

Supplementary Table 2

Master_id_first	PsitelnProtein	diff	FDR
P84098	S189	-1.670489281	2.81E-04
Q9NRR4	S217_S221	-1.123886299	0.042133206
O15318	S157	-1.011674643	3.33E-06
Q96RK0	S436	-0.955248207	0.005313429
P11388	S1377	-0.890785582	4.64E-04
Q9UHB7	SSPRPTAEK	-0.744292381	0.012496476
Q86YS7	S295	-0.724145059	2.27E-04
Q12830	S1310	-0.720303255	2.75E-05
Q9UQ35	S222	-0.714191545	3.12E-04
Q58WW2	S847_S850	-0.710412426	0.045456295
P55327	S36	-0.702963265	6.88E-05
O95359	S2317	-0.650783117	5.09E-04
Q9UGJ0	S71	-0.644264742	6.58E-05
Q9UPW6	S39	-0.64230502	0.006363026
Q96T58	T1619	-0.628494198	0.048388526
Q5VZ89	S1225	0.601622749	0.001690327
A6H8Y1	S100	0.605046658	0.004766805
O43765	S305	0.605124332	9.75E-04
P50747	S79	0.6067396	0.003477569
Q9UBH6	S676	0.612103292	0.002709452
Q8NFC6	S1077	0.612352517	1.69E-05
Q9BW71	S269	0.616715941	0.013969911
P20794	Y159	0.621540647	0.035205852
Q8IWN7	S1671	0.621928479	0.004060354
Q16644	T201	0.636610904	0.001132634
Q9NW75	S129	0.638526346	0.016179326
P15056	S429	0.642108443	0.001173429
O43395	T172	0.64835402	0.001919793
P50548	T526	0.64887295	4.81E-05
Q2NKG8	FDASTPKNDISPP	0.649344065	9.14E-04
Q8NDI1	S649	0.656249797	2.60E-04
Q86VW2	S556_T565_S/T	0.658502784	7.94E-04
Q96D71	S709	0.658641067	1.08E-04
Q9Y3R0	S996	0.660215311	0.003485813
O75152	S758	0.672977667	3.11E-05
Q7LBC6	T1270	0.673014802	4.55E-04
Q5T3J3	T389	0.677200574	3.26E-04
Q8TF72	S285	0.690682971	6.32E-04
Q7Z4S6	S853	0.691495771	1.70E-04
P55196	T1352	0.694573587	3.27E-05
O75151	S1056_S/T	0.696891464	0.043890913
Q6PKG0	T376	0.697518943	8.85E-06
Q2LD37	S4124	0.698401708	8.41E-04
Q5T5P2	S474	0.699718392	4.38E-05
Q8IY17	S372	0.700691502	0.003477569
Q8IWB9	S270_S/T	0.706474401	0.008155201
P23588	S230	0.70734361	6.87E-04
Q9NZJ0	S557	0.713796214	7.21E-04
Q5U5Q3	S537_S/T	0.714020432	0.0123582
Q9P266	S1194	0.714525412	1.13E-04
Q9Y6Y0	T330	0.719599582	3.82E-06
O94854	S578	0.727203002	2.54E-04
O75151	S625	0.727555284	0.017760793
Q9HCH0	S1295	0.736508525	0.009349702

