

Accession	Peptides	Score	Anova (p)*	Fold	Description	Average Normalised Abundances		
						Control	SDS_HaCaT	Triton_HaCaT
P25705	23 (21)	2997.10	0.02	2.79	ATP synthase subunit alpha, mitochondrial (ATPA_HUMAN)	1.20e+007	3.34E+07	2.42e+007
P36952	15	2000.00	3.65e-005	5.19	Serpin B5 (SPB5_HUMAN)	2.01E+07	1.47E+07	3.88e+006
P05120	10	1098.74	1.19e-004	5.53	Plasminogen activator inhibitor 2 (PAI2_HUMAN)	3.50e+006	1.16e+006	6.32e+005
P18621	6	699.14	1.25e-005	3.73	60S ribosomal protein L17 (RL17_HUMAN)	2.12e+006	1.56e+006	5.67e+005
O43776	5	500.00	1.96e-004	7.42	Asparagine--tRNA ligase, cytoplasmic (SYNC_HUMAN)	8.47e+005	6.29e+005	1.14e+005
Q96KP4	4	400.00	1.63e-003	3.76	Cytosolic non-specific dipeptidase (CNDP2_HUMAN)	1.38e+006	5.83e+005	3.67e+005
O14950	4 (3)	399.91	0.37	1.43	Myosin regulatory light chain 12B (ML12B_HUMAN)	1.58e+006	1.92e+006	1.35e+006
Q9P0L0	2	300.00	0.57	2.79	Vesicle-associated membrane protein-associated protein A (VAPA_HUMAN)	1.56E+05	4.37e+005	3.31E+05
Q9UIV8	3	300.00	5.19e-003	5.42	Serpin B13 (SPB13_HUMAN)	1.26e+006	6.26e+005	2.33e+005
P49257	3	299.95	0.70	2.42	Protein ERGIC-53 (LMAN1_HUMAN)	1.30e+005	2.38e+005	3.14e+005
Q14126	2	200.00	0.39	2.87	Desmoglein-2 (DSG2_HUMAN)	1.33e+005	1.12e+005	4.62e+004
O43396	2	200.00	0.22	1.86	Thioredoxin-like protein 1 (TXNL1_HUMAN)	1.38e+005	1.00e+005	7.42e+004
P32926	1	100.00	0.87	1.98	Desmoglein-3 (DSG3_HUMAN)	1.20e+005	6.07e+004	9.29e+004
Q02487	1	100.00	4.58e-003	59.88	Desmocollin-2 (DSC2_HUMAN)	2.86e+004	3.21e+004	536.01
Q15555	1	100.00	9.71e-005	7202	Microtubule-associated protein RP/EB family member 2 (MARE2_HUMAN)	6.35e+004	6276.12	8.82
P19404	1	100.00	0.31	5.41	NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial (NDUV2_HUMAN)	4.35e+004	2.35e+005	2.19e+005
P00167	1	100.00	0.11	1.27	Cytochrome b5 (CYB5_HUMAN)	2.68e+004	2.11e+004	2.23e+004
P62314	1	100.00	0.22	18.63	Small nuclear ribonucleoprotein Sm D1 (SMD1_HUMAN)	4.41e+004	8650.38	2367.15