

Supplement Table 1. Identified secreted proteins from Alexander, HepG2, SK-Hep-1 and HCC-S102 cell lines

| Cell lines | Bands | Accession numbers | Names | Score | Peptide matches | Functions |
|------------|------------|---------------------------------|--|-------|-----------------|-----------------------------------|
| Alexander | 1 | TTBK1_HUMAN | Tau-tubulin kinase 1 | 46 | 1 | Metabolism |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 71 | 7 | Protein synthesis and degradation |
| | | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 39 | 2 | Cytoskeleton/mobility |
| | 2 | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 80 | 5 | Cytoskeleton/mobility |
| | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 76 | 8 | Cytoskeleton/mobility |
| | | ALBU_HUMAN | Human serum albumin | 70 | 3 | Transport/binding proteins |
| | 3 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 95 | 6 | Cytoskeleton/mobility |
| | | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 89 | 5 | Cytoskeleton/mobility |
| | | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 39 | 3 | Cytoskeleton/mobility |
| | | RAE1_HUMAN | Rab proteins geranylgeranyltransferase component A 1 | 35 | 1 | Transport/binding proteins |
| | | ALBU_HUMAN | Human serum albumin | 37 | 2 | Transport/binding proteins |
| | 4 | ALBU_HUMAN | Human serum albumin | 37 | 2 | Transport/binding proteins |
| | | FINC_HUMAN | Fibronectin | 36 | 2 | Cytoskeleton/mobility |
| | | CNGB3_HUMAN | Cyclic nucleotide-gated cation channel beta-3 | 36 | 1 | Ion channels |
| | | CCL28_HUMAN | C-C motif chemokine 28 | 35 | 1 | Immunological response |
| | 5 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 76 | 4 | Cytoskeleton/mobility |
| | | K2C1B_HUMAN | Keratin, type II cytoskeletal 1b | 45 | 2 | Cytoskeleton/mobility |
| | | LOXH1_HUMAN | Lipoxygenase homology domain-containing protein 1 | 35 | 1 | Transport/binding proteins |
| | | ALBU_HUMAN | Human serum albumin | 38 | 2 | Transport/binding proteins |
| | 6 | ALBU_HUMAN | Human serum albumin | 38 | 3 | Transport/binding proteins |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 56 | 9 | Protein synthesis and degradation |
| | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 91 | 1 | Cytoskeleton/mobility |
| | | LG3BP_HUMAN | Galectin-3-binding protein | 55 | 4 | Transport/binding proteins |
| | 7 | MYOZ2_HUMAN | Myozenin-2 | 39 | 1 | Transport/binding proteins |
| | | BGAT_HUMAN | Histo-blood group ABO system transferase | 38 | 1 | Metabolism |
| | | ALBU_HUMAN | Human serum albumin | 36 | 2 | Transport/binding proteins |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 37 | 1 | Protein synthesis and degradation |
| | 8 | CERU_HUMAN | Ceruloplasmin | 40 | 2 | Transport/binding proteins |
| | | SNX32_HUMAN | Sorting nexin-32 | 37 | 1 | Transport/binding proteins |
| | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 44 | 2 | Cytoskeleton/mobility |
| | 9 | CNTN1_HUMAN | Contactin-1 | 56 | 3 | Signal transduction |
| | | NRCAM_HUMAN | Neuronal cell adhesion molecule | 55 | 4 | Transport/binding proteins |
| | | ALBU_HUMAN | Human serum albumin | 37 | 2 | Transport/binding proteins |
| 10 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 182 | 8 | | Cytoskeleton/mobility |

| | | | | | |
|----|--------------|--|-----|---|-----------------------------------|
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 100 | 5 | Cytoskeleton/mobility |
| | LG3BP_HUMAN | Galectin-3-binding protein | 42 | 4 | Transport/binding proteins |
| | CERU_HUMAN | Ceruloplasmin | 38 | 1 | Transport/binding proteins |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 61 | 3 | Cytoskeleton/mobility |
| 11 | Q2A130_HUMAN | Autogenous vein graft remodeling associated protein 5 | 39 | 1 | DNA replication/gene regulation |
| | LG3BP_HUMAN | Galectin-3-binding protein | 41 | 2 | Transport/binding proteins |
| | ANKAR_HUMAN | Ankyrin and armadillo repeat-containing protein | 37 | 2 | Transport/binding proteins |
| 12 | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 67 | 4 | Cytoskeleton/mobility |
| | LG3BP_HUMAN | Galectin-3-binding protein | 120 | 8 | Transport/binding proteins |
| 13 | Q13876_HUMAN | Bone-derived growth factor | 43 | 2 | Metabolism |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 38 | 2 | Cytoskeleton/mobility |
| | LG3BP_HUMAN | Galectin-3-binding protein | 44 | 3 | Transport/binding proteins |
| | gi_119628058 | Mitochondrial trans-2-enoyl-CoA reductase, isoform CRA_a | 37 | 1 | Metabolism |
| | LOXH1_HUMAN | Lipoxygenase homology domain-containing protein 1 | 35 | 2 | Transport/binding proteins |
| 14 | TRFE_HUMAN | Serotransferrin | 70 | 5 | Transport/binding proteins |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 40 | 4 | Cytoskeleton/mobility |
| | RRAGB_HUMAN | Ras-related GTP-binding protein B | 35 | 1 | Signal transduction |
| | CSN1_HUMAN | COP9 signalosome complex subunit 1 | 35 | 1 | Cell cycle |
| 15 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 47 | 3 | Cytoskeleton/mobility |
| | K2C1B_HUMAN | Keratin, type II cytoskeletal 1b | 43 | 2 | Cytoskeleton/mobility |
| | ERFI_HUMAN | Eukaryotic peptide chain release factor subunit 1 | 36 | 1 | Protein synthesis and degradation |
| | TRFE_HUMAN | Serotransferrin | 45 | 3 | Transport/binding proteins |
| | RGS14_HUMAN | Regulator of G-protein signaling 14 | 39 | 1 | Signal transduction |
| 16 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 39 | 3 | Cytoskeleton/mobility |
| | SNX32_HUMAN | Sorting nexin-32 | 37 | 1 | Transport/binding proteins |
| | gi_119583243 | hCG1784572 | 39 | 1 | Unknown |
| | MA2C1_HUMAN | Alpha-mannosidase 2C1 | 35 | 1 | Metabolism |
| 17 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 46 | 3 | Cytoskeleton/mobility |
| | Z780B_HUMAN | Zinc finger protein 780B | 39 | 1 | DNA replication/gene regulation |
| | IN80D_HUMAN | INO80 complex subunit D | 42 | 1 | DNA replication/gene regulation |
| | CNGB3_HUMAN | Cyclic nucleotide-gated cation channel beta-3 | 36 | 1 | Ion channels |
| | B3GT6_HUMAN | Beta-1,3-galactosyltransferase 6 | 38 | 1 | Metabolism |
| 18 | A1AT_HUMAN | Alpha-1-antitrypsin | 63 | 2 | Protein synthesis and degradation |
| | ANGT_HUMAN | Angiotensinogen | 112 | 5 | Signal Transduction |
| | BGH3_HUMAN | Transforming growth factor-beta-induced protein ig-h3 | 37 | 2 | Transport/binding proteins |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 36 | 2 | Cytoskeleton/mobility |
| 19 | AMBP_HUMAN | Protein AMBP | 46 | 1 | Protein synthesis and degradation |
| | A1AT_HUMAN | Alpha-1-antitrypsin | 90 | 4 | Protein synthesis and degradation |

