	0	0
6321809	RPS27B	9.76	1	5	0	0	0	0	0	0	0	0	0
6321012	CHD1	5.72	7	8	0	0	0	0	0	0	0	0	0
6320499	SSD1	4.88	5	8	0	0	0	0	0	0	0	0	0
6323072	TRX1	45.63	4	105	9.71	1	1	24.27	2	8	0	0	0
6325238	VTC3	2.75	2	2	1.56	1	1	10.18	6	32	0	0	0
398365319	MAK21	10.15	7	16	0	0	0	0	0	0	0	0	0
398364327	LSB3	22.88	8	72	0	0	0	7.84	2	2	0	0	0
6319513	ATP3	18.97	6	44	8.68	2	6	17.04	4	6	0	0	0
6324140	SRP1	5.35	3	27	2.40	1	1	2.40	1	14	0	0	0
6324297	HHT2	0	0	0	5.15	1	1	0	0	0	0	0	0
6323306	DBP9	10.61	6	7	0	0	0	0	0	0	0	0	0
6324039	RFC3	17.35	5	7	0	0	0	3.24	1	1	0	0	0
6321193	RTF1	14.87	8	127	1.97	1	1	2.33	1	4	0	0	0
398366149	TRX2	58.65	5	183	9.62	1	1	12.50	1	4	0	0	0
6322452	CCT8	19.72	10	32	13.03	6	15	6.34	3	46	0	0	0
6321679	PFK1	24.11	19	339	10.03	7	29	10.64	8	46	0	0	0
6324219	NOP15	4.09	1	1	0	0	0	0	0	0	0	0	0
330443717	CTR9	16.06	15	62	0	0	0	0	0	0	0	0	0
6323755	ILV2	9.75	5	49	6.26	3	10	10.77	4	7	0	0	0
6320276	PAA1	9.95	2	3	6.81	1	9	0	0	0	0	0	0
6681849	RPL42A	0	0	0	0	0	0	0	0	0	0	0	0
14318517	SEC4	15.81	3	6	5.12	1	2	8.37	1	10	0	0	0
6320400	MSS116	3.92	2	3	0	0	0	0	0	0	0	0	0
6321389	TIF4632	4.92	3	6	0	0	0	0	0	0	0	0	0
6324081	RPA49	32.53	11	52	0	0	0	0	0	0	0	0	0
398366347	MES1	17.44	12	33	0	0	0	3.46	2	7	0	0	0
6320418	TCP1	19.50	10	146	10.38	5	20	10.02	4	24	0	0	0
398365595	PIL1	34.51	8	113	27.73	6	33	34.51	7	42	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398364417	SAH1	13.14	5	94	2.67	1	1	3.34	1	1	0	0	0
6320396	RVB1	16.41	6	26	2.59	1	1	5.40	2	2	0	0	0
6320077	CDC48	13.05	8	44	8.02	5	32	9.10	5	12	0	0	0
6325025	FAS2	13.94	21	150	14.84	20	126	18.02	23	146	0	0	0
6324517	THI20	14.52	6	32	30.67	15	548	31.22	14	411	0	0	0
6323030	NOC3	1.96	1	1	0	0	0	0	0	0	0	0	0
6320143	TSR1	4.44	3	5	0	0	0	0	0	0	0	0	0
6321318	PRP43	7.17	5	10	0	0	0	0	0	0	0	0	0
6319618	MAK5	9.44	7	11	0	0	0	0	0	0	0	0	0
398364755	URB1	0	0	0	0	0	0	0	0	0	0	0	0
6320780	POL5	4.79	4	4	0	0	0	1.47	1	1	0	0	0
6325021	RVB2	18.47	7	34	5.10	2	2	6.58	2	3	0	0	0
6322350	CCT7	15.64	6	48	16.36	7	66	8.91	4	21	0	0	0
14270688	UTP20	1.20	3	3	0	0	0	0	0	0	0	0	0
6324197	KRE33	5.40	4	5	0	0	0	0	0	0	0	0	0
6321816	RPN1	13.39	11	53	2.52	2	2	5.24	4	9	0	0	0
6322901	NAP1	9.83	3	5	4.08	1	1	8.15	2	10	0	0	0
6320030	GLT1	6.90	12	135	5.59	10	19	0.65	1	10	0	0	0
6323470	CAR2	6.84	2	9	14.39	3	12	11.79	3	12	0	0	0
6320956	KAP123	16.80	13	66	10.15	8	42	8.09	7	34	0	0	0
398365941	SEC7	4.13	7	9	1.99	2	2	0.70	1	1	0	0	0
6323324	GSP1	14.61	3	27	9.59	2	6	5.02	1	1	0	0	0
6322697	GPM1	36.44	7	433	29.15	6	24	27.94	5	123	0	0	0
6325013	SRP68	2.17	1	11	1.34	1	2	0	0	0	0	0	0
6322103	AVT7	0	0	0	1.63	1	14	0	0	0	0	0	0
6322049	CCT2	32.26	12	51	20.30	8	38	14.99	5	28	0	0	0
330443694	SLA2	19.42	14	86	3.00	2	2	0	0	0	0	0	0
6321804	YSC84	12.18	5	20	0	0	0	0	0	0	0	0	0
398366021	CCT6	8.24	4	52	11.17	5	15	3.66	1	1	0	0	0
6324042	SEC21	8.66	5	6	3.32	2	5	5.45	4	14	0	0	0
398364741	SCS2	11.48	2	5	11.89	2	9	27.05	4	16	0	0	0
6325253	LSP1	29.91	8	86	13.20	3	17	10.26	2	20	0	0	0
6321390	RPT6	15.06	5	17	10.37	3	28	14.07	4	27	0	0	0
398364807	PNC1	13.43	2	129	13.43	2	7	9.26	1	1	0	0	0
6325164	NOG1	17.62	8	12	2.78	1	12	2.01	1	1	0	0	0
6319814	GRX1	20.00	2	6	25.45	2	14	11.82	1	1	0	0	0
6322823	MAE1	5.68	3	4	1.20	1	1	2.39	1	2	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6320255	TPI1	15.32	3	139	0	0	0	14.52	2	2	0	0	0
6320983	GDI1	3.10	1	15	4.21	1	1	3.10	1	6	0	0	0
398365205	RPT5	24.65	8	29	12.21	4	12	13.13	4	10	0	0	0
6324992	KEL3	6.30	3	4	0	0	0	0	0	0	0	0	0
6323028	DNM1	1.59	1	1	0	0	0	1.85	1	1	0	0	0
398365977	RPT4	10.07	4	15	10.98	4	26	2.52	1	4	0	0	0
6320058	CCT4	11.74	6	70	9.66	4	48	10.98	4	48	0	0	0
6320379	HMO1	16.67	3	29	4.88	1	2	0	0	0	0	0	0
6322508	CYC1	7.34	1	1	0	0	0	0	0	0	0	0	0
6325137	VPS30	0	0	0	0	0	0	0	0	0	0	0	0
6321301	SEC27	8.10	6	10	3.04	2	5	2.70	1	1	0	0	0
6321673	YHB1	6.77	2	15	0	0	0	3.51	1	1	0	0	0
27469359	SYN8	13.33	3	6	0	0	0	5.88	1	6	0	0	0
6324433	CDC33	25.82	6	94	0	0	0	12.68	2	12	0	0	0
6323571	POB3	3.62	2	2	0	0	0	0	0	0	0	0	0
398365157	NOP2	19.09	9	65	0	0	0	0	0	0	0	0	0
6321228	YPT32	30.18	5	6	16.22	3	7	5.86	1	1	0	0	0
