

<b>Peak name</b>	<b>P value</b>	<b>Fold change</b>	<b>Fold change</b>
sp   Q9P0J7   KCMF1_HUMAN	0.02399	4.388536121	0.227866417
sp   O60669   MOT2_HUMAN	0.021	3.523680792	0.283794151
tr   K7ERI7   K7ERI7_HUMAN	0.04805	3.362868163	0.297365211
sp   O95071   UBR5_HUMAN	0.00746	3.358724506	0.29773207
sp   Q9NVT9   ARMC1_HUMAN	0.03904	3.233497399	0.309262658
sp   Q14012   KCC1A_HUMAN	0.03054	3.088629694	0.323768175
sp   O43395   PRPF3_HUMAN	0.03612	2.950485271	0.338927298
sp   Q969S3   ZN622_HUMAN	0.02318	2.939511779	0.340192547
tr   B3KTM8   B3KTM8_HUMAN	0.03706	2.802683281	0.35680093
sp   Q9BPX3   CND3_HUMAN	0.00807	2.68237663	0.372803725
sp   P14174   MIF_HUMAN	0.0231	2.681218833	0.372964708
sp   Q96GM8   TOE1_HUMAN	0.02589	2.590172969	0.386074603
tr   F8VUA2   F8VUA2_HUMAN	8.06E-05	2.439941074	0.409845963
sp   Q5R3I4   TTC38_HUMAN	0.00198	2.407488207	0.415370674
sp   Q12768   STRUM_HUMAN	0.0186	2.403012932	0.416144244
sp   Q8N3V7-2   SYNPO_HUMAN	0.00208	2.351134561	0.425326571
sp   P62942   FKB1A_HUMAN	0.04451	2.34978258	0.425571288
sp   Q9UPY8   MARE3_HUMAN	0.03397	2.27840259	0.438903995
sp   Q9ULX3   NOB1_HUMAN	0.00065	2.260527698	0.442374584
sp   O60645   EXOC3_HUMAN	0.03527	2.248761678	0.444689186
sp   Q8N8A6   DDX51_HUMAN	0.02855	2.193398716	0.455913461
sp   Q71RC2   LARP4_HUMAN	0.03038	2.101897854	0.475760512
sp   O75494   SRS10_HUMAN	0.00546	2.069175785	0.483284217
sp   Q8NI36   WDR36_HUMAN	0.02345	2.06035618	0.485352974
sp   Q9H8H2   DDX31_HUMAN	0.03107	2.043938129	0.4892516
sp   Q96FJ2   DYL2_HUMAN	0.00371	2.03326856	0.491818946
sp   Q16537   2A5E_HUMAN	0.00644	2.030602864	0.492464587
sp   Q92615   LAR4B_HUMAN	0.02111	1.998889333	0.500277821

sp   P02751   FINC_HUMAN	0.02949	1.99625418	0.500938212
sp   Q9Y333   LSM2_HUMAN	0.02813	1.993944575	0.501518454
sp   Q6RW13-2   ATRAP_HUMAN	0.02376	1.95968704	0.510285561
sp   Q9GZT8   GTPC1_HUMAN	0.00057	1.959621852	0.510302536
sp   Q9H2P9-5   DPH5_HUMAN	0.00015	1.922039035	0.520280796
sp   Q9NPD3   EXOS4_HUMAN	0.04064	1.91376698	0.522529655
sp   P25685   DNJB1_HUMAN	0.00723	1.913635232	0.52256563
sp   P00325   ADH1B_HUMAN	0.02254	1.913122946	0.52270556
tr   A0A0A0MRM9   A0A0A0MRM9_HUMAN	0.01033	1.910266287	0.523487226
tr   R4GMX8   R4GMX8_HUMAN	0.03129	1.903923129	0.525231289
tr   A0A087WZV0   A0A087WZV0_HUMAN	0.02244	1.902836752	0.525531157
sp   Q96AE4   FUBP1_HUMAN	0.04956	1.901539901	0.52588957
sp   A5YKK6   CNOT1_HUMAN	0.02078	1.90006793	0.526296973