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|----|--------------|---|-----|----|-----------------------------------|--|
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| 20 | AMBP_HUMAN | Protein AMBP | 46 | 1 | Protein synthesis and degradation | |
| | A1AT_HUMAN | Alpha-1-antitrypsin | 169 | 8 | Protein synthesis and degradation | |
| | HERC6_HUMAN | Probable E3 ubiquitin-protein ligase HERC6 | 42 | 1 | Protein synthesis and degradation | |
| | gi 119609527 | hCG1641439 | 38 | 1 | Unknown | |
| 21 | A1AT_HUMAN | Alpha-1-antitrypsin | 141 | 8 | Protein synthesis and degradation | |
| | NSUN4_HUMAN | Putative methyltransferase NSUN4 | 48 | 1 | Methyltransferase activity | |
| | AMBP_HUMAN | Protein AMBP | 46 | 1 | Cell adhesion | |
| | ANGT_HUMAN | Angiotensinogen | 54 | 4 | Signal Transduction | |
| 22 | IDH3A_HUMAN | Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial | 39 | 1 | Metabolism | |
| | A1AT_HUMAN | Alpha-1-antitrypsin | 231 | 8 | Protein synthesis and degradation | |
| 23 | IDH3A_HUMAN | Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial | 39 | 1 | Metabolism | |
| | AMBP_HUMAN | Protein AMBP | 58 | 3 | Protein synthesis and degradation | |
| | FIBB_HUMAN | Fibrinogen beta chain | 107 | 3 | Protein synthesis and degradation | |
| 24 | ENOA_HUMAN | Alpha-enolase | 81 | 4 | Metabolism | |
| | AMBP_HUMAN | Protein AMBP | 46 | 1 | Protein synthesis and degradation | |
| 25 | CBPN_HUMAN | Carboxypeptidase N catalytic chain | 41 | 1 | Protein synthesis and degradation | |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 38 | 1 | Cytoskeleton/mobility | |
| | DAG1_HUMAN | Dystroglycan | 36 | 1 | Cytoskeleton/mobility | |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 45 | 2 | Cytoskeleton/mobility | |
| 26 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 49 | 5 | Cytoskeleton/mobility | |
| | AATC_HUMAN | Aspartate aminotransferase, cytoplasmic | 45 | 3 | Metabolism | |
| | gi 28336 | Mutant beta-actin | 43 | 4 | Cytoskeleton/mobility | |
| | AMBP_HUMAN | Protein AMBP | 54 | 2 | Protein synthesis and degradation | |
| | APOE_HUMAN | Apolipoprotein E | 39 | 1 | Transport/binding proteins | |
| 27 | K1C18_HUMAN | Keratin, type I cytoskeletal 18 | 45 | 2 | Cytoskeleton/mobility | |
| | RIF1_HUMAN | Telomere-associated protein RIF1 | 35 | 2 | Cell cycle | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 65 | 5 | Cytoskeleton/mobility | |
| | DYH11_HUMAN | Dynein heavy chain 11, axonemal | 39 | 3 | Cytoskeleton/mobility | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 237 | 10 | Cytoskeleton/mobility | |
| 28 | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 122 | 8 | Cytoskeleton/mobility | |
| | APOE_HUMAN | Apolipoprotein E | 46 | 2 | Transport/binding proteins | |
| | AMBP_HUMAN | Protein AMBP | 46 | 1 | Protein synthesis and degradation | |
| 29 | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 36 | 1 | Cytoskeleton/mobility | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 68 | 5 | Cytoskeleton/mobility | |
| | APOE_HUMAN | Apolipoprotein E | 38 | 2 | Transport/binding proteins | |
| 29 | AMBP_HUMAN | Protein AMBP | 52 | 1 | Protein synthesis and degradation | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 165 | 8 | Cytoskeleton/mobility | |
| | K1C9_HUMAN | Keratin type, I cytoskeletal 9 | 61 | 4 | Cytoskeleton/mobility | |

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|----|--------------|---|-----|----|-----------------------------------|
| | K1C10_HUMAN | Keratin type, I cytoskeletal 10 | 81 | 8 | Cytoskeleton/mobility |
| | APOE_HUMAN | Apolipoprotein E | 144 | 4 | Transport/binding proteins |
| 30 | AMBP_HUMAN | Protein AMBP | 108 | 3 | Protein synthesis and degradation |
| | K1C18_HUMAN | Keratin, type I cytoskeletal 18 | 40 | 4 | Cytoskeleton/mobility |
| | G3P_HUMAN | Glyceraldehyde-3-phosphate dehydrogenase | 39 | 1 | Metabolism |
| | AMBP_HUMAN | Protein AMBP | 78 | 3 | Protein synthesis and degradation |
| 31 | ANXA1_HUMAN | Annexin A1 | 48 | 1 | Signal transduction |
| | CLUS_HUMAN | Clusterin | 58 | 3 | Transport/binding proteins |
| | 1433E_HUMAN | 14-3-3 protein epsilon | 39 | 1 | Signal transduction |
| | K2C8_HUMAN | Keratin, type II cytoskeletal 8 | 48 | 3 | Cytoskeleton/mobility |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 70 | 8 | Cytoskeleton/mobility |
| 32 | AMBP_HUMAN | Protein AMBP | 78 | 3 | Protein synthesis and degradation |
| | K2C8_HUMAN | Keratin, type II cytoskeletal 8 | 39 | 1 | Cytoskeleton/mobility |
| | P12L1_HUMAN | POM121-like protein 1 | 40 | 1 | Unknown |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 64 | 4 | Cytoskeleton/mobility |
| 33 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 50 | 5 | Cytoskeleton/mobility |
| | gi 119581427 | hCG1744108 | 47 | 1 | Unknown |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 172 | 10 | Cytoskeleton/mobility |
| 34 | TPIS_HUMAN | Triosephosphate isomerase | 92 | 3 | Metabolism |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 120 | 10 | Cytoskeleton/mobility |
| | BMS1_HUMAN | Ribosome biogenesis protein BMS1 homolog | 35 | 1 | Protein synthesis and degradation |
| | Q59F02_HUMAN | Phosphomannomutase 2 variant | 39 | 1 | Metabolism |
| | K22E_HUMAN | Keratin, type II cytoskeletal 2 epidermal | 159 | 6 | Cytoskeleton/mobility |
| | K2C8_HUMAN | Keratin, type II cytoskeletal 8 | 45 | 2 | Cytoskeleton/mobility |
| | AMBP_HUMAN | Protein AMBP | 36 | 1 | Protein synthesis and degradation |
| | APOA1_HUMAN | Apolipoprotein A-I | 57 | 5 | Transport/binding proteins |
| 35 | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 61 | 4 | Cytoskeleton/mobility |
| | APOE_HUMAN | Apolipoprotein E | 57 | 2 | Transport/binding proteins |
| | PSA2_HUMAN | Proteasome subunit alpha type-2 | 40 | 1 | Protein synthesis and degradation |
| | GSTP1_HUMAN | Glutathione S-transferase P | 95 | 3 | Metabolism |
| 36 | gi 119615628 | hCG17324, isoform CRA_b | 36 | 1 | Unknown |
| | gi 27368977 | Immunoglobulin kappa light chain variable region | 37 | 1 | Immunological response |
| | gi 119568507 | Chromosome 6 open reading frame 173, isoform CRA_c | 37 | 1 | DNA replication/gene regulation |
| | gi 129561187 | Alpha-helix coiled-coil rod homologue | 35 | 1 | DNA replication/gene regulation |
| | PEX14_HUMAN | Peroxisomal membrane protein PEX14 | 41 | 1 | Transport/binding proteins |
| 37 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 89 | 6 | Cytoskeleton/mobility |
| | K2C1B_HUMAN | Keratin, type II cytoskeletal 1b | 40 | 2 | Cytoskeleton/mobility |
| | A2A391_HUMAN | Obscurin, cytoskeletal calmodulin and titin-interacting | 35 | 2 | Metabolism |