398366559	RPT3	4.91	2	10	7.01	2	13	8.64	2	3	0	0	0
6322704	RPT1	11.56	5	27	10.71	4	9	5.78	2	5	0	0	0
398366379	PRT1	13.50	8	14	2.62	2	2	6.95	3	6	0	0	0
398365125	PDA1	2.14	1	1	4.76	2	18	2.62	1	6	0	0	0
398364913	GND1	19.43	8	91	5.52	2	11	6.34	2	9	0	0	0
6322637	SAC1	0	0	0	4.82	2	6	7.54	3	11	0	0	0
398365933	CYS4	25.05	12	124	3.35	1	11	16.77	5	13	0	0	0
6323903	FAA4	7.06	4	28	5.33	3	7	3.46	2	34	0	0	0
6319806	PDI1	17.62	7	59	9.39	3	23	19.54	6	69	0	0	0
6323512	PHO84	10.73	6	73	9.88	3	33	17.21	8	305	0	0	0
6325367	RPC40	17.01	4	34	5.37	1	2	0	0	0	0	0	0
6319835	GBP2	5.15	2	3	0	0	0	0	0	0	0	0	0
6319290	MYO4	0.68	1	1	0	0	0	1.22	1	1	0	0	0
14318531	GCN20	11.44	8	43	0	0	0	1.86	1	1	0	0	0
6320869	YPT31	16.14	3	4	16.14	3	7	11.21	2	2	0	0	0
398365617	UTP22	5.25	5	7	0	0	0	2.18	2	2	0	0	0
398366209	RTN1	9.83	2	9	9.83	2	10	9.83	2	3	0	0	0
6325097	CDC60	11.74	10	24	4.50	4	14	3.12	3	5	0	0	0
6323294	RPS28B	31.34	2	80	13.43	1	1	17.91	1	6	0	0	0
6325151	SSE1	23.95	15	74	15.15	8	24	14.29	7	43	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398366337	APL6	10.38	8	12	0	0	0	0	0	0	0	0	0
6320242	KRS1	17.26	9	15	0	0	0	0	0	0	0	0	0
6324705	YOR131C	0	0	0	0	0	0	0	0	0	0	0	0
6319760	SSH1	0	0	0	4.29	1	1	0	0	0	0	0	0
6323335	ACO1	12.85	8	11	3.34	2	3	0	0	0	0	0	0
330443748	GLN1	9.19	3	5	0	0	0	0	0	0	0	0	0
6319464	SLA1	8.68	6	16	0	0	0	0	0	0	0	0	0
14627173	BBC1	12.53	12	27	0	0	0	0	0	0	0	0	0
398364931	URB2	0	0	0	0	0	0	0	0	0	0	0	0
6325423	RHO1	0	0	0	0	0	0	0	0	0	0	0	0
6321837	AAP1	2.69	2	2	0	0	0	1.64	1	2	0	0	0
6319281	RBG1	11.11	3	3	0	0	0	4.34	1	3	0	0	0
6323379	KAP95	4.88	4	9	0	0	0	3.72	2	3	0	0	0
398366047	RNR4	24.64	7	122	5.51	1	2	10.14	2	11	0	0	0
6322446	CCT3	29.59	14	69	4.12	2	28	10.49	4	15	0	0	0
398365063	LAT1	7.26	3	5	13.69	6	59	27.80	7	48	0	0	0
6322427	HCA4	3.64	2	7	0.91	1	1	0	0	0	0	0	0
6322073	RHO3	0	0	0	5.63	1	1	0	0	0	0	0	0
6324630	NOB1	1.96	1	2	0	0	0	0	0	0	0	0	0
398364409	RET2	2.93	2	3	0	0	0	0	0	0	0	0	0
6319796	SPB1	4.52	3	4	0	0	0	0	0	0	0	0	0
6323968	PSE1	2.20	2	4	0	0	0	0	0	0	0	0	0
6322934	RPF2	2.62	1	9	0	0	0	0	0	0	0	0	0
398365233	RPS21A	25.29	2	22	0	0	0	0	0	0	0	0	0
6322325	RPS21B	25.29	2	22	0	0	0	0	0	0	0	0	0
398366473	SSF2	0	0	0	0	0	0	0	0	0	0	0	0
86559466	CTT1	0	0	0	0	0	0	0	0	0	0	0	0
6325449	QCR2	2.72	1	1	8.42	2	4	4.08	1	1	0	0	0
6323537	TSL1	5.46	5	5	3.73	3	7	1.46	1	1	0	0	0
6320914	RNR1	9.12	6	12	1.58	1	1	1.46	1	2	0	0	0
6323182	ACS2	10.25	7	25	2.64	1	1	5.27	2	2	0	0	0
6319432	URA7	22.11	11	81	2.42	1	3	7.25	3	4	0	0	0
6323135	MDN1	4.13	15	28	0.73	3	5	0.86	3	5	0	0	0
398365477	ATP2	16.83	7	27	23.87	9	51	7.44	2	8	0	0	0
6322988	VPS13	12.15	30	70	1.11	3	3	2.54	5	10	0	0	0
6322794	TMA19	25.15	4	120	9.58	1	6	17.96	2	25	0	0	0
6323037	RLP24	5.03	1	1	6.53	1	1	0	0	0	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398364441	NOT3	5.38	4	6	1.67	1	1	0	0	0	0	0	0
398366191	RPN13	21.15	2	6	21.15	2	4	0	0	0	0	0	0
6321453	YGR016W	5.26	1	2	0	0	0	0	0	0	0	0	0
6320930	ILV1	1.39	1	2	0	0	0	0	0	0	0	0	0
6323016	PUF3	1.93	2	3	0	0	0	0	0	0	0	0	0
37362700	SRP72	7.19	3	3	0	0	0	0	0	0	0	0	0
6321326	TAF6	8.91	3	7	0	0	0	0	0	0	0	0	0
6322565	ADO1	11.47	3	10	0	0	0	0	0	0	0	0	0
6319722	ISW1	6.29	6	11	0	0	0	0	0	0	0	0	0
330443605	SET2	9.96	6	12	0	0	0	0	0	0	0	0	0
398365101	ARP3	7.35	3	58	0	0	0	0	0	0	0	0	0
6321780	STE20	25.77	17	85	0	0	0	0	0	0	0	0	0
6321776	PRS3	6.56	2	2	2.81	1	1	5.94	1	1	0	0	0
6323214	RPL37A	0	0	0	11.36	1	1	11.36	1	1	0	0	0
6321757	ECM29	4.66	7	10	1.93	3	5	1.28	2	2	0	0	0
398365069	ADH5	2.28	1	69	2.28	1	6	2.28	1	2	0	0	0
398365695	CPR8	0	0	0	0	0	0	4.22	1	4	0	0	0
6319854	RVS161	16.60	4	21	11.70	2	4	13.21	3	6	0	0	0
6324620	DBP5	14.52	7	26	2.70	1	2	7.47	3	7	0	0	0
6321399	YGL039W	9.20	2	2	0	0	0	13.51	3	12	0	0	0
398365061	SUP45	11.90	5	11	5.49	2	5	9.61	3	14	0	0	0
6324673	KTR1	4.58	2	2	0	0	0	11.45	4	34	0	0	0
6323387	ILV5	23.04	8	32	10.13	3	9	21.77	6	42	0	0	0
6323795	TIF34	17.58	5	10	4.32	1	1	0	0	0	0	0	0
6322667	PRS1	9.84	3	9	2.58	1	6	0	0	0	0	0	0
6320910	RGI1	8.07	1	1	0	0	0	0	0	0	0	0	0
398365575	HOM6	8.91	3	4	0	0	0	0	0	0	0	0	0