sp   P48556   PSMD8_HUMAN	0.01607	1.894492083	0.527845964
sp   Q96CP2   FWCH2_HUMAN	0.00341	1.89185461	0.528581845
sp   Q9H3Q1   BORG4_HUMAN	0.00343	1.881061621	0.531614695
sp   Q9H1B7   I2BPL_HUMAN	0.02233	1.861858917	0.537097624
sp   Q96GX2   A7L3B_HUMAN	0.02183	1.856150068	0.538749543
sp   Q9BT73   PSMG3_HUMAN	0.04747	1.829301431	0.546656764
sp   Q9BY43-2   CHM4A_HUMAN	0.03085	1.815739473	0.550739803
sp   Q00796   DHSO_HUMAN	4.45E-05	1.813294208	0.551482487
tr   Q4R9M9   Q4R9M9_HUMAN	0.03739	1.808972334	0.552800052
sp   P05412   JUN_HUMAN	0.00124	1.80342987	0.554498967
sp   P05387   RLA2_HUMAN	0.01095	1.800733614	0.555329224
sp   Q8TF05   PP4R1_HUMAN	0.01869	1.795963102	0.556804312
sp   O95487   SC24B_HUMAN	0.02136	1.794041763	0.557400625
sp   O60443   DFNA5_HUMAN	0.00031	1.785855186	0.559955817
sp   Q96KP1   EXOC2_HUMAN	0.02155	1.784849415	0.560271355
tr   G3V599   G3V599_HUMAN	0.00056	1.782562413	0.560990175

sp   Q16222   UAP1_HUMAN	0.00407	1.778208832	0.562363645
sp   Q99685   MGLL_HUMAN	0.00319	1.771246546	0.564574143
tr   U3KQP1   U3KQP1_HUMAN	0.00499	1.761006412	0.567857103
sp   Q96T76   MMS19_HUMAN	0.04308	1.752137233	0.570731551
sp   P42226   STAT6_HUMAN	0.04649	1.752103481	0.570742545
sp   O95336   6PGL_HUMAN	0.00294	1.7263613	0.579253022
sp   O15294   OGT1_HUMAN	0.00231	1.718489145	0.581906498
tr   H0Y7L6   H0Y7L6_HUMAN	0.03111	1.711638521	0.584235508
sp   O75683   SURF6_HUMAN	0.02606	1.711502989	0.584281773
tr   A6NMQ3   A6NMQ3_HUMAN	0.03923	1.710805804	0.584519878
sp   O15084   ANR28_HUMAN	0.04928	1.709161616	0.585082178
sp   Q8NEU8   DP13B_HUMAN	0.04233	1.709016327	0.585131917
sp   Q9HA64   KT3K_HUMAN	0.02993	1.706945382	0.585841826
tr   H6UYS7   H6UYS7_HUMAN	0.04904	1.70321122	0.58712624
sp   Q8IVF2-3   AHNK2_HUMAN	0.03209	1.693766952	0.590399995
sp   A6NDU8   CE051_HUMAN	0.04084	1.692265967	0.590923661
sp   Q96DG6   CMBL_HUMAN	0.00816	1.685490672	0.593299041
sp   P22392-2   NDKB_HUMAN	0.00353	1.679625083	0.595370961
sp   Q8WTS6   SETD7_HUMAN	0.03867	1.677399543	0.596160887
sp   Q9GZL7   WDR12_HUMAN	0.00057	1.666038353	0.600226278
sp   P08397   HEM3_HUMAN	1.48E-05	1.663136018	0.601273732
sp   Q9H223   EHD4_HUMAN	0.01285	1.661111146	0.602006676
sp   Q6BCY4   NB5R2_HUMAN	0.03523	1.654225507	0.604512502
sp   Q53FT3   HIKES_HUMAN	0.02947	1.652138985	0.605275954
tr   E9PD53   E9PD53_HUMAN	0.04452	1.648468064	0.606623823
sp   Q15102   PA1B3_HUMAN	0.004	1.644249025	0.608180382