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|----|--------------|---|----|---|------------------------------------|
| | | RhoGEF | | | |
| | WWP2_HUMAN | NEDD4-like E3 ubiquitin-protein ligase WWP2 | 35 | 1 | Protein synthesis and degradation |
| | MYO9A_HUMAN | Myosin-IXa | 35 | 1 | Signal transduction |
| 39 | PPIC_HUMAN | Peptidyl-prolyl cis-trans isomerase C | 41 | 1 | Signal transduction |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 38 | 1 | Cytoskeleton/mobility |
| | gi 119583243 | hCG1784572 | 37 | 1 | Unknown |
| | gi 10637712 | Immunoglobulin lambda light chain variable region | 35 | 1 | Immunological response |
| 40 | TBC17_HUMAN | TBC1 domain family member 17 | 38 | 1 | Signal transduction |
| | PRDX1_HUMAN | Peroxiredoxin-1 | 58 | 2 | Protection and detoxification |
| | BBS7_HUMAN | Bardet-Biedl syndrome 7 protein | 35 | 1 | Transport/binding proteins |
| 41 | S14L2_HUMAN | SEC14-like protein 2 | 26 | 2 | DNA replication/gene regulation |
| | FAM3C_HUMAN | Protein FAM3C | 45 | 1 | Signal transduction |
| | GGCT_HUMAN | Gamma-glutamyl cyclotransferase | 38 | 1 | Metabolism |
| | gi 8895212 | Zinc finger DNA binding protein | 36 | 1 | DNA replication/gene regulation |
| 42 | PPIB_HUMAN | Peptidyl-prolyl cis-trans isomerase B | 40 | 1 | Signal transduction |
| | EGFR_HUMAN | Epidermal growth factor receptor | 40 | 1 | Metabolism |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 48 | 2 | Cytoskeleton/mobility |
| | AHSA1_HUMAN | Activator of 90 kDa heat shock protein ATPase homolog 1 | 37 | 1 | Chaperone/stress response |
| 43 | CIDEA_HUMAN | Cell death activator CIDE-A | 39 | 1 | Signal transduction |
| | PPIB_HUMAN | Peptidyl-prolyl cis-trans isomerase B | 41 | 2 | Signal transduction |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 38 | 2 | Cytoskeleton/mobility |
| | SODC_HUMAN | Superoxide dismutase [Cu-Zn] | 42 | 1 | Protection and detoxification |
| 44 | PPIB_HUMAN | Peptidyl-prolyl cis-trans isomerase B | 61 | 4 | Signal transduction |
| 45 | K22E_HUMAN | Keratin, type II cytoskeletal 2 epidermal | 43 | 2 | Cytoskeleton/mobility |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 41 | 3 | Cytoskeleton/mobility |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 55 | 7 | Cytoskeleton/mobility |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 36 | 5 | Cytoskeleton/mobility Cytoskeleton |
| 46 | PROF1_HUMAN | Profilin-1 | 47 | 2 | Cytoskeleton/mobility |
| | THIO_HUMAN | Thioredoxin | 41 | 1 | Protection and detoxification |
| | PPIA_HUMAN | Peptidyl-prolyl cis-trans isomerase A | 38 | 1 | Signal transduction |
| 47 | HBB_HUMAN | Hemoglobin subunit beta | 38 | 2 | Transport/binding proteins |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 77 | 3 | Cytoskeleton/mobility |
| | LYSC_HUMAN | Lysozyme C | 38 | 1 | Protein synthesis and degradation |
| 48 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 45 | 4 | Cytoskeleton/mobility |
| | gi 119619136 | hCG1790904, isoform CRA_b | 40 | 2 | Unknown |
| | HEM6_HUMAN | Coproporphyrinogen-III oxidase, mitochondrial | 36 | 1 | Metabolism |
| | gi 182311 | Fructose-1,6-bisphosphatase | 35 | 1 | Metabolism |
| 49 | NEK3_HUMAN | Serine/threonine-protein kinase Nek3 | 36 | 1 | Cell cycle |

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|--------------|----|--------------|---|------|----|-----------------------------------|
| | | UBIQ_HUMAN | Ubiquitin | 42 | 3 | Protein synthesis and degradation |
| | | Q7Z2R6_HUMAN | MSTP132 | 35 | 1 | Unknown |
| HepG2 | 1 | FINC_HUMAN | Fibronectin | 48 | 3 | Cytoskeleton/mobility |
| | | DNLI3_HUMAN | DNA ligase 3 | 36 | 1 | Cell cycle |
| | 2 | FINC_HUMAN | Fibronectin | 70 | 7 | Cytoskeleton/mobility |
| | | ITIH2_HUMAN | Inter-alpha-trypsin inhibitor heavy chain H2 | 114 | 7 | Metabolism |
| | | RELN_HUMAN | Reelin | 103 | 7 | Transport/binding proteins |
| | 3 | FINC_HUMAN | Fibronectin | 243 | 14 | Cytoskeleton/mobility |
| | 4 | FINC_HUMAN | Fibronectin | 324 | 20 | Cytoskeleton/mobility |
| | 5 | FINC_HUMAN | Fibronectin | 71 | 8 | Cytoskeleton/mobility |
| | | AT8B4_HUMAN | Probable phospholipid-transporting ATPase IM | 39 | 1 | Metabolism |
| | | A2RQR4_HUMAN | Reverse transcriptase-like protein | 38 | 1 | DNA replication/gene regulation |
| | 6 | FINC_HUMAN | Fibronectin | 70 | 8 | Cytoskeleton/mobility |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 134 | 13 | Protein synthesis and degradation |
| | 7 | A2MG_HUMAN | Alpha-2-macroglobulin | 203 | 12 | Protein synthesis and degradation |
| | 8 | 2DMA_HUMAN | HLA class II histocompatibility antigen, DM alpha chain | 38 | 1 | Immunological response |
| | | NID1_HUMAN | Nidogen-1 | 39 | 1 | Transport/binding proteins |
| | | CERU_HUMAN | Ceruloplasmin | 55 | 2 | Transport/binding proteins |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 225 | 19 | Protein synthesis and degradation |
| | | A2RTY6_HUMAN | Inter-alpha (globulin) inhibitor H2 | 36 | 1 | Metabolism |
| | | FINC_HUMAN | Fibronectin | 59 | 7 | Cytoskeleton/mobility |
| | | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 45 | 2 | Cytoskeleton/mobility |
| | 9 | CHK2_HUMAN | Serine/threonine-protein kinase Chk2 | 0.11 | 1 | Cell cycle |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 177 | 10 | Protein synthesis and degradation |
| | | CERU_HUMAN | Ceruloplasmin | 42 | 3 | Transport/binding proteins |
| | 10 | CO3_HUMAN | Complement C3 | 217 | 10 | Immunological response |
| | | FINC_HUMAN | Fibronectin | 35 | 3 | Cytoskeleton/mobility |
| | 11 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 51 | 4 | Cytoskeleton/mobility |
| | | MTMR4_HUMAN | Myotubularin-related protein 4 | 35 | 1 | |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 45 | 3 | Protein synthesis and degradation |
| | 12 | LG3BP_HUMAN | Galectin-3-binding protein | 177 | 7 | Transport/binding proteins |
| | 13 | ALBU_HUMAN | Human serum albumin | 44 | 4 | Transport/binding proteins |
| | | CTRO_HUMAN | Citron Rho-interacting kinase | 54 | 1 | Cell cycle |
| | | gi 306891 | 90 kDa Heat shock protein | 44 | 4 | Chaperone/stress response |
| | | LG3BP_HUMAN | Galectin-3-binding protein | 40 | 3 | Transport/binding proteins |
| | | TRFE_HUMAN | Serotransferrin | 73 | 6 | Transport/binding proteins |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 48 | 3 | Protein synthesis and degradation |
| | | ARMX2_HUMAN | Armadillo repeat-containing, X-linked protein 2 | 36 | 1 | Transport/binding proteins |