6323916	TIF11	26.14	4	5	0	0	0	0	0	0	0	0	0
6321612	RBG2	17.93	5	8	0	0	0	0	0	0	0	0	0
6323225	PWP1	13.54	6	13	0	0	0	0	0	0	0	0	0
6324737	SSP1	21.81	4	31	0	0	0	0	0	0	0	0	0
6323450	CDC73	25.95	8	78	0	0	0	0	0	0	0	0	0
398365461	SRP40	0	0	0	0	0	0	0	0	0	0	0	0
6320708	RPL37B	0	0	0	0	0	0	0	0	0	0	0	0
398364957	MKT1	2.53	1	1	0	0	0	1.81	1	1	0	0	0
6320079	YDL124W	11.54	3	6	8.33	2	2	4.49	1	2	0	0	0
398364371	RSC8	13.29	6	7	0	0	0	5.57	2	2	0	0	0

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6325418	GPH1	1.11	1	1	2.66	2	2	1.44	1	3	0	0	0
6321859	DYS1	2.33	1	3	7.49	2	2	7.49	2	3	0	0	0
6323967	GAS1	7.51	3	3	2.50	1	1	2.50	1	7	0	0	0
6319695	PYC2	6.27	7	13	0.85	1	1	3.56	3	8	0	0	0
6323642	YPT7	0	0	0	12.02	2	5	18.27	2	9	0	0	0
6320552	HXT3	2.12	1	1	2.12	1	2	12.35	4	11	0	0	0
6325196	ALD6	25.60	10	75	6.60	2	3	9.80	3	12	0	0	0
6319315	EFB1	42.72	6	153	8.25	1	1	8.25	1	14	0	0	0
10383811	CDC39	0.90	2	2	0.76	1	1	0	0	0	0	0	0
6321753	SBP1	6.80	2	2	5.78	1	1	0	0	0	0	0	0
6320764	DLD3	9.68	4	12	3.23	1	1	0	0	0	0	0	0
398366413	ERV29	0	0	0	7.74	2	2	0	0	0	0	0	0
6320969	GLO3	2.23	1	1	1.42	1	12	0	0	0	0	0	0
6319602	TPS1	2.02	1	1	8.48	3	28	0	0	0	0	0	0
6579193	NMD2	0.83	1	1	0	0	0	0	0	0	0	0	0
6321990	UTP9	1.22	1	2	0	0	0	0	0	0	0	0	0
398366561	SXM1	2.22	2	2	0	0	0	0	0	0	0	0	0
6320157	NPC2	6.94	1	2	0	0	0	0	0	0	0	0	0
398364687	NOP9	8.71	5	5	0	0	0	0	0	0	0	0	0
22532997	BUD27	2.51	2	5	0	0	0	0	0	0	0	0	0
398366449	BFR2	9.18	4	6	0	0	0	0	0	0	0	0	0
398365717	TRM1	10.88	5	11	0	0	0	0	0	0	0	0	0
6323362	CHS5	11.03	5	12	0	0	0	0	0	0	0	0	0
398364573	PAN1	15.34	16	70	0	0	0	0	0	0	0	0	0
6320720	GRX2	0	0	0	0	0	0	0	0	0	0	0	0
6321595	PTI1	0	0	0	0	0	0	0	0	0	0	0	0
398364853	OCA1	0	0	0	0	0	0	0	0	0	0	0	0
6323852	YMR196	0	0	0	0	0	0	0	0	0	0	0	0
33438816	YILOO2W-	0	0	0	0	0	0	0	0	0	0	0	0
398366575	RRP17	0	0	0	0	0	0	0	0	0	0	0	0
6323359	TMA10	0	0	0	0	0	0	0	0	0	0	0	0
398365853	WTM1	13.96	5	20	1.83	1	1	3.20	1	1	0	0	0
6320181	GPD1	7.67	3	13	8.44	3	11	3.58	1	1	0	0	0
6323518	NAB6	1.23	1	1	0	0	0	1.06	1	1	0	0	0
6321221	KIP3	1.12	1	1	0	0	0	1.86	1	1	0	0	0
398365131	RFC2	9.63	3	4	0	0	0	2.27	1	1	0	0	0
6322294	ERG20	9.09	3	12	0	0	0	3.98	1	1	0	0	0

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6324550	IFM1	1.78	1	1	0	0	0	0	0	0	0	0	0
6324950	ALD4	1.73	1	1	0	0	0	0	0	0	0	0	0
6324635	CKA2	2.65	1	1	0	0	0	0	0	0	0	0	0
6319849	YCP4	4.86	1	1	0	0	0	0	0	0	0	0	0
6325202	LGE1	2.71	1	1	0	0	0	0	0	0	0	0	0
398366027	SUR4	4.93	1	1	0	0	0	0	0	0	0	0	0
6321641	PCT1	6.60	2	2	0	0	0	0	0	0	0	0	0
6325280	EAF3	6.48	2	2	0	0	0	0	0	0	0	0	0
6320803	MCM3	2.88	2	2	0	0	0	0	0	0	0	0	0
6323872	SKY1	4.58	2	2	0	0	0	0	0	0	0	0	0
6325140	IDI1	4.51	1	2	0	0	0	0	0	0	0	0	0
6320049	SAS10	10.82	4	4	0	0	0	0	0	0	0	0	0
6324989	DIM1	11.01	3	5	0	0	0	0	0	0	0	0	0
398366039	PBP1	9.97	5	7	0	0	0	0	0	0	0	0	0
6323088	REX2	9.67	2	8	0	0	0	0	0	0	0	0	0
6324899	PRO2	11.18	4	10	0	0	0	0	0	0	0	0	0
398365467	YNL010W	7.47	2	16	0	0	0	0	0	0	0	0	0
41629671	SEA4	13.58	11	21	0	0	0	0	0	0	0	0	0
6320704	PUF6	0	0	0	0	0	0	0	0	0	0	0	0
6323650	PLB1	0	0	0	0	0	0	0	0	0	0	0	0
6323158	DIP2	0	0	0	0	0	0	0	0	0	0	0	0
6322727	SRP21	0	0	0	0	0	0	0	0	0	0	0	0
14318504	MDJ1	0	0	0	0	0	0	0	0	0	0	0	0
6322878	GCN3	0	0	0	0	0	0	0	0	0	0	0	0
330443380	PSK1	0	0	0	0	0	0	0	0	0	0	0	0
398364789	UFD4	0	0	0	0	0	0	0	0	0	0	0	0
6319903	PWP2	0	0	0	0	0	0	0	0	0	0	0	0
6324613	CKB2	0	0	0	0	0	0	0	0	0	0	0	0
6324568	TOP1	0	0	0	0	0	0	0	0	0	0	0	0
398365393	MDM32	0	0	0	0	0	0	0	0	0	0	0	0
6322731	VPH2	0	0	0	0	0	0	0	0	0	0	0	0
6323653	HXT2	0	0	0	0	0	0	0	0	0	0	0	0
6320236	MRH1	0	0	0	0	0	0	0	0	0	0	0	0
398364559	NSP1	0	0	0	1.22	1	1	0.97	1	1	0	0	0
6321498	ADE6	8.39	9	14	1.25	1	2	1.18	1	1	0	0	0
6319326	ADE1	7.19	2	2	0	0	0	3.92	1	1	0	0	0
6321812	MYO1	1.30	2	2	0	0	0	0.57	1	1	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6324734	MTR10	0	0	0	0	0	0	0	0	0	0	0	0
6320284	PET100	0	0	0	0	0	0	0	0	0	0	0	0
6323411	SEC61	0	0	0	0	0	0	0	0	0	0	0	0
6320883	ERG28	0	0	0	0	0	0	0	0	0	0	0	0
6324593	YOR019W	1.78	1	5	1.78	1	1	1.23	1	1	0	0	0
6320432	ADK1	24.32	5	89	4.50	1	1	4.50	1	1	0	0	0
6322919	CCP1	0	0	0	2.77	1	1	4.71	1	1	0	0	0