sp   A6NHR9   SMHD1_HUMAN	0.02317	1.636666033	0.6109982
sp   Q7Z4W1   DCXR_HUMAN	0.01592	1.635054212	0.611600516
tr   F8VXU5   F8VXU5_HUMAN	0.01754	1.628447472	0.614081828

sp   Q8NBT2   SPC24_HUMAN	0.01425	1.627679895	0.614371415
sp   Q05048   CSTF1_HUMAN	0.03528	1.625929239	0.615032915
sp   Q9NY27   PP4R2_HUMAN	0.01337	1.624298876	0.615650245
tr   C9JBI3   C9JBI3_HUMAN	0.0177	1.620756753	0.616995732
sp   Q13085-4   ACACA_HUMAN	0.01491	1.620649523	0.617036556
sp   P19338   NUCL_HUMAN	0.04769	1.620349867	0.617150666
sp   Q9H3S7   PTN23_HUMAN	1.12E-05	1.616741215	0.618528179
sp   P29401   TKT_HUMAN	0.0089	1.612648346	0.620097991
sp   O95182   NDUA7_HUMAN	0.04562	1.61128946	0.620620953
sp   P36639   8ODP_HUMAN	0.00637	1.609876187	0.621165782
tr   A8MT02   A8MT02_HUMAN	0.00757	1.589947838	0.628951451
tr   S4R3N1   S4R3N1_HUMAN	0.03045	1.589881234	0.628977799
sp   P40925-3   MDHC_HUMAN	0.02353	1.577216483	0.634028373
sp   P16930   FAAA_HUMAN	5.47E-06	1.571837022	0.636198274
sp   Q13126   MTAP_HUMAN	0.02959	1.556710279	0.64238029
sp   P18669   PGAM1_HUMAN	0.03456	1.542737365	0.64819847
tr   E5RG17   E5RG17_HUMAN	0.03073	1.537448424	0.650428323
sp   Q96GS4   CQ059_HUMAN	0.02054	1.536210534	0.650952443
sp   Q9H3K6   BOLA2_HUMAN	0.01752	1.535997829	0.651042587
tr   H0YEB6   H0YEB6_HUMAN	0.00371	1.534676055	0.651603312
sp   Q8WXD5   GEMI6_HUMAN	0.0107	1.532711596	0.652438464
tr   E9PS17   E9PS17_HUMAN	0.02648	1.530021627	0.653585533
sp   Q9BPX5   ARP5L_HUMAN	0.0251	1.524814756	0.655817368
sp   Q15654   TRIP6_HUMAN	0.01349	1.521377446	0.657299083
sp   O75223   GGCT_HUMAN	0.01663	1.515503656	0.659846643
tr   A0A0A0MTG4   A0A0A0MTG4_HUMAN	0.00139	1.514071464	0.660470806
sp   Q8NC51   PAIRB_HUMAN	0.01795	1.509497696	0.662472028
sp   O00629   IMA3_HUMAN	0.03877	0.664178869	1.505618512
sp   A6NIH7   U119B_HUMAN	0.03176	0.66325627	1.507712847

sp   Q9HCE5   MET14_HUMAN	0.01047	0.662702962	1.508971678
sp   Q8WVV9-5   HNRLH_HUMAN	0.01456	0.661801307	1.511027539
sp   Q9UIQ6   LCAP_HUMAN	0.00541	0.659364917	1.51661087
tr   E5RHW4   E5RHW4_HUMAN	0.0056	0.657058378	1.521934782
sp   O43896   KIF1C_HUMAN	0.00341	0.656736289	1.522681199
tr   A0A087WXS7   A0A087WXS7_HUMAN	0.02628	0.656410568	1.523436777
sp   Q03001   DYST_HUMAN	0.01681	0.650597097	1.537049587
sp   Q5JPE7   NOMO2_HUMAN	0.03207	0.650008346	1.538441786
sp   Q96S59   RANB9_HUMAN	0.02831	0.648961039	1.540924554
sp   Q96A33   CCD47_HUMAN	0.00329	0.648792552	1.541324723