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|--|----|-------------|---|-----|----|-----------------------------------|
| | 14 | TRFE_HUMAN | Serotransferrin | 289 | 12 | Transport/binding proteins |
| | 15 | TRFE_HUMAN | Serotransferrin | 540 | 14 | Transport/binding proteins |
| | | THRΒ_HUMAN | Prothrombin | 38 | 1 | Protein synthesis and degradation |
| | | gi 3282001 | Immunoglobulin heavy chain variable region | 37 | 1 | Immunological response |
| | | NPY5R_HUMAN | Nneuropeptide Y receptor type 5 | 35 | 1 | Signal transduction |
| | 16 | CO3_HUMAN | Complement C3 | 375 | 17 | Immunological response |
| | | CO4A_HUMAN | Complement C4-A | 72 | 6 | Immunological response |
| | | AACT_HUMAN | Alpha-1-antichymotrypsin | 87 | 5 | Protein synthesis and degradation |
| | | FIBA_HUMAN | Fibrinogen alpha chain | 38 | 1 | Protein synthesis and degradation |
| | | FETA_HUMAN | Alpha-fetoprotein | 40 | 2 | Transport/binding proteins |
| | | TRFE_HUMAN | Serotransferrin | 146 | 9 | Transport/binding proteins |
| | | ALBU_HUMAN | Human serum albumin | 106 | 5 | Transport/binding proteins |
| | 17 | FETA_HUMAN | Alpha-fetoprotein | 421 | 11 | Transport/binding proteins |
| | | AACT_HUMAN | Alpha-1-antichymotrypsin | 54 | 2 | Protein synthesis and degradation |
| | | ALBU_HUMAN | Human serum albumin | 490 | 15 | Transport/binding proteins |
| | | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 39 | 1 | Cytoskeleton/mobility |
| | 18 | ALBU_HUMAN | Human serum albumin | 289 | 11 | Transport/binding proteins |
| | | FETA_HUMAN | Alpha-fetoprotein | 332 | 10 | Transport/binding proteins |
| | | F261_HUMAN | 6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1 | 41 | 1 | Metabolism |
| | | TOPB1_HUMAN | DNA topoisomerase 2-binding protein 1 | 36 | 1 | DNA replication/gene regulation |
| | 19 | FETA_HUMAN | Alpha-fetoprotein | 97 | 7 | Transport/binding proteins |
| | | ANGT_HUMAN | Angiotensinogen | 145 | 4 | Signal transduction |
| | | ALBU_HUMAN | Human serum albumin | 206 | 11 | Transport/binding proteins |
| | | FETUA_HUMAN | Alpha-2-HS-glycoprotein | 41 | 1 | Protein synthesis and degradation |
| | 20 | A1AT_HUMAN | Alpha-1-antitrypsin | 312 | 10 | Protein synthesis and degradation |
| | | ANGT_HUMAN | Angiotensinogen | 145 | 6 | Signal transduction |
| | | CBPE_HUMAN | Carboxypeptidase E | 47 | 1 | Protein synthesis and degradation |
| | | SPA11_HUMAN | Serpin A11 | 37 | 2 | Protein synthesis and degradation |
| | 21 | A1AT_HUMAN | Alpha-1-antitrypsin | 250 | 9 | Protein synthesis and degradation |
| | | IDH3A_HUMAN | Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial | 46 | 1 | Metabolism |
| | | FETA_HUMAN | Alpha-fetoprotein | 43 | 1 | Transport/binding proteins |
| | | FIBG_HUMAN | Fibrinogen gamma chain | 49 | 3 | Protein synthesis and degradation |
| | 22 | A1AT_HUMAN | Alpha-1-antitrypsin | 483 | 16 | Protein synthesis and degradation |
| | | APOH_HUMAN | Beta-2-glycoprotein 1 | 49 | 1 | Transport/binding proteins |
| | | ALBU_HUMAN | Human serum albumin | 68 | 8 | Transport/binding proteins |
| | | FETA_HUMAN | Alpha-fetoprotein | 102 | 6 | Transport/binding proteins |
| | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 61 | 8 | Cytoskeleton/mobility |
| | | IDH3A_HUMAN | Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial | 48 | 1 | Metabolism |

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|--|----|-------------|--|-----|---|-----------------------------------|
| | | PEDF_HUMAN | Pigment epithelium-derived factor | 190 | 5 | Protein synthesis and degradation |
| | | MATN3_HUMAN | Matrilin-3 | 73 | 4 | Extracellular matrix |
| | | ALBU_HUMAN | Human serum albumin | 59 | 4 | Transport/binding proteins |
| | 23 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 172 | 8 | Cytoskeleton/mobility |
| | | PAI1_HUMAN | Plasminogen activator inhibitor 1 | 39 | 6 | Protein synthesis and degradation |
| | | ALBU_HUMAN | Human serum albumin | 66 | 4 | Transport/binding proteins |
| | 24 | K22E_HUMAN | Keratin, type II cytoskeletal 2 epidermal | 95 | 5 | Cytoskeleton/mobility |
| | | TACC1_HUMAN | Transforming acidic coiled-coil-containing protein 1 | 35 | 1 | Cell cycle |
| | | A1AG2_HUMAN | Alpha-1-acid glycoprotein 2 | 46 | 1 | Immunological response |
| | | A1AG1_HUMAN | Alpha-1-acid glycoprotein 1 | 46 | 1 | Immunological response |
| | 25 | AIFM2_HUMAN | Apoptosis-inducing factor 2 | 37 | 1 | Metabolism |
| | | ACTB_HUMAN | Actin, cytoplasmic 1 | 48 | 5 | Cytoskeleton/mobility |
| | | ADRO_HUMAN | NADPH:adrenodoxin oxidoreductase, mitochondrial | 43 | 1 | Metabolism |
| | 26 | TRFE_HUMAN | Serotransferrin | 38 | 1 | Transport/binding proteins |
| | | CO3_HUMAN | Complement C3 | 49 | 3 | Immunological response |
| | | SPB5_HUMAN | Serpin B5 | 47 | 2 | Cytoskeleton/mobility |
| | 27 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 38 | 1 | Cytoskeleton/mobility |
| | | SYNE1_HUMAN | Nesprin-1 | 36 | 1 | Cytoskeleton/mobility |
| | | TNR4_HUMAN | Tumor necrosis factor receptor superfamily member 4 | 38 | 1 | Immunological response |
| | | IL1B_HUMAN | Interleukin-1 beta | 38 | 1 | Signal transduction |
| | | CLUS_HUMAN | Clusterin | 37 | 2 | Immunological response |
| | 28 | ALBU_HUMAN | Human serum albumin | 75 | 3 | Transport/binding proteins |
| | | APOE_HUMAN | Apolipoprotein E | 37 | 4 | Transport/binding proteins |
| | | gi 33318880 | Ig heavy chain variable region, VH3 family | 43 | 2 | Immunological response |
| | | gi 266250 | Tetanus toxoid-specific T-cell receptor beta chain | 42 | 1 | Unknown |
| | 29 | L1CAM_HUMAN | Neural cell adhesion molecule L1 | 35 | 1 | Transport/binding proteins |
| | | ALDOA_HUMAN | Fructose-bisphosphate aldolase A | 38 | 1 | Metabolism |
| | | APOE_HUMAN | Apolipoprotein E | 102 | 6 | Transport/binding proteins |
| | 30 | 1B49_HUMAN | HLA class I histocompatibility antigen, B-49 alpha chain | 43 | 1 | Immunological response |
| | | 1B50_HUMAN | HLA class I histocompatibility antigen, B-50 alpha chain | 43 | 1 | Immunological response |
| | | 1B45_HUMAN | HLA class I histocompatibility antigen, B-45 alpha chain | 43 | 1 | Immunological response |
| | | 1B41_HUMAN | HLA class I histocompatibility antigen, B-41 alpha chain | 43 | 1 | Immunological response |
| | | ALBU_HUMAN | Human serum albumin | 77 | 6 | Transport/binding proteins |
| | | APOE_HUMAN | Apolipoprotein E | 62 | 4 | Transport/binding proteins |
| | | IBP1_HUMAN | Insulin-like growth factor-binding protein 1 | 43 | 2 | Signal transduction |
| | 31 | IBP1_HUMAN | Insulin-like growth factor-binding protein 1 | 63 | 4 | Signal transduction |
| | | AMBP_HUMAN | Protein AMBP | 46 | 2 | Protein synthesis and degradation |
| | 32 | IBP1_HUMAN | Insulin-like growth factor-binding protein 1 | 70 | 5 | Signal transduction |