6320204	NTH1	0	0	0	1.73	1	2	1.73	1	1	0	0	0
6320279	TPS2	1.67	1	2	6.14	4	9	1.67	1	1	0	0	0
6323681	CCS1	6.02	1	1	0	0	0	4.02	1	1	0	0	0
398366097	ADH6	8.06	2	4	0	0	0	5.00	1	1	0	0	0
6324381	NOG2	9.67	4	7	0	0	0	2.88	1	1	0	0	0
6323974	PRE5	22.65	5	7	0	0	0	5.98	1	1	0	0	0
6320090	ATG20	30.00	15	41	0	0	0	2.19	1	1	0	0	0
37362683	SCJ1	0	0	0	0	0	0	4.24	1	1	0	0	0
6323676	ARP9	0	0	0	0	0	0	2.78	1	1	0	0	0
398364547	SER3	2.35	1	1	0	0	0	7.04	2	2	0	0	0
6319439	HEK2	6.04	2	2	0	0	0	6.30	2	2	0	0	0
6321964	NMD3	6.56	3	7	2.70	1	1	2.70	1	3	0	0	0
6321984	ERG9	0	0	0	4.73	2	3	2.70	1	3	0	0	0
6323768	ADE17	15.88	7	33	0	0	0	2.53	1	3	0	0	0
398365401	CPA2	1.79	2	3	0	0	0	1.43	1	4	0	0	0
42742253	ATG27	11.44	3	7	0	0	0	11.07	2	5	0	0	0
6324191	SRV2	16.16	6	17	0	0	0	6.08	2	7	0	0	0
6322476	ILV3	10.26	6	53	1.54	1	5	2.39	1	9	0	0	0
6321900	GGA2	4.79	2	4	0	0	0	7.52	3	11	0	0	0
330443535	STE2	6.03	2	14	3.25	1	1	18.56	6	16	0	0	0
6324343	ACC1	9.54	19	54	4.75	8	23	4.12	6	18	0	0	0
6319434	APL3	4.00	3	3	1.07	1	1	0	0	0	0	0	0
398364593	MET6	14.86	10	14	1.17	1	1	0	0	0	0	0	0
6323751	YPK2	0	0	0	1.92	1	1	0	0	0	0	0	0
330443495	AIM6	0	0	0	3.59	1	1	0	0	0	0	0	0
6323752	PGM2	0	0	0	2.28	1	1	0	0	0	0	0	0
82795256	SEC28	3.38	1	1	5.41	1	2	0	0	0	0	0	0
6322215	YIR024C	0	0	0	4.63	1	2	0	0	0	0	0	0
6323291	YPT6	5.12	1	1	5.12	1	3	0	0	0	0	0	0
6321632	PDX1	4.63	2	2	12.20	2	5	0	0	0	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6320843	PAC2	0	0	0	0	0	0	0	0	0	0	0	0
398365529	SIR1	0	0	0	0	0	0	0	0	0	0	0	0
14318435	TIP20	0	0	0	0	0	0	0	0	0	0	0	0
6323645	AIM34	0	0	0	0	0	0	0	0	0	0	0	0
330443386	BUD14	0	0	0	0	0	0	0	0	0	0	0	0
6319751	CHK1	0	0	0	0	0	0	0	0	0	0	0	0
6319452	APN2	0	0	0	0	0	0	0	0	0	0	0	0
10383783	YCR015C	0	0	0	0	0	0	0	0	0	0	0	0
33438820	YJL133C-A	0	0	0	0	0	0	0	0	0	0	0	0
398365435	ERG25	0	0	0	0	0	0	0	0	0	0	0	0
6321363	MPS2	0	0	0	0	0	0	0	0	0	0	0	0
6323180	PCD1	0	0	0	0	0	0	0	0	0	0	0	0
398364799	PMD1	0	0	0	0	0	0	0	0	0	0	0	0
6322709	TGL1	0	0	0	0	0	0	0	0	0	0	0	0
6325305	TAH18	0	0	0	0	0	0	0	0	0	0	0	0
6322698	YKL151C	0	0	0	0	0	0	0	0	0	0	0	0
6322341	RPE1	0	0	0	0	0	0	0	0	0	0	0	0
6323336	STT4	0	0	0	0	0	0	0	0	0	0	0	0
6320542	MSN5	0	0	0	0	0	0	0	0	0	0	0	0
6325171	ELP3	0	0	0	0	0	0	0	0	0	0	0	0
6319274	ERV46	0	0	0	0	0	0	0	0	0	0	0	0
398366207	CRM1	2.58	3	4	1.38	1	1	1.66	1	1	0	0	0
6321198	CSE1	0.94	1	1	1.25	1	3	1.25	1	1	0	0	0
398366491	SKP1	12.89	1	1	0	0	0	8.76	1	1	0	0	0
6323819	MLH1	1.69	1	1	0	0	0	1.69	1	1	0	0	0
398365033	IRA1	0.36	1	1	0	0	0	0.32	1	1	0	0	0
398366427	SRP101	1.93	1	1	0	0	0	2.25	1	1	0	0	0
6320008	NUS1	4.00	1	1	0	0	0	1.87	1	1	0	0	0
6324098	PDR16	7.69	2	2	0	0	0	3.42	1	1	0	0	0
6324093	SIN4	2.36	2	2	0	0	0	1.75	1	1	0	0	0
6322063	MET18	2.23	2	2	0	0	0	1.74	1	1	0	0	0
6319901	THR4	7.00	3	4	0	0	0	3.70	1	1	0	0	0
6321899	CDC12	11.79	4	4	0	0	0	3.19	1	1	0	0	0
398364305	YND1	1.27	1	9	0	0	0	2.22	1	1	0	0	0
6325365	RPN7	19.11	9	39	0	0	0	2.10	1	1	0	0	0
6322288	RFA3	53.28	4	74	0	0	0	12.30	1	1	0	0	0
6323832	YMR178	0	0	0	0	0	0	6.20	1	1	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6321327	NSA1	0	0	0	0	0	0	3.02	1	1	0	0	0
398366645	PKH1	0	0	0	0	0	0	2.09	1	1	0	0	0
6323507	MSC1	0	0	0	0	0	0	1.75	1	1	0	0	0
398366107	YPP1	0	0	0	0	0	0	1.22	1	1	0	0	0
6323586	SPC2	0	0	0	0	0	0	6.74	1	1	0	0	0
6324079	RAD50	0	0	0	0	0	0	0.91	1	1	0	0	0
6320865	GAL83	0	0	0	0	0	0	2.64	1	1	0	0	0
14318465	YFL054C	0	0	0	0	0	0	2.17	1	1	0	0	0
6325349	NVJ2	0	0	0	0	0	0	2.21	1	1	0	0	0
6323962	UBP15	0.98	1	1	0.98	1	1	0.98	1	2	0	0	0
6323891	RNA1	6.88	2	3	3.19	1	1	7.86	2	2	0	0	0
6323273	MAP1	9.30	3	4	3.10	1	1	3.10	1	2	0	0	0
6320935	TRP2	12.43	5	6	1.97	1	1	6.71	2	2	0	0	0
6324994	YPL260W	2.54	1	1	1.81	1	2	4.72	2	2	0	0	0
27469358	PMT2	0	0	0	3.03	2	11	1.58	1	2	0	0	0
398365969	CDC31	8.70	1	1	0	0	0	8.70	1	2	0	0	0
6320214	SNQ2	1.87	2	4	0	0	0	0.80	1	2	0	0	0
6322263	PHO90	2.04	2	2	1.70	1	1	2.95	2	3	0	0	0
42742305	YNL208W	0	0	0	17.09	2	2	9.05	1	3	0	0	0
6321432	PMC1	0	0	0	2.13	2	4	2.81	2	3	0	0	0
6323221	HCR1	8.30	2	4	4.15	1	5	4.15	1	3	0	0	0
6322722	PGM1	2.46	1	1	0	0	0	2.28	1	3	0	0	0