sp   P07954-2   FUMH_HUMAN	0.00274	0.646788726	1.546099923
sp   P36542   ATPG_HUMAN	0.00663	0.64626732	1.547347311
sp   Q08752   PPID_HUMAN	9.69E-05	0.642985791	1.55524432
sp   P11279   LAMP1_HUMAN	0.01338	0.641165624	1.559659411
sp   P11047   LAMC1_HUMAN	1.51E-05	0.63864071	1.565825641
sp   Q4G0N4   NAKD2_HUMAN	0.04539	0.637530704	1.568551905
sp   P38646   GRP75_HUMAN	0.00183	0.636186659	1.57186572
sp   Q13423   NNTM_HUMAN	0.00024	0.635717299	1.573026251
sp   Q15084-5   PDIA6_HUMAN	8.56E-05	0.634701338	1.575544181
tr   A0A087X0S5   A0A087X0S5_HUMAN	0.04424	0.633816254	1.577744329
tr   D6RF48   D6RF48_HUMAN	0.00095	0.631885129	1.582566124
ENSEMBL:ENSBTAP00000018574	0.01318	0.631237019	1.584190994
sp   P14854   CX6B1_HUMAN	0.0332	0.630431984	1.586213938
sp   P08962   CD63_HUMAN	0.01134	0.629781303	1.587852791
sp   P30084   ECHM_HUMAN	0.00846	0.627300127	1.594133265
sp   Q6L8Q7   PDE12_HUMAN	0.0256	0.626890969	1.595173722
sp   P24390   ERD21_HUMAN	0.00272	0.625095194	1.599756341
sp   Q9NX14   NDUBB_HUMAN	0.00366	0.622394315	1.606698479
sp   O95302   FKBP9_HUMAN	0.02505	0.620705501	1.611069983

tr   A0A087WU53   A0A087WU53_HUMAN	0.0468	0.620294325	1.612137915
sp   P21796   VDAC1_HUMAN	0.03887	0.619748981	1.613556506
tr   V9GZ56   V9GZ56_HUMAN	0.03473	0.618291217	1.617360837
sp   Q8ND56   LS14A_HUMAN	0.00469	0.61443111	1.627521758
sp   Q96I99   SUCB2_HUMAN	0.00225	0.613511061	1.629962464
sp   Q6GMV2   SMYD5_HUMAN	0.00765	0.613030655	1.631239796
sp   P55084   ECHB_HUMAN	0.01036	0.612446385	1.632795986
sp   P53007   TXTP_HUMAN	0.0076	0.611008545	1.636638321
sp   Q9UJZ1   STML2_HUMAN	0.03788	0.609016471	1.641991717
sp   P15559   NQO1_HUMAN	0.03688	0.608471445	1.643462496
sp   O75306   NDUS2_HUMAN	0.03571	0.608118329	1.644416806
sp   Q08722-4   CD47_HUMAN	0.01717	0.607846112	1.645153239
sp   Q96PY5   FMNL2_HUMAN	0.03997	0.596234998	1.677191046
sp   P25705   ATPA_HUMAN	0.01855	0.595076844	1.68045524
tr   F5GX77   F5GX77_HUMAN	8.81E-05	0.592801985	1.686903933
sp   P43235   CATK_HUMAN	0.03658	0.592756518	1.687033326
tr   E9PC15   E9PC15_HUMAN	0.0008	0.591497944	1.690622951
sp   Q9H9B4   SFXN1_HUMAN	0.03331	0.589382506	1.69669101
sp   Q969V3   NCLN_HUMAN	0.00264	0.584833142	1.709889418
sp   P46977   STT3A_HUMAN	0.02123	0.583086453	1.715011546
sp   P35908   K22E_HUMAN	9.22E-08	0.582843616	1.715726094
sp   P39656   OST48_HUMAN	0.0225	0.581381997	1.7200395
sp   Q12981   SEC20_HUMAN	0.026	0.579021224	1.72705241