| | | | | | |
|----|--------------|---|-----|----|-----------------------------------|
| | ALBU_HUMAN | Human serum albumin | 43 | 1 | Transport/binding proteins |
| | AMBP_HUMAN | Protein AMBP | 47 | 1 | Protein synthesis and degradation |
| 33 | ALBU_HUMAN | Human serum albumin | 48 | 3 | Transport/binding proteins |
| | CN180_HUMAN | Transmembrane protein C14orf180 | 42 | 1 | Membrane proteins |
| | AMBP_HUMAN | Protein AMBP | 38 | 1 | Protein synthesis and degradation |
| 34 | APOA1_HUMAN | Apolipoprotein A-I | 99 | 6 | Transport/binding proteins |
| | ALBU_HUMAN | Human serum albumin | 39 | 2 | Transport/binding proteins |
| | FER_HUMAN | Proto-oncogene tyrosine-protein kinase FER | 37 | 2 | Metabolism |
| | gi 46811348 | Immunoglobulin heavy chain CDR3 | 40 | 1 | Immunological response |
| 35 | APOA1_HUMAN | Apolipoprotein A-I | 471 | 15 | Transport/binding proteins |
| | CCR6_HUMAN | C-C chemokine receptor type 6 | 42 | 1 | Signal transduction |
| 36 | APOA1_HUMAN | Apolipoprotein A-I | 40 | 4 | Transport/binding proteins |
| | ZN292_HUMAN | Zinc finger protein 292 | 36 | 2 | Transcriptional regulation |
| | B0QYE8_HUMAN | Phospholipase A2, group VI (Cytosolic, calcium-independent) | 38 | 1 | Metabolism |
| 37 | UBP15_HUMAN | Ubiquitin carboxyl-terminal hydrolase 15 | 39 | 1 | Protein synthesis and degradation |
| | HECD1_HUMAN | E3 ubiquitin-protein ligase HECTD1 | 35 | 1 | Protein synthesis and degradation |
| | ZN787_HUMAN | Zinc finger protein 787 | 38 | 1 | Transcriptional regulation |
| 38 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 52 | 4 | Cytoskeleton/mobility |
| | K2C1B_HUMAN | Keratin, type II cytoskeletal 1b | 46 | 1 | Cytoskeleton/mobility |
| | NOL11_HUMAN | Nucleolar protein 11 | 36 | 2 | DNA replication/gene regulation |
| 39 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 46 | 1 | Cytoskeleton/mobility |
| | SUIS_HUMAN | Sucrase-isomaltase, intestinal | 39 | 1 | Metabolism |
| | HP1B3_HUMAN | Heterochromatin protein 1-binding protein 3 | 49 | 2 | Transcriptional regulation |
| 40 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 56 | 4 | Cytoskeleton/mobility |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 52 | 1 | Cytoskeleton/mobility |
| | gi 169166439 | PRDICTED: HPX-153 homeobox | 38 | 2 | Transcriptional regulation |
| | gi 16198460 | GAA protein | 48 | 1 | Metabolism |
| 41 | LTBP1_HUMAN | Latent-transforming growth factor beta-binding protein 1 | 35 | 2 | Transport/binding proteins |
| | SSX4_HUMAN | Protein SSX4 | 44 | 1 | Transcriptional regulation |
| | TEX2_HUMAN | Testis-expressed sequence 2 protein | 39 | 1 | Signal transduction |
| 42 | RET4_HUMAN | Retinol-binding protein 4 | 38 | 2 | Transport/binding proteins |
| 43 | PPIB_HUMAN | Peptidyl-prolyl cis-trans isomerase B | 43 | 3 | Signal transduction |
| | gi 119626643 | Collagen, type XXV, alpha 1, isoform CRA_a | 36 | 1 | Protein synthesis and degradation |
| | SCTR_HUMAN | Secretin receptor | 38 | 1 | Signal transduction |
| | FGFR3_HUMAN | Fibroblast growth factor receptor 3 | 35 | 1 | Signal transduction |
| 44 | LYSC_HUMAN | Lysozyme C | 48 | 1 | Protein synthesis and degradation |
| | ALBU_HUMAN | Human serum albumin | 39 | 1 | Transport/binding proteins |

| | | | | | | |
|----------|--------------|---|---|-----|-----------------------------------|----------------------------|
| | | COF1_HUMAN | Cofilin-1 | 42 | 1 | Cytoskeleton/mobility |
| 45 | TTHY_HUMAN | Transthyretin | 39 | 1 | Transport/binding proteins | |
| | RNAS4_HUMAN | Ribonuclease 4 | 37 | 1 | DNA replication/gene regulation | |
| | gi 386772 | Histone H3 | 38 | 3 | DNA replication/gene regulation | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 84 | 9 | Cytoskeleton/mobility | |
| 46 | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 43 | 3 | Cytoskeleton/mobility | |
| | gi 46811771 | Immunoglobulin heavy chain CDR3 | 38 | 3 | Immunological response | |
| 47 | LYSC_HUMAN | Lysozyme C | 89 | 3 | Protein synthesis and degradation | |
| 48 | MYCN_HUMAN | N-myc proto-oncogene protein | 49 | 1 | Transcription regulation | |
| | FREM1_HUMAN | FRAS1-related extracellular matrix protein 1 | 37 | 1 | Transport/binding proteins | |
| | GRB10_HUMAN | Growth factor receptor-bound protein 10 | 41 | 1 | Signal transduction | |
| 49 | gi 119583891 | Kinesin family member 13B, isoform CRA_b | 45 | 1 | Cytoskeleton/mobility | |
| | gi 619383 | Apolipoprotein D, apoD [human, plasma, Peptide, 246 aa] | 52 | 5 | Transport/binding proteins | |
| SK-Hep-1 | 1 | TSP1_HUMAN | Thrombospondin-1 | 49 | 3 | Cytoskeleton/mobility |
| | 2 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 126 | 6 | Cytoskeleton/mobility |
| | | TSP1_HUMAN | Thrombospondin-1 | 41 | 2 | Cytoskeleton/mobility |
| | 3 | FINC_HUMAN | Fibronectin | 143 | 9 | Cytoskeleton/mobility |
| | | EMR1_HUMAN | EGF-like module-containing mucin-like hormone receptor-like 1 | 36 | 1 | Signal transduction |
| | 4 | TSP1_HUMAN | Thrombospondin-1 | 38 | 1 | Cytoskeleton/mobility |
| | | L1CAM_HUMAN | Neural cell adhesion molecule L1 | 130 | 8 | Membrane proteins |
| | | gi 119602648 | Gasdermin domain containing 1, isoform CRA_a | 42 | 1 | Unknown |
| | 6 | FINC_HUMAN | Fibronectin | 45 | 4 | Cytoskeleton/mobility |
| | 7 | TSP1_HUMAN | Thrombospondin-1 | 100 | 9 | Cytoskeleton/mobility |
| | | 31873147 | Interferon gamma receptor 1 | 36 | 1 | Immunological response |
| | | SPRC_HUMAN | SPARC | 41 | 1 | Extracellular matrix |
| | | CL030_HUMAN | N-terminal acetyltransferase B complex subunit MDM20 | 36 | 1 | Transport/binding proteins |
| | | FINC_HUMAN | Fibronectin | 36 | 2 | Cytoskeleton/mobility |
| | 8 | TSP1_HUMAN | Thrombospondin-1 | 39 | 2 | Cytoskeleton/mobility |
| | | CO6A1_HUMAN | Collagen alpha-1 (VI) chain | 37 | 1 | Transport/binding proteins |
| | | A16L1_HUMAN | Autophagy-related protein 16-1 APG16 like 1 | 43 | 1 | Transport/binding proteins |
| | | DOS_HUMAN | Protein Dos | 42 | 1 | Membrane proteins |
| | | HS90B_HUMAN | Heat shock protein HSP 90-beta | 42 | 2 | Chaperone/stress response |
| | | TRAP1_HUMAN | Heat shock protein 75 kDa, mitochondrial | 42 | 1 | Chaperone/stress response |
| 9 | ITA3_HUMAN | Integrin alpha-3 | 39 | 2 | Cytoskeleton/mobility | |
| | ZFP57_HUMAN | Zinc finger protein 57 homolog | 64 | 1 | DNA replication/gene regulation | |
| | IQGA2_HUMAN | Ras GTPase-activating-like protein IQGAP2 | 36 | 1 | Signal transduction | |