6323089	FRS1	4.71	2	2	0	0	0	4.20	2	3	0	0	0
6321271	PMR1	3.47	3	3	0	0	0	3.37	2	3	0	0	0
6322097	LYS12	20.49	6	35	0	0	0	7.01	2	3	0	0	0
6320151	SLC1	0	0	0	0	0	0	5.94	1	3	0	0	0
6319312	ERP2	0	0	0	0	0	0	12.09	1	3	0	0	0
398365779	YCF1	1.98	2	2	0	0	0	2.05	2	4	0	0	0
6324082	YNL247W	3.91	3	3	0	0	0	3.39	2	4	0	0	0
6324625	ETT1	0	0	0	0	0	0	2.91	1	4	0	0	0
6321492	MUP1	0	0	0	0	0	0	1.74	1	4	0	0	0
6322434	RNR2	9.52	4	13	2.76	1	7	7.02	2	5	0	0	0
6320245	ENA1	1.01	1	2	0	0	0	2.66	2	5	0	0	0
6323853	VTI1	24.42	4	7	0	0	0	14.75	2	5	0	0	0
6319837	ILV6	5.18	1	6	5.18	1	1	9.06	2	6	0	0	0
6319441	PET9	3.46	1	1	16.04	5	14	3.14	1	6	0	0	0
398365855	KGD2	2.81	1	1	6.26	2	5	10.58	3	7	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6324988	ACM1	0	0	0	0	0	0	0	0	0	0	0	0
330443417	RRN6	0	0	0	0	0	0	0	0	0	0	0	0
6325398	TAZ1	0	0	0	0	0	0	0	0	0	0	0	0
6321828	MSC7	0	0	0	0	0	0	0	0	0	0	0	0
6323320	GUF1	0	0	0	0	0	0	0	0	0	0	0	0
16740526	GYP8	0	0	0	0	0	0	0	0	0	0	0	0
6324783	NPT1	2.56	1	1	2.56	1	1	3.96	1	1	0	0	0
6319974	HOP	5.97	2	2	2.73	1	1	2.73	1	1	0	0	0
6319931	ABP1	15.20	6	18	2.03	1	1	2.20	1	1	0	0	0
6321787	OSH7	0	0	0	1.83	1	1	1.83	1	1	0	0	0
6319512	CHS2	0	0	0	0.83	1	1	1.66	1	1	0	0	0
6324912	KRE5	0	0	0	0.88	1	1	0.73	1	1	0	0	0
398365337	SFL1	0	0	0	6.14	1	1	2.74	1	1	0	0	0
398364433	CAJ1	7.16	2	10	3.58	1	2	3.58	1	1	0	0	0
6322547	EMC2	0	0	0	3.77	1	2	6.16	1	1	0	0	0
6324225	LEU4	3.55	1	1	3.88	2	4	2.42	1	1	0	0	0
6322537	MIR1	0	0	0	3.86	1	4	3.86	1	1	0	0	0
6321795	ERG11	0	0	0	9.81	3	15	3.40	1	1	0	0	0
6323953	YMR295C	0	0	0	15.74	1	28	12.18	1	1	0	0	0
398366013	YDR186C	1.94	1	1	0	0	0	2.17	1	1	0	0	0
6323943	DSS1	1.14	1	1	0	0	0	1.14	1	1	0	0	0
6320851	HEM14	1.48	1	1	0	0	0	2.78	1	1	0	0	0
6322116	SER33	2.35	1	1	0	0	0	2.99	1	1	0	0	0
83578105	TMA46	2.90	1	1	0	0	0	3.19	1	1	0	0	0
6321960	CDC23	2.08	1	1	0	0	0	1.76	1	1	0	0	0
6320470	AKR1	1.70	1	1	0	0	0	3.40	1	1	0	0	0
398365797	YNR048W	3.05	1	1	0	0	0	3.05	1	1	0	0	0
6322387	SMC3	0.81	1	1	0	0	0	1.22	1	1	0	0	0
6321806	ARG4	1.94	1	1	0	0	0	2.59	1	1	0	0	0
6324055	GOR1	2.57	1	1	0	0	0	3.43	1	1	0	0	0
6324554	YOL019W	2.18	1	1	0	0	0	2.72	1	1	0	0	0
6324897	PMT3	1.59	1	1	0	0	0	1.73	1	1	0	0	0
6323553	ALO1	1.90	1	1	0	0	0	2.66	1	1	0	0	0
6323958	ADE4	4.12	2	2	0	0	0	2.35	1	1	0	0	0
6322585	ENT3	2.70	1	2	0	0	0	3.92	1	1	0	0	0
6322732	SBA1	4.17	1	2	0	0	0	4.17	1	1	0	0	0
6319767	BSD2	11.53	1	2	0	0	0	4.05	1	1	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398365317	GPT2	2.02	1	2	0	0	0	2.02	1	1	0	0	0
82795241	APE3	6.33	2	2	0	0	0	4.10	1	1	0	0	0
398365865	GIR2	5.28	1	2	0	0	0	4.15	1	1	0	0	0
6322509	UTR1	2.45	1	2	0	0	0	2.45	1	1	0	0	0
6322500	GEF1	3.21	2	2	0	0	0	2.31	1	1	0	0	0
6320893	HOM3	2.66	1	2	0	0	0	2.09	1	1	0	0	0
82795247	LSM6	34.88	3	3	0	0	0	20.93	1	1	0	0	0
6320249	HEM13	9.45	3	3	0	0	0	6.10	1	1	0	0	0
6322287	KRE9	3.99	1	3	0	0	0	9.78	1	1	0	0	0
6324580	SGT2	13.58	4	5	0	0	0	4.05	1	1	0	0	0
6322870	YKRO18C	6.34	4	7	0	0	0	2.07	1	1	0	0	0
14318569	DUG1	8.94	3	14	0	0	0	3.33	1	1	0	0	0
6319509	PDX3	17.98	3	31	0	0	0	5.26	1	1	0	0	0
6324697	LEO1	19.83	7	50	0	0	0	1.94	1	1	0	0	0
6319764	YBR287W	0	0	0	0	0	0	6.09	1	1	0	0	0
6324484	MPD2	0	0	0	0	0	0	5.42	1	1	0	0	0
6323510	PGA3	0	0	0	0	0	0	4.49	1	1	0	0	0
6321646	CIR1	0	0	0	0	0	0	6.13	1	1	0	0	0
6324059	ALP1	0	0	0	0	0	0	2.44	1	1	0	0	0
6320516	SUM1	0	0	0	0	0	0	1.51	1	1	0	0	0
37362655	COG2	0	0	0	0	0	0	2.67	1	1	0	0	0
398366091	ART10	0	0	0	0	0	0	2.32	1	1	0	0	0
6320327	KIN1	0	0	0	0	0	0	1.60	1	1	0	0	0
6324609	SHE4	0	0	0	0	0	0	1.39	1	1	0	0	0
6324132	WHI3	0	0	0	0	0	0	2.42	1	1	0	0	0
6322550	JSN1	0	0	0	0	0	0	1.65	1	1	0	0	0
6320107	PMT1	0	0	0	0	0	0	1.59	1	1	0	0	0
6319413	SHP1	0	0	0	0	0	0	4.02	1	1	0	0	0
6321756	GOS1	0	0	0	0	0	0	11.66	1	1	0	0	0
6321337	YGL101W	0	0	0	0	0	0	5.12	1	1	0	0	0
6325199	PDR12	0	0	0	0	0	0	0.99	1	1	0	0	0
6322782	NUP100	0	0	0	0	0	0	1.25	1	1	0	0	0
6324859	RDL1	0	0	0	0	0	0	10.79	1	1	0	0	0
6323899	ZRC1	0	0	0	0	0	0	3.85	1	1	0	0	0
6319392	NUP170	0	0	0	0	0	0	1.33	1	1	0	0	0
398365585	PCS60	0	0	0	0	0	0	3.31	1	1	0	0	0
6322186	EPS1	0	0	0	0	0	0	2.57	1	1	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398366407	YOR1	0	0	0	0	0	0	0.95	1	1	0	0	0