tr   A0A087X0W7   A0A087X0W7_HUMAN	0.00341	0.578093076	1.72982525
tr   A2A274   A2A274_HUMAN	0.025	0.578006144	1.730085416
sp   P24844   MYL9_HUMAN	0.00403	0.577445931	1.73176387
tr   M0R0R2   M0R0R2_HUMAN	0.00811	0.574044514	1.742025184
sp   P45877   PPIC_HUMAN	0.00908	0.573035047	1.745093962
sp   Q9UBP6   TRMB_HUMAN	0.00012	0.5720933	1.747966635

sp   Q9H0K6   PUS7L_HUMAN	0.01479	0.570477957	1.752916107
sp   P08559   ODP A_HUMAN	0.02116	0.565939539	1.766973203
sp   P07355-2   ANXA2_HUMAN	0.0116	0.562550626	1.777617789
sp   P05141   ADT2_HUMAN	0.00107	0.559447266	1.787478573
P00735	5.33E-08	0.555501034	1.800176666
tr   M0R0A9   M0R0A9_HUMAN	0.02629	0.555426974	1.800416701
sp   P24752   THIL_HUMAN	0.00187	0.555398254	1.8005098
sp   Q9UHQ9   NB5R1_HUMAN	0.01263	0.553744794	1.805886053
sp   Q96N67   DOCK7_HUMAN	0.01396	0.550472462	1.816621301
sp   P26038   MOES_HUMAN	0.00944	0.548704199	1.822475573
sp   P04114   APOB_HUMAN	0.01367	0.547966709	1.824928383
sp   Q8TCT8   SPP2A_HUMAN	0.0308	0.54770958	1.825785117
sp   Q9NTJ5   SAC1_HUMAN	0.03282	0.545772688	1.832264643
tr   E9PKP7   E9PKP7_HUMAN	0.01882	0.537451564	1.860632784
Q3MHN5	0.01275	0.535840529	1.866226882
sp   Q8NDI1-3   EHBP1_HUMAN	0.01967	0.53502268	1.869079643
P41361	0.00272	0.532033151	1.879582124
P00978	7.67E-05	0.527271941	1.896554554
Q3SZR3	0.02345	0.526914705	1.897840371
sp   P16401   H15_HUMAN	0.02761	0.522909521	1.912376729
sp   Q9BQC6   RT63_HUMAN	0.02701	0.522728784	1.913037946
sp   Q3SY69   AL1L2_HUMAN	0.01389	0.521998644	1.915713788
Q2KJF1	0.003	0.521286364	1.918331398
sp   P53597   SUCA_HUMAN	0.00185	0.520961219	1.91952868
P17690	0.00419	0.520288454	1.922010745
sp   Q9UBB5   MBD2_HUMAN	0.01255	0.519002343	1.926773576
sp   Q9C0J8   WDR33_HUMAN	0.0006	0.512647787	1.950657011
sp   Q2TAA5   ALG11_HUMAN	0.00948	0.511038104	1.956801249
sp   P67812   SC11A_HUMAN	6.31E-05	0.510836953	1.957571774

sp   P08727   K1C19_HUMAN	0.02119	0.509528747	1.962597806
sp   P45880   VDAC2_HUMAN	0.0013	0.506703426	1.973541027
sp   Q15165-1   PON2_HUMAN	0.01949	0.506023263	1.976193732
Q03247	0.00301	0.504728713	1.981262356
sp   O75380   NDUS6_HUMAN	1.00E-05	0.503553671	1.985885632
sp   P28331   NDUS1_HUMAN	0.00402	0.50191666	1.992362637
sp   Q8N6L1   KTAP2_HUMAN	0.01745	0.497081424	2.01174285
P34955	0.02898	0.493291526	2.027198822
tr   H0Y892   H0Y892_HUMAN	0.01778	0.491393167	2.035030333
Q3SX14	0.0176	0.491075762	2.036345668
Q0VCM5	0.00436	0.481894531	2.075142871