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|----|-------------|---|-----|----|----------------------------|
| | LR37A_HUMAN | Leucine-rich repeat-containing protein 37A3 | 36 | 1 | Transport/binding proteins |
| | PTPRF_HUMAN | Receptor-type tyrosine-protein phosphatase F | 52 | 4 | Transport/binding proteins |
| 10 | A4_HUMAN | Amyloid beta A4 protein | 44 | 2 | Cytoskeleton/mobility |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 79 | 4 | Cytoskeleton/mobility |
| | K22E_HUMAN | Keratin, type II cytoskeletal 2 epidermal | 47 | 3 | Cytoskeleton/mobility |
| | CSTN1_HUMAN | Calsyntenin-1 | 97 | 5 | Transport/binding proteins |
| | SPRC_HUMAN | SPARC | 39 | 1 | Extracellular matrix |
| 11 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 98 | 5 | Cytoskeleton/mobility |
| | LG3BP_HUMAN | Galectin-3-binding protein | 55 | 6 | Transport/binding proteins |
| | K2C8_HUMAN | Keratin, type II cytoskeletal 8 | 37 | 1 | Cytoskeleton/mobility |
| 12 | LG3BP_HUMAN | Galectin-3-binding protein | 497 | 11 | Transport/binding proteins |
| | ACTN1_HUMAN | Alpha-actinin-1 | 55 | 4 | Cytoskeleton/mobility |
| | ACTN4_HUMAN | Alpha-actinin-4 | 50 | 3 | Transport/binding proteins |
| | SPRC_HUMAN | SPARC | 39 | 1 | Extracellular matrix |
| | QSOX1_HUMAN | Sulphydryl oxidase 1 | 35 | 1 | Metabolism |
| 13 | CD166_HUMAN | CD166 antigen | 41 | 1 | Signal transduction |
| | LG3BP_HUMAN | Galectin-3-binding protein | 275 | 10 | Transport/binding proteins |
| | QSOX1_HUMAN | Sulphydryl oxidase 1 | 136 | 7 | Metabolism |
| | ALBU_HUMAN | Human serum albumin | 40 | 2 | Transport/binding proteins |
| 14 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 185 | 9 | Cytoskeleton/mobility |
| | MOES_HUMAN | Moesin | 39 | 1 | Cytoskeleton/mobility |
| | RADI_HUMAN | Radixin | 39 | 1 | Cytoskeleton/mobility |
| | EZRI_HUMAN | Ezrin | 80 | 6 | Cytoskeleton/mobility |
| | TGM2_HUMAN | Protein-glutamine gamma-glutamyltransferase 2 | 41 | 1 | Metabolism |
| | K22E_HUMAN | Keratin, type II cytoskeletal 2 epidermal | 84 | 4 | Cytoskeleton/mobility |
| | SPRC_HUMAN | SPARC | 39 | 1 | Extracellular matrix |
| | QSOX1_HUMAN | Sulphydryl oxidase 1 | 86 | 7 | Metabolism |
| | MYO7A_HUMAN | Myosin-VIIa | 35 | 1 | Cytoskeleton/mobility |
| 15 | EXT1_HUMAN | Exostosin-1 | 37 | 1 | Metabolism |
| | CL011_HUMAN | Uncharacterized protein C12orf11 | 35 | 1 | Unknown |
| | QSOX1_HUMAN | Sulphydryl oxidase 1 | 89 | 6 | Metabolism |
| | TFR1_HUMAN | Transferrin receptor protein 1 | 45 | 1 | Transport/binding proteins |
| | FETA_HUMAN | Alpha-fetoprotein | 57 | 2 | Transport/binding proteins |
| 16 | TRFL_HUMAN | Lactotransferrin | 47 | 2 | Transport/binding proteins |
| | UFO_HUMAN | Tyrosine protein kinase receptor UFO | 40 | 2 | Signal transduction |
| | MOES_HUMAN | Moesin | 51 | 1 | Cytoskeleton/mobility |
| 17 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 35 | 2 | Cytoskeleton/mobility |
| | SNX32_HUMAN | Sorting nexin-32 | 37 | 1 | Unknown |

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|----|--------------|---|-----|---|-----------------------------------|
| | gi 62896815 | Heat shock 70kDa protein 8 isoform 2 variant | 113 | 7 | Chaperone/stress response |
| | ALBU_HUMAN | Human serum albumin | 48 | 3 | Transport/binding proteins |
| 18 | ALBU_HUMAN | Human serum albumin | 141 | 4 | Transport/binding proteins |
| | QSOX1_HUMAN | Sulfhydryl oxidase 1 | 35 | 2 | Metabolism |
| 19 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 36 | 3 | Cytoskeleton/mobility |
| | ALBU_HUMAN | Human serum albumin | 39 | 1 | Transport/binding proteins |
| 20 | AF273054_1 | CTCL tumor antigen | 37 | 1 | Transport/binding proteins |
| | CJ118_HUMAN | Uncharacterized protein C10orf118 | 35 | 1 | Unknown |
| | PDIA1_HUMAN | Protein disulfide-isomerase | 39 | 1 | Chaperone/stress response |
| | K0284_HUMAN | Protein KIAA0284 | 37 | 1 | Unknown |
| | KPYM_HUMAN | Pyruvate kinase isozymes M1/M2 | 78 | 4 | Metabolism |
| 21 | Q96JS8_HUMAN | ZNF589 protein | 49 | 1 | DNA replication/gene regulation |
| | gi 340021 | Alpha-tubulin | 61 | 4 | Cytoskeleton/mobility |
| | gi 340021 | Alpha-tubulin | 281 | 6 | Cytoskeleton/mobility |
| 22 | INADL_HUMAN | InaD like protein | 42 | 2 | Transport/binding proteins |
| | K22E_HUMAN | Keratin, type II cytoskeletal 2 epidermal | 62 | 2 | Cytoskeleton/mobility |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 60 | 3 | Cytoskeleton/mobility |
| 23 | gi 37492 | Alpha-tubulin | 38 | 2 | Cytoskeleton/mobility |
| | EF1A1_HUMAN | Elongation factor 1-alpha 1 | 121 | 6 | Protein synthesis and degradation |
| | NGN3_HUMAN | Neurogenin-3 | 69 | 1 | Protein synthesis and degradation |
| 24 | ENOA_HUMAN | Alpha-enolase | 98 | 3 | Metabolism |
| | ENOG_HUMAN | Gamma-enolase | 46 | 1 | Metabolism |
| | gi 338695 | Beta-tubulin | 52 | 4 | Cytoskeleton/mobility |
| | VIGLN_HUMAN | Vigilin | 37 | 1 | Metabolism |
| | GDN_HUMAN | Glia-derived nexin | 54 | 4 | Protein synthesis and degradation |
| | ENOA_HUMAN | Alpha-enolase | 264 | 5 | Metabolism |
| | EF1A1_HUMAN | Elongation factor 1-alpha 1 | 54 | 3 | Protein synthesis and degradation |
| | EF1A2_HUMAN | Elongation factor 1-alpha 2 | 51 | 1 | Protein synthesis and degradation |
| | IF4A1_HUMAN | Eukaryotic initiation factor 4A-I | 47 | 1 | Protein synthesis and degradation |
| | IF4A2_HUMAN | Eukaryotic initiation factor 4A-II | 47 | 1 | Protein synthesis and degradation |
| 25 | HAUS6_HUMAN | HAUS augmin-like complex subunit 6 | 45 | 1 | Cell cycle |
| | TRFL_HUMAN | Lactotransferrin | 47 | 2 | Transport/binding proteins |
| 26 | UFO_HUMAN | Tyrosine-protein kinase receptor UFO | 40 | 2 | Signal transduction |
| | ENOG_HUMAN | Gamma-enolase | 56 | 2 | Metabolism |
| | ENOA_HUMAN | Alpha-enolase | 104 | 5 | Metabolism |
| | PAI1_HUMAN | Plasminogen activator inhibitor 1 | 83 | 5 | Protein synthesis and degradation |
| 27 | CATB_HUMAN | Cathepsin B | 37 | 1 | Protein synthesis and degradation |
| | ANS1B_HUMAN | Ankyrin repeat and sterile alpha motif domain-containing 1B | 83 | 1 | Protein synthesis and degradation |