37362695	OST2	0	0	0	0	0	0	13.08	1	1	0	0	0
6324353	SEC12	0	0	0	0	0	0	2.12	1	1	0	0	0
330443368	FLC2	0	0	0	0	0	0	2.17	1	1	0	0	0
398365223	MSP1	0	0	0	0	0	0	3.87	1	1	0	0	0
6321449	YGR012W	0	0	0	0	0	0	3.56	1	1	0	0	0
6322310	SNA3	0	0	0	0	0	0	24.81	1	1	0	0	0
6323413	CTF3	0	0	0	0	0	0	1.23	1	1	0	0	0
6321225	SKI8	0	0	0	0	0	0	2.02	1	1	0	0	0
6325101	PRM4	0	0	0	0	0	0	5.28	1	1	0	0	0
6321345	SPC105	0	0	0	0	0	0	1.64	1	1	0	0	0
398366519	EAF1	0	0	0	0	0	0	1.43	1	1	0	0	0
398365305	PAM17	0	0	0	0	0	0	9.64	1	1	0	0	0
6319334	PRM9	0	0	0	0	0	0	4.70	1	1	0	0	0
83722565	PTM1	0	0	0	0	0	0	1.91	1	1	0	0	0
6322825	TCD2	0	0	0	0	0	0	3.58	1	1	0	0	0
6323936	CAT8	0	0	0	0	0	0	1.05	1	1	0	0	0
6324567	CSI2	0	0	0	0	0	0	5.28	1	1	0	0	0
6319516	CST26	0	0	0	0	0	0	4.28	1	1	0	0	0
6321275	RAD54	0	0	0	0	0	0	1.56	1	1	0	0	0
398366375	DON1	0	0	0	0	0	0	4.11	1	1	0	0	0
398365229	EAF6	0	0	0	0	0	0	10.62	1	1	0	0	0
398365911	ESA1	0	0	0	0	0	0	2.47	1	1	0	0	0
6323375	GAS2	0	0	0	0	0	0	1.44	1	1	0	0	0
398366357	GDS1	0	0	0	0	0	0	2.87	1	1	0	0	0
330443484	YCL002C	0	0	0	0	0	0	6.08	1	1	0	0	0
6321588	YGR149W	0	0	0	0	0	0	3.01	1	1	0	0	0
37362691	ITR2	0	0	0	0	0	0	2.13	1	1	0	0	0
398364419	MRF1	0	0	0	0	0	0	3.15	1	1	0	0	0
6324909	MRS2	0	0	0	0	0	0	3.40	1	1	0	0	0
398364395	PEX14	0	0	0	0	0	0	4.69	1	1	0	0	0
14318528	YFR006W	0	0	0	0	0	0	3.36	1	1	0	0	0
6321021	RAD24	0	0	0	0	0	0	2.58	1	1	0	0	0
6319451	RFT1	0	0	0	0	0	0	2.09	1	1	0	0	0
398365357	RRG1	0	0	0	0	0	0	2.74	1	1	0	0	0
14318556	RRT5	0	0	0	0	0	0	3.81	1	1	0	0	0
6321943	SKG6	0	0	0	0	0	0	1.77	1	1	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398365903	SSP2	0	0	0	0	0	0	2.70	1	1	0	0	0
398366103	CST9	0	0	0	0	0	0	1.66	1	1	0	0	0
398364315	TOS3	0	0	0	0	0	0	2.50	1	1	0	0	0
6321577	TPO2	0	0	0	0	0	0	3.75	1	1	0	0	0
6320230	VPS54	0	0	0	0	0	0	1.80	1	1	0	0	0
6321759	WSC4	0	0	0	0	0	0	2.81	1	1	0	0	0
6322315	IDS2	0	0	0	0	0	0	2.13	1	1	0	0	0
6319635	IFA38	0	0	0	0	0	0	3.17	1	1	0	0	0
6323014	BPT1	0	0	0	0	0	0	1.15	1	1	0	0	0
6323657	ERG5	0	0	0	0	0	0	2.79	1	1	0	0	0
6323920	CUE1	7.39	1	1	7.39	1	1	7.39	1	2	0	0	0
86558908	CBR1	2.82	1	1	3.87	1	1	3.87	1	2	0	0	0
6323859	TOM40	0	0	0	4.13	1	1	4.13	1	2	0	0	0
398364295	WBP1	0	0	0	2.56	1	1	6.74	2	2	0	0	0
6322066	KGD1	0.99	1	1	2.37	2	2	2.47	2	2	0	0	0
6323593	GSF2	0	0	0	6.95	2	3	7.94	2	2	0	0	0
6322127	ARC15	32.47	3	12	6.49	1	4	6.49	1	2	0	0	0
10383789	NPP1	1.62	1	1	0	0	0	1.75	1	2	0	0	0
6319477	RCR1	7.04	1	1	0	0	0	6.10	1	2	0	0	0
398366155	AHA1	8.86	2	2	0	0	0	5.14	1	2	0	0	0
398365557	VPS27	4.18	2	3	0	0	0	2.25	1	2	0	0	0
6320054	RPN5	4.04	2	3	0	0	0	4.27	1	2	0	0	0
398365953	SUP35	5.84	3	3	0	0	0	4.96	2	2	0	0	0
37362677	AAT2	9.33	3	4	0	0	0	3.11	1	2	0	0	0
6324511	PRS5	11.29	4	4	0	0	0	3.63	1	2	0	0	0
6324463	ZEO1	24.78	3	6	0	0	0	29.20	2	2	0	0	0
6322338	MTC1	4.60	2	14	0	0	0	2.30	1	2	0	0	0
6324198	TOM22	0	0	0	0	0	0	11.18	1	2	0	0	0
330443603	YJL171C	0	0	0	0	0	0	4.29	1	2	0	0	0
6322173	SNL1	0	0	0	0	0	0	10.69	1	2	0	0	0
6323876	ERG8	0	0	0	0	0	0	6.87	2	2	0	0	0
6321817	DAP2	0	0	0	0	0	0	1.83	1	2	0	0	0
6324071	DSL1	0	0	0	0	0	0	1.99	1	2	0	0	0
6325204	KTR6	0	0	0	0	0	0	6.73	2	2	0	0	0
398365745	MTC4	0	0	0	0	0	0	3.60	1	2	0	0	0
398366001	PNT1	0	0	0	0	0	0	3.07	1	2	0	0	0
6320424	SPR28	0	0	0	0	0	0	2.60	1	2	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6321400	OCH1	0	0	0	0	0	0	3.75	1	2	0	0	0
6321416	STT3	0	0	0	0	0	0	1.95	1	2	0	0	0
6321206	EMC4	0	0	0	0	0	0	11.05	1	2	0	0	0
6320071	LYS21	22.50	7	15	2.27	1	1	5.45	2	3	0	0	0
330443405	CDC27	0	0	0	1.45	1	1	1.72	1	3	0	0	0
14318471	EMP47	0	0	0	2.70	1	1	5.39	2	3	0	0	0
6322765	MDH1	9.88	2	2	4.19	1	2	4.19	1	3	0	0	0
6322421	NUP192	0	0	0	1.01	1	2	0.77	1	3	0	0	0
6323619	APT1	5.35	1	1	0	0	0	5.35	1	3	0	0	0
6321355	SCY1	1.37	1	1	0	0	0	4.10	3	3	0	0	0
6324610	PEP12	32.64	7	16	0	0	0	4.86	1	3	0	0	0
6320019	LYS20	29.44	9	17	0	0	0	5.61	2	3	0	0	0
6323056	ADE16	9.98	5	30	0	0	0	2.54	1	3	0	0	0
6321748	EFM1	0	0	0	0	0	0	2.56	1	3	0	0	0
6324110	ALG9	0	0	0	0	0	0	3.42	1	3	0	0	0
6319869	HSP30	0	0	0	0	0	0	6.63	1	3	0	0	0
6325200	SUR1	0	0	0	0	0	0	4.19	1	3	0	0	0