Q28065	0.00254	0.480786534	2.079925143
sp   P24821   TENA_HUMAN	0.01746	0.479075638	2.087353063
sp   P05120   PAI2_HUMAN	0.03793	0.474769693	2.106284406
tr   A0A087WUB9   A0A087WUB9_HUMAN	0.00092	0.474327941	2.108246034
Q3SZ57	0.00098	0.471832943	2.119394194
sp   Q00325-2   MPCP_HUMAN	0.00277	0.470543288	2.125202984
Q32PI4	0.00253	0.469362028	2.130551557
Q3KUS7	0.00092	0.459701152	2.175326289
sp   Q9NVE7   PANK4_HUMAN	0.00286	0.456552438	2.190328901
P81644	0.01803	0.455422507	2.195763242
Q9TTE1	0.0014	0.453202202	2.20652061
Q0V8M9	0.00099	0.451496806	2.214855094
sp   Q9H845   ACAD9_HUMAN	0.01246	0.447432945	2.234971767
sp   Q9UMY1   NOL7_HUMAN	0.02271	0.43831339	2.281472622
sp   P02462   CO4A1_HUMAN	0.00903	0.436694878	2.289928394
sp   P08574   CY1_HUMAN	0.02782	0.432996311	2.309488496
sp   Q9H788   SH24A_HUMAN	0.02692	0.431085013	2.319728058
P28800	0.00043	0.430015484	2.325497653

sp   O95168   NDUB4_HUMAN	3.46E-05	0.429005902	2.330970262
Q2UVX4	1.08E-06	0.428157289	2.335590274
sp   Q96RN5   MED15_HUMAN	0.02082	0.428100913	2.335897847
sp   Q9BVG9   PTSS2_HUMAN	0.00045	0.425440613	2.350504323
ENSEMBL:ENSBTAP00000024466	0.01333	0.424295546	2.356847743
tr   Q5T4U5   Q5T4U5_HUMAN	0.01332	0.410495177	2.436082213
sp   Q9H267   VP33B_HUMAN	0.02976	0.402444929	2.484812027
tr   A0A087WTW0   A0A087WTW0_HUMAN	0.01835	0.401516902	2.490555178
sp   Q86WJ1   CHD1L_HUMAN	0.00255	0.399378775	2.503888695
sp   Q16832   DDR2_HUMAN	0.00683	0.39557676	2.527954373
sp   Q00059   TFAM_HUMAN	0.00051	0.39150534	2.554243578
sp   P09669   COX6C_HUMAN	0.00122	0.389072591	2.570214461
tr   J3KQ18   J3KQ18_HUMAN	0.00994	0.377188561	2.65119387
sp   P05997   CO5A2_HUMAN	0.01949	0.372873456	2.681875003
tr   A0A087X0M4   A0A087X0M4_HUMAN	0.00964	0.365989419	2.732319428
sp   Q9BSJ2   GCP2_HUMAN	0.00808	0.365667079	2.734728003
Q3T052	0.0007	0.365160797	2.7385196
sp   Q96G21   IMP4_HUMAN	0.04255	0.347603996	2.876836891
sp   Q13425   SNTB2_HUMAN	0.02182	0.338907122	2.950660917
P15497	0.00043	0.323340598	3.092714015
sp   P47914   RL29_HUMAN	0.00053	0.313375544	3.191059479
Q58D62	6.20E-06	0.281597996	3.551161635
sp   Q53T59   H1BP3_HUMAN	0.00028	0.273547702	3.655669528
sp   Q99570   PI3R4_HUMAN	2.11E-05	0.262776216	3.805519441
P06868	0.00164	0.26100822	3.831296966
sp   Q10570   CPSF1_HUMAN	3.88E-05	0.257586777	3.882186862
sp   Q92542   NICA_HUMAN	0.04992	0.210466584	4.751348088

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