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|----|--------------|--|-------------------------------------|-----|-----------------------------------|------------------------|
| | | SPRC_HUMAN | SPARC | 39 | 1 | Extracellular matrix |
| | | PTX3_HUMAN | Pentraxin-related protein PTX3 | 56 | 1 | Immunological response |
| | | ACTG_HUMAN | Actin, cytoplasmic 2 | 91 | 3 | Cytoskeleton/mobility |
| | | ACTB_HUMAN | Actin, cytoplasmic 1 | 362 | 8 | Cytoskeleton/mobility |
| | | ACTG_HUMAN | Actin, cytoplasmic 2 | 189 | 4 | Cytoskeleton/mobility |
| | | POTEE_HUMAN | POTE ankyrin domain family member E | 162 | 5 | Membrane protein |
| | | Q5T8M8_HUMAN | Actin, alpha 1, skeletal muscle | 219 | 7 | Cytoskeleton/mobility |
| 26 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 39 | 4 | Cytoskeleton/mobility | |
| | gi 181192 | procathepsin B | 44 | 2 | Protein synthesis and degradation | |
| | VIME_HUMAN | Vimentin | 48 | 2 | Cytoskeleton/mobility | |
| | RSSA_HUMAN | 40S ribosomal protein SA | 44 | 2 | Protein synthesis and degradation | |
| | SPRC_HUMAN | SPARC | 44 | 2 | Extracellular matrix | |
| 27 | SPRC_HUMAN | SPARC | 40 | 1 | Extracellular matrix | |
| | NDF4_HUMAN | Neurogenic differentiation factor 4 | 41 | 1 | Protein synthesis and degradation | |
| 28 | SPRC_HUMAN | SPARC | 40 | 2 | Extracellular matrix | |
| | EPCR_HUMAN | Endothelial protein C receptor | 43 | 1 | Immunological response | |
| | ALDOA_HUMAN | Fructose-bisphosphate aldolase A | 38 | 1 | Metabolism | |
| 29 | SPRC_HUMAN | SPARC | 40 | 1 | Extracellular matrix | |
| | ROA2_HUMAN | Heterogeneous nuclear ribonucleoproteins A2/B1 | 39 | 1 | DNA replication/gene regulation | |
| | TOP2B_HUMAN | DNA topoisomerase 2-beta | 36 | 1 | DNA replication/gene regulation | |
| 30 | G3P_HUMAN | Glyceraldehyde-3-phosphate dehydrogenase | 38 | 2 | Metabolism | |
| 31 | LDHA_HUMAN | L-lactate dehydrogenase A chain | 49 | 3 | Metabolism | |
| 32 | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 43 | 4 | Cytoskeleton/mobility | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 37 | 3 | Cytoskeleton/mobility | |
| | LANC3_HUMAN | LanC-like protein 3 | 35 | 2 | Protein synthesis and degradation | |
| | gi 189481 | p58/GTA protein kinase | 42 | 1 | Cell cycle | |
| | RNF17_HUMAN | RING finger protein 17 | 37 | 1 | Protein synthesis and degradation | |
| 33 | gi 86438948 | Immunoglobulin heavy chain | 41 | 1 | Immunological response | |
| | Q6P6B6_HUMAN | ZNFX1 protein | 36 | 1 | Unknown | |
| | MLE1_HUMAN | Myosin light chain 1, skeletal muscle isoform | 48 | 1 | Cytoskeleton/mobility | |
| | TIMP1_HUMAN | Metalloproteinase inhibitor 1 | 44 | 2 | Protein synthesis and degradation | |
| 34 | Q8WU60_HUMAN | NADH dehydrogenase (Ubiquinone) flavoprotein 3, 10kDa | 43 | 1 | Metabolism | |
| | gi 2623068 | Yotiao | 37 | 1 | Metabolism | |
| | VIP2_HUMAN | Inositol hexakisphosphate and diphosphoinositol-pentakisphosphate kinase 2 | 38 | 1 | Metabolism | |
| | gi 553309 | Golgin-165 | 38 | 1 | Unknown | |
| | TIMP1_HUMAN | Metalloproteinase inhibitor 1 | 48 | 1 | Protein synthesis and degradation | |
| 35 | TIMP1_HUMAN | Metalloproteinase inhibitor 1 | 48 | 2 | Protein synthesis and degradation | |

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|----|--------------|---|-----|----|-----------------------------------|
| | OTOF_HUMAN | Otoferlin | 36 | 1 | Membrane protein |
| | gi 38383015 | TAF3 protein | 39 | 1 | Protein synthesis and degradation |
| | gi 2181869 | DFFRY | 35 | 1 | Protein synthesis and degradation |
| 37 | AFF4_HUMAN | AF4/FMR2 family member 4 | 41 | 1 | Protein synthesis and degradation |
| | DOK3_HUMAN | Docking protein 3 | 48 | 1 | Protein synthesis and degradation |
| | F154A_HUMAN | Protein FAM154A | 40 | 1 | Unknown |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 38 | 1 | Cytoskeleton/mobility |
| | LRP1B_HUMAN | Low-density lipoprotein receptor-related protein 1B | 37 | 1 | Transport/binding protein |
| 38 | 34528479 | Unnamed protein product | 38 | 1 | Unknown |
| 39 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 80 | 5 | Cytoskeleton/mobility |
| | TSR1_HUMAN | Pre-rRNA-processing protein TSR1 homolog | 52 | 1 | Transport/binding protein |
| | ZKSC3_HUMAN | Zinc finger with KRAB and SCAN domains 3 | 50 | 1 | Protein synthesis and degradation |
| 40 | FXL19_HUMAN | F-box/LRR-repeat protein 19 | 48 | 1 | DNA replication/gene regulation |
| 41 | MYT2_HUMAN | Replication initiation-like protein | 37 | 1 | DNA replication/gene regulation |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 81 | 7 | Cytoskeleton/mobility |
| | TIMP2_HUMAN | Metalloproteinase inhibitor 2 precursor | 43 | 4 | Protein synthesis and degradation |
| 42 | NDK8_HUMAN | Putative nucleoside diphosphate kinase | 36 | 1 | Metabolism |
| | Q32Q12_HUMAN | Nucleoside diphosphate kinase | 39 | 3 | Metabolism |
| | gi 67190163 | Proteoglycan 4 isoform A | 42 | 1 | Transport/binding protein |
| | ZN256_HUMAN | Zinc finger protein 256 | 38 | 1 | Cell cycle |
| 43 | K2C1_HUMAN | Keratin type II cytoskeletal 1 | 185 | 11 | Cytoskeleton/mobility |
| | RAP1A_HUMAN | Ras-related protein Rap-1A | 42 | 1 | Cell cycle |
| | RAP1B_HUMAN | Ras-related protein Rap-1b | 42 | 1 | Cell cycle |
| | 825723 | tropomyosin 227 AA | 42 | 1 | Cytoskeleton/mobility |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 97 | 5 | Cytoskeleton/mobility |
| | PPIB_HUMAN | Peptidylprolyl cistrans isomerase B | 63 | 4 | Metabolism |
| 44 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 37 | 4 | Cytoskeleton/mobility |
| | K1H5_HUMAN | Keratin, type I cuticular Ha5 | 43 | 1 | Cytoskeleton/mobility |
| 45 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 232 | 12 | Cytoskeleton/mobility |
| | PPIA_HUMAN | Peptidyl-prolyl cis-trans isomerase A | 95 | 3 | Metabolism |
| | COF2_HUMAN | Cofilin-2 | 42 | 2 | Cytoskeleton/mobility |
| | COF1_HUMAN | Cofilin-1 | 42 | 2 | Cytoskeleton/mobility |
| | Q32Q12_HUMAN | Nucleoside diphosphate kinase | 49 | 2 | Metabolism |
| | 114319005 | Transthyretin Homo sapiens | 42 | 1 | Transport/binding protein |
| | gi 6166493 | Thioredoxin peroxidase PMP20 | 48 | 2 | Metabolism |
| 46 | TTHY_HUMAN | Transthyretin | 38 | 1 | Membrane protein |
| | PRDX5_HUMAN | Peroxiredoxin-5, mitochondrial | 37 | 3 | Cytoskeleton/mobility |
| | CUTA_HUMAN | Protein CutA | 40 | 2 | Protein synthesis and degradation |