6320451	MNN10	0	0	0	0	0	0	5.85	2	3	0	0	0
6319914	CPR4	0	0	0	0	0	0	4.72	1	3	0	0	0
6323998	AAD14	0	0	0	0	0	0	4.52	1	3	0	0	0
6323509	ERG13	8.96	4	31	3.26	1	1	6.52	2	4	0	0	0
6320560	TRR1	21.32	4	9	4.70	1	3	4.70	1	4	0	0	0
398364773	TOM70	1.94	1	1	0	0	0	4.21	2	4	0	0	0
6319497	CHS3	0.77	1	1	0	0	0	6.01	3	4	0	0	0
398365065	DID2	3.92	1	2	0	0	0	6.86	1	4	0	0	0
330443526	ALD5	17.12	7	8	0	0	0	5.19	2	4	0	0	0
6319421	SEC17	19.52	4	11	0	0	0	9.59	2	4	0	0	0
398366327	VMA6	14.78	4	24	0	0	0	5.51	1	4	0	0	0
6325220	EGD1	0	0	0	0	0	0	12.10	1	4	0	0	0
6324160	PSD1	0	0	0	0	0	0	7.00	2	4	0	0	0
398364599	BAP2	0	0	0	0	0	0	3.45	1	4	0	0	0
6325035	FLC1	0	0	0	0	0	0	1.89	1	4	0	0	0
6322683	MCD4	0	0	0	0	0	0	1.31	1	4	0	0	0
6321502	VHT1	0	0	0	0	0	0	4.38	2	4	0	0	0
6321437	ERG26	0	0	0	0	0	0	4.87	1	4	0	0	0
6320793	GDA1	11.00	4	5	1.93	1	1	5.79	2	5	0	0	0
398365875	SKN1	1.95	1	8	1.82	1	1	1.95	1	5	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398364717	OLE1	0	0	0	2.35	1	1	8.63	3	5	0	0	0
398364715	OST1	0	0	0	2.94	1	2	6.30	2	5	0	0	0
6320639	CYM1	0.81	1	1	0	0	0	1.01	1	5	0	0	0
6324109	ADE12	3.70	2	2	0	0	0	4.39	2	5	0	0	0
330443698	CHS1	2.12	2	2	0	0	0	2.65	2	5	0	0	0
398365123	AQR1	2.56	1	2	0	0	0	4.61	3	5	0	0	0
6324745	LCB4	4.81	2	2	0	0	0	5.29	2	5	0	0	0
398366587	RPN9	11.20	5	5	0	0	0	3.82	1	5	0	0	0
6320837	MNN1	0	0	0	0	0	0	3.54	2	5	0	0	0
6324594	HSP10	19.81	2	4	10.38	1	2	10.38	1	6	0	0	0
6320859	RPN3	13.00	5	9	5.16	2	3	5.16	2	6	0	0	0
398366611	NHX1	3.16	1	1	0	0	0	4.58	1	6	0	0	0
6320395	SLY1	2.85	2	2	0	0	0	6.91	3	6	0	0	0
41629692	YIP4	8.51	2	4	0	0	0	20.43	2	6	0	0	0
330443374	DRS2	3.62	4	7	0	0	0	1.99	2	6	0	0	0
6325207	MNN9	0	0	0	0	0	0	2.78	1	6	0	0	0
6323858	ERG20	0	0	0	0	0	0	6.31	1	6	0	0	0
6319489	MNN2	8.38	3	3	2.35	1	1	4.02	2	7	0	0	0
6319726	ARO4	10.27	3	12	4.32	1	1	4.32	1	7	0	0	0
398364279	VAC8	1.90	1	1	9.17	4	8	2.42	1	7	0	0	0
6324555	TLG2	13.35	4	9	0	0	0	13.85	4	7	0	0	0
398366239	GNT1	1.22	1	19	0	0	0	5.09	2	7	0	0	0
6322249	OPT1	0	0	0	0	0	0	3.38	2	7	0	0	0
6319454	PEP1	14.06	18	90	2.03	2	2	1.71	2	8	0	0	0
6322787	YKL063C	0	0	0	0	0	0	8.38	1	8	0	0	0
6324645	NRT1	0	0	0	0	0	0	5.02	2	8	0	0	0
6323270	YLR241W	1.28	1	2	2.94	2	4	2.94	2	10	0	0	0
14318577	RPN12	10.22	2	5	0	0	0	10.22	2	10	0	0	0
6324061	LYP1	0	0	0	0	0	0	4.42	2	10	0	0	0
6320993	FTR1	0	0	0	0	0	0	9.90	3	10	0	0	0
6320705	ITR1	0	0	0	0	0	0	4.62	2	11	0	0	0
6319319	ERP1	5.48	1	2	5.48	1	6	17.81	3	12	0	0	0
6324828	SEC63	0	0	0	3.47	2	6	9.95	5	13	0	0	0
6319429	FUI1	0	0	0	2.35	1	3	8.45	4	16	0	0	0
6325293	VMA13	8.16	4	18	0	0	0	12.34	4	16	0	0	0
10383762	RNQ1	4.20	1	1	4.20	1	1	9.88	2	18	0	0	0
6320691	KRE2	10.41	3	3	2.94	1	3	17.19	5	19	0	0	0

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Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6323630	ERV25	6.16	1	8	13.74	2	9	6.16	1	19	0	0	0
6323716	TVP18	8.38	1	2	0	0	0	8.38	1	19	0	0	0
6319289	SNC1	0	0	0	0	0	0	13.68	1	24	0	0	0
398364281	EMP24	0	0	0	6.90	1	3	22.17	3	25	0	0	0
6320298	DNF2	0.87	1	1	1.55	1	4	5.77	6	26	0	0	0
330443597	MNN5	1.88	1	1	0	0	0	9.56	4	26	0	0	0
6320897	FCY2	3.00	1	2	0	0	0	10.51	5	31	0	0	0
330443542	SAP155	0.90	1	1	1.60	1	1	0	0	0	0	0	0
14318521	SPB4	1.49	1	1	1.82	1	1	0	0	0	0	0	0
6323367	NUP2	2.50	1	1	1.53	1	1	0	0	0	0	0	0
6319303	TPD3	3.62	1	1	1.89	1	1	0	0	0	0	0	0
6324166	RIA1	1.17	1	1	0.72	1	1	0	0	0	0	0	0
398364557	AIM21	1.62	1	1	3.39	1	1	0	0	0	0	0	0
6322369	SRS2	0.68	1	1	1.02	1	1	0	0	0	0	0	0
6324088	ZWF1	3.37	1	1	3.96	1	1	0	0	0	0	0	0
6319890	TAF2	0.92	1	1	0.57	1	1	0	0	0	0	0	0
6320033	SFA1	2.33	1	1	2.07	1	1	0	0	0	0	0	0
6325429	BSP1	1.74	1	1	1.39	1	1	0	0	0	0	0	0
6323022	YLL007C	2.41	1	1	1.50	1	1	0	0	0	0	0	0
6324918	YOR342C	3.76	1	1	3.76	1	1	0	0	0	0	0	0
6320091	TRM3	0.63	1	1	0.70	1	1	0	0	0	0	0	0
6322925	SIS2	4.27	2	2	1.78	1	1	0	0	0	0	0	0
6321999	SCH9	1.82	1	2	1.58	1	1	0	0	0	0	0	0
6320106	RPN6	2.76	1	2	2.76	1	1	0	0	0	0	0	0
6320934	PTC2	4.53	2	2	4.53	1	1	0	0	0	0	0	0
6322545	STE18	15.45	1	2	8.18	1	1	0	0	0	0	0	0
27808714	LAP3	2.42	1	2	3.08	1	1	0	0	0	0	0	0
6322410	IRC8	2.07	2	2	0.97	1	1	0	0	0	0	0	0