Q9P2B2	S875	0.740392655	0.037535384
Q5SW79	S1160_S1165	0.743910193	3.32E-06
Q15691	S155	0.745540061	3.82E-06
Q96RK0	S173	0.74998776	5.09E-04
Q9UGU5	S197	0.759878196	1.55E-04
Q9C0B5	S409	0.774603299	3.25E-06
Q5SW79	T174	0.776978059	0.019969504
Q86SQ0	MISSISGRDDLMD	0.778628242	0.001297773
Q6EEV4	LRAPSSR	0.780686954	8.99E-04
Q9UKA4	S433	0.794625722	0.003326958
Q8WXD9	S826	0.798325002	5.05E-05
O43379	S49	0.803109157	1.84E-04
Q9BY44	S526	0.808955982	9.14E-04
Q07889	S1134	0.811811249	9.68E-05
Q9UPP1	S1021	0.820420955	4.39E-04
Q9H6F5	S217	0.821211657	0.002724479
Q9Y6X9	S739_S743	0.825922634	0.006955041
P0CAP2	S179	0.829747668	0.019228924
P09884	T219	0.836571736	5.24E-07
O14686	S2726	0.863333634	1.55E-04
Q9UPN7	S726	0.867783888	0.006027765
Q2LD37	S3653	0.884869812	1.54E-05
P51812	S369_S375	0.887843423	1.33E-06
O60934	T497	0.892160715	2.16E-04
Q14934	S272	0.892917386	0.001536624
Q765P7	S601	0.900876186	5.38E-07
Q14160	T1342_S1348	0.930576518	2.55E-04
O15357	T958	0.933498436	1.68E-05
P10398	T181	0.955592066	7.81E-06
Q5TKA1	S65	0.961395971	2.39E-04
Q86SQ0	S212	0.961723677	7.87E-04
Q5VT52	S581	0.965151101	2.09E-04
Q9UKJ3	T293	0.969989181	6.66E-07
Q8N5U6	S128	0.974062795	6.66E-07
O76039	S407	1.001394554	5.85E-08
P41162	TESSPGSR	1.003331325	0.002537304
P47974	S125	1.020107506	2.47E-04
Q8TF72	S1173	1.02690173	7.86E-06
P53367	T361	1.030749738	1.37E-05
Q8NG27	S265	1.040476507	9.36E-05
P30260	S364	1.048750617	3.85E-05
Q8N5C8	S645	1.049056194	9.50E-04
Q7L804	S388	1.090364347	1.88E-06
O94804	T952	1.091401297	2.53E-06
Q9UHI6	S500	1.101983264	2.09E-06
O14924	S172	1.113490681	4.74E-05
P23443	T444	1.11543865	8.48E-07
P35658	S430_S433	1.123403476	6.66E-07
Q02880	S1466_S1476	1.131573865	0.001148546
Q6ZRI6	T364	1.151053588	3.20E-07
P78536	T735	1.151662174	4.76E-09
Q9HCM7	S487	1.155552962	5.32E-05
Q09666	S5110	1.161055922	5.05E-05
Q6ZSZ5	T1345	1.18162819	5.50E-09
Q9NUL3	S440	1.199038899	2.53E-06
O14974	S507	1.199310364	2.83E-06

Q14680	T428	1.208454976	5.85E-08
Q7RTP6	RKTSQSEEEEEAP	1.209598439	4.24E-06
Q9Y6J0	T1927	1.213465802	4.64E-05
P30307	T48	1.229095945	2.56E-07
Q6Y7W6	S26_S30	1.241505956	5.01E-09
Q92995	S40	1.24511433	2.72E-05
Q8TF72	S1279	1.270024865	4.88E-05
Q92609	S558_T/S	1.294311834	7.57E-07
Q86SQ0	S415	1.311617773	9.80E-06
Q15154	S1283	1.320951525	5.59E-05
Q8IWC1	S524	1.337802706	3.14E-07
P49327	S831	1.362757505	1.14E-04
Q15418	S363_S369	1.387172446	3.23E-08
Q9UJY4	S400	1.396751421	2.24E-04
Q5SW79	S1160	1.419707148	6.33E-08
Q8TF72	S403	1.475842451	1.11E-06
O43295	S858	1.484343041	4.24E-06
Q15418	S732	1.528840505	1.08E-05
Q8TF72	S816	1.53438654	5.69E-08
Q9UQC2	S543	1.544255432	8.54E-08
Q7Z5J4	T1068	1.551286529	5.69E-08
P15056	S151	1.558659437	3.23E-08
P12270	S2155	1.581266963	7.02E-09
Q15418	S380	1.622483945	5.85E-08
P28482	T185	1.908205633	1.99E-08
P12270	T2116	1.935996532	1.13E-08
P27361	Y204	2.028335793	3.23E-08
Q9UJY4	S177	2.051775366	3.67E-06
Q9UDY2	S163	2.053682259	3.52E-07
Q9NZN5	T703	2.496942393	3.48E-04
P28482	Y187	2.826457168	7.02E-09
P27361	T202_Y204	3.452269656	2.60E-10
P28482	T185_Y187	3.653298586	2.60E-10