14318532	UBP6	6.01	2	3	3.21	1	1	0	0	0	0	0	0
6320129	BRE1	5.14	3	3	1.29	1	1	0	0	0	0	0	0
6324287	COG6	2.74	2	3	1.31	1	1	0	0	0	0	0	0
6320846	NTF2	25.60	3	3	12.80	1	1	0	0	0	0	0	0
6324508	MET22	5.60	2	3	6.44	1	1	0	0	0	0	0	0
6319975	GCS1	8.52	2	4	5.40	1	1	0	0	0	0	0	0
63535562	CHA1	10.83	4	4	3.89	1	1	0	0	0	0	0	0
6322141	NEO1	2.17	3	5	1.48	1	1	0	0	0	0	0	0
398365175	CDC28	11.07	3	7	2.35	1	1	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
398366131	ADE3	10.04	8	13	1.59	1	1	0	0	0	0	0	0
6322020	VTH1	2.84	4	39	1.55	1	1	0	0	0	0	0	0
6324916	RPA43	21.17	6	87	3.68	1	1	0	0	0	0	0	0
6323757	HFD1	0	0	0	1.69	1	1	0	0	0	0	0	0
6323256	ADY4	0	0	0	2.23	1	1	0	0	0	0	0	0
6320504	ATP5	0	0	0	6.13	1	1	0	0	0	0	0	0
6324144	YML19	0	0	0	13.29	1	1	0	0	0	0	0	0
398365747	DAM1	0	0	0	2.62	1	1	0	0	0	0	0	0
398365791	MTQ2	0	0	0	3.62	1	1	0	0	0	0	0	0
6323109	EMP46	0	0	0	3.15	1	1	0	0	0	0	0	0
398365027	RPN10	0	0	0	7.09	1	1	0	0	0	0	0	0
6322634	OXP1	0	0	0	0.70	1	1	0	0	0	0	0	0
6323485	SST2	0	0	0	1.86	1	1	0	0	0	0	0	0
6322729	OAC1	0	0	0	4.94	1	1	0	0	0	0	0	0
6323798	SWP1	0	0	0	3.85	1	1	0	0	0	0	0	0
6321694	COQ6	0	0	0	2.71	1	1	0	0	0	0	0	0
6321251	COX4	0	0	0	12.90	1	1	0	0	0	0	0	0
50593504	YPL109C	0	0	0	2.89	1	1	0	0	0	0	0	0
6324006	LEM3	0	0	0	2.66	1	1	0	0	0	0	0	0
6323581	NTE1	0	0	0	1.01	1	1	0	0	0	0	0	0
6320666	HEH2	0	0	0	1.06	1	1	0	0	0	0	0	0
6323377	YLR345W	0	0	0	1.77	1	1	0	0	0	0	0	0
6325244	MRPS16	0	0	0	10.74	1	1	0	0	0	0	0	0
398365817	STE13	0	0	0	1.61	1	1	0	0	0	0	0	0
6322786	MNR2	0	0	0	1.34	1	1	0	0	0	0	0	0
6323299	DCS1	0	0	0	4.00	1	1	0	0	0	0	0	0
6323777	POM152	0	0	0	1.27	1	1	0	0	0	0	0	0
6323521	VAN1	0	0	0	2.43	1	1	0	0	0	0	0	0
6321838	YHK8	0	0	0	1.75	1	1	0	0	0	0	0	0
6321688	MGA1	0	0	0	1.97	1	1	0	0	0	0	0	0
6323415	SMC6	0	0	0	0.72	1	1	0	0	0	0	0	0
398365989	YLR361C-	0	0	0	12.24	1	1	0	0	0	0	0	0
6320856	ISC1	0	0	0	2.10	1	1	0	0	0	0	0	0
6323515	NDI1	0	0	0	3.90	1	1	0	0	0	0	0	0
6322031	POT1	0	0	0	1.68	1	1	0	0	0	0	0	0
14318490	CAK1	0	0	0	2.45	1	1	0	0	0	0	0	0
6325257	CIT3	0	0	0	2.47	1	1	0	0	0	0	0	0

Table S1 - Smith-Kinnaman, *et al* (2014) Abbreviations: %SC = percent sequence coverage, UP = unique peptides, PSM = peptide spectrum matches

Accession	Acronym	%SC FLAG_Rtr 1_1	UP FLAG_Rtr1 _1	# PSM FLAG_Rtr 1_1	%SC FLAG_Rtr1_ ctk1_deleti on	UP FLAG_Rtr 1_ctk1_d eletion	# PSM FLAG_Rtr 1_ctk1_d eletion	%SC FLAG_Rtr1 _2	UP FLAG_Rtr 1_2	# PSM FLAG_Rtr 1_2	%SC Rpb3_TA P_WT	UP Rpb3_TA P_WT	# PSM Rpb3_TA P_WT
6323257	ECM22	0	0	0	1.23	1	1	0	0	0	0	0	0
6323492	GAB1	0	0	0	2.54	1	1	0	0	0	0	0	0
6322779	YKL071W	0	0	0	3.13	1	1	0	0	0	0	0	0
330443687	YMR317	0	0	0	1.67	1	1	0	0	0	0	0	0
6322216	MND2	0	0	0	4.08	1	1	0	0	0	0	0	0
9755337	NYV1	0	0	0	3.95	1	1	0	0	0	0	0	0
398366349	RIF2	0	0	0	4.05	1	1	0	0	0	0	0	0
6323548	RPM2	0	0	0	0.75	1	1	0	0	0	0	0	0
398364459	SCS3	0	0	0	3.95	1	1	0	0	0	0	0	0
6323956	LIP1	0	0	0	6.00	1	1	0	0	0	0	0	0
37362686	STB1	0	0	0	2.62	1	1	0	0	0	0	0	0
6321342	TOS8	0	0	0	3.99	1	1	0	0	0	0	0	0
6323000	TPO1	0	0	0	2.73	1	1	0	0	0	0	0	0
398364289	VAB2	0	0	0	3.90	1	1	0	0	0	0	0	0
6324803	WTM2	0	0	0	2.36	1	1	0	0	0	0	0	0
398366213	LYS4	0	0	0	2.74	1	1	0	0	0	0	0	0
6323285	HAP1	0	0	0	1.07	1	1	0	0	0	0	0	0
6321912	MSH1	0	0	0	0.94	1	1	0	0	0	0	0	0
10383785	CWH43	0	0	0	1.26	1	1	0	0	0	0	0	0
398364257	SPF1	0	0	0	2.06	1	1	0	0	0	0	0	0
6323304	MCM5	1.03	1	1	1.94	1	2	0	0	0	0	0	0
398366185	YLR419W	0.77	1	1	0.70	1	2	0	0	0	0	0	0
6319366	SFT2	5.58	1	1	5.58	1	2	0	0	0	0	0	0
6324749	ALE1	2.26	1	1	2.26	1	2	0	0	0	0	0	0
6325112	KES1	2.30	1	2	3.23	1	2	0	0	0	0	0	0
6323682	SUB1	14.04	3	3	4.79	1	2	0	0	0	0	0	0
6325049	TYW1	2.84	2	3	1.23	1	2	0	0	0	0	0	0
6325179	ATP4	17.62	4	4	12.30	2	2	0	0	0	0	0	0
6323882	YMR226C	19.85	4	5	10.11	2	2	0	0	0	0	0	0
6323505	ERO1	0	0	0	2.31	1	2	0	0	0	0	0	0
37362643	GET2	0	0	0	3.51	1	2	0	0	0	0	0	0
14318552	PTR3	0	0	0	1.47	1	2	0	0	0	0	0	0
6320179	RTK1	2.26	1	1	1.13	1	3	0	0	0	0	0	0
6324513	GPD2	4.77	2	2	3.41	1	3	0	0	0	0	0	0
398364869	APC11	0	0	0	4.24	1	3	0	0	0	0	0	0
6319912	SED4	0	0	0	5.16	2	3	0	0	0	0	0	0
6325029	ALG5	2.69	1	2	2.69	1	4	0	0	0	0	0	0