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|----------|----|--------------|--|-----|---|-----------------------------------|
| HCC-S102 | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 36 | 2 | Transport/binding proteins |
| | | CYTC_HUMAN | Cystatin-C | 38 | 2 | Cytoskeleton/mobility |
| | 47 | AT12A_HUMAN | Potassium-transporting ATPase alpha chain 2 | 35 | 1 | Transport/binding proteins |
| | | HBB_HUMAN | Hemoglobin subunit beta | 35 | 1 | Transport/binding proteins |
| | | PROF1_HUMAN | Profilin-1 | 178 | 3 | Cytoskeleton/mobility |
| | | CYTC_HUMAN | Cystatin-C | 37 | 1 | Protein synthesis and degradation |
| | | H4_HUMAN | Histone H4 | 42 | 3 | DNA replication/gene regulation |
| | 48 | HBB_HUMAN | Hemoglobin subunit beta | 35 | 1 | Transport/binding proteins |
| | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 45 | 1 | Cytoskeleton/mobility |
| | | B2MG_HUMAN | Beta-2-microglobulin | 54 | 2 | Transport/binding proteins |
| | | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 54 | 1 | Cytoskeleton/mobility |
| | | CH10_HUMAN | 10 kDa heat shock protein, mitochondrial | 35 | 1 | Chaperone/stress response |
| | | DSC3_HUMAN | Desmocollin-3 | 47 | 1 | Transport/binding proteins |
| | | RS28_HUMAN | 40S ribosomal protein S28 | 42 | 1 | Protein synthesis and degradation |
| | | UBIQ_HUMAN | Ubiquitin | 98 | 3 | Transport/binding proteins |
| | 49 | UBIQ_HUMAN | Ubiquitin | 58 | 4 | Transport/binding proteins |
| | 1 | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 53 | 4 | Cytoskeleton/mobility |
| | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 104 | 7 | Cytoskeleton/mobility |
| | | K2C4_HUMAN | Keratin, type II cytoskeletal 4 | 38 | 2 | Cytoskeleton/mobility |
| | 2 | 85397980 | Coiled coil domain containing 144B Homo sapiens | 37 | 1 | Unknown |
| | | 149363642 | Coiled coil domain containing 144AHomo sapiens | 37 | 1 | Unknown |
| | | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 77 | 4 | Cytoskeleton/mobility |
| | 3 | IF4G2_HUMAN | Eukaryotic translation initiation factor 4 gamma 2 | 35 | 1 | Protein synthesis and degradation |
| | | XPO2_HUMAN | Exportin-2 | 35 | 1 | Transport/binding proteins |
| | 4 | 85397980 | Coiled coil domain containing 144B Homo sapiens | 37 | 1 | Unknown |
| | | 149363642 | Coiled coil domain containing 144AHomo sapiens | 37 | 1 | Unknown |
| | | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 35 | 1 | Cytoskeleton/mobility |
| | | KIF6_HUMAN | Kinesin-like protein KIF6 | 35 | 1 | Cytoskeleton/mobility |
| | 5 | GNDS_HUMAN | Ral guanine nucleotide dissociation stimulator | 38 | 1 | Signal transduction |
| | 6 | ALKB5_HUMAN | Alkylated DNA repair protein alkB homolog 5 | 37 | 1 | Membrane proteins |
| | | GGT1_HUMAN | Gamma-glutamyltransferase type 1 | 37 | 1 | Metabolism |
| | | gi 119624248 | hCG2041343 | 38 | 1 | Unknown |
| | | MIB2_HUMAN | E3 ubiquitin-protein ligase MIB2 | 39 | 2 | Signal transduction |
| | 7 | CO6_HUMAN | Complement component C6 | 45 | 3 | Immunological response |
| | | MYH10_HUMAN | Myosin-10 | 36 | 1 | Cytoskeleton/mobility |
| | | COPG2_HUMAN | Coatomer subunit gamma-2 | 38 | 1 | Transport/binding proteins |
| | 8 | CERU_HUMAN | Ceruloplasmin | 49 | 4 | Transport/binding proteins |
| | | A2MG_HUMAN | Alpha-2-macroglobulin | 36 | 2 | Protein synthesis and degradation |

| | | | | | | |
|----|--------------|---|---------------------------------|----|-----------------------------------|-----------------------|
| | 9 | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 44 | 2 | Cytoskeleton/mobility |
| | | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 40 | 5 | Cytoskeleton/mobility |
| 10 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 51 | 3 | Cytoskeleton/mobility | |
| | CSTN1_HUMAN | Calsyntenin-1 | 42 | 2 | Transport/binding proteins | |
| | ALBU_HUMAN | Human serum albumin | 38 | 2 | Transport/binding proteins | |
| 11 | CSTN1_HUMAN | Calsyntenin-1 | 41 | 1 | Transport/binding proteins | |
| | LG3BP_HUMAN | Galectin-3-binding protein | 79 | 5 | Transport/binding proteins | |
| 12 | LG3BP_HUMAN | Galectin-3-binding protein | 230 | 8 | Transport/binding proteins | |
| | EF2_HUMAN | Elongation factor 2 | 37 | 1 | Protein synthesis and degradation | |
| 13 | HS90B_HUMAN | Heat shock protein HSP 90-beta | 39 | 1 | Chaperone/stress response | |
| | gi 62914009 | HSP90AA1 protein | 47 | 4 | Chaperone/stress response | |
| | LG3BP_HUMAN | Galectin-3-binding protein | 53 | 4 | Transport/binding proteins | |
| 14 | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 48 | 1 | Cytoskeleton/mobility | |
| | HS90B_HUMAN | Heat shock protein HSP 90-beta | 52 | 4 | Chaperone/stress response | |
| | QSOX1_HUMAN | Sulphydryl oxidase 1 | 44 | 4 | Metabolism | |
| | EPHB1_HUMAN | Ephrin type-B receptor 1 | 42 | 1 | Transport/binding proteins | |
| 15 | RCCD1_HUMAN | RCC1 domain-containing protein 1 | 41 | 1 | Unknown | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 39 | 1 | Cytoskeleton/mobility | |
| | B4DYM1_HUMAN | cDNA FLJ60372, highly similar to Myosin-18A | 37 | 1 | Unknown | |
| 16 | ALBU_HUMAN | Human serum albumin | 69 | 5 | Transport/binding proteins | |
| | GFAP_HUMAN | Glial fibrillary acidic protein | 40 | 1 | Transport/binding proteins | |
| | K2C7_HUMAN | Keratin, type II cytoskeletal 7 | 40 | 1 | Cytoskeleton/mobility | |
| | K2C8_HUMAN | Keratin, type II cytoskeletal 8 | 40 | 1 | Cytoskeleton/mobility | |
| | K2C4_HUMAN | Keratin, type II cytoskeletal 4 | 40 | 1 | Cytoskeleton/mobility | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 72 | 1 | Cytoskeleton/mobility | |
| 17 | ALBU_HUMAN | Human serum albumin | 145 | 4 | Transport/binding proteins | |
| 18 | ALBU_HUMAN | Human serum albumin | 135 | 9 | Transport/binding proteins | |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 105 | 6 | Cytoskeleton/mobility | |
| | R4RL1_HUMAN | Reticulon-4 receptor-like 1 | 48 | 1 | Transport/binding proteins | |
| | IGF1A_HUMAN | Insulin-like growth factor IA | 44 | 1 | DNA replication/gene regulation | |
| | TIP60_HUMAN | Histone acetyltransferase KAT5 | 46 | 1 | DNA replication/gene regulation | |
| 19 | S12A7_HUMAN | Solute carrier family 12 member 7 | 42 | 1 | Transport/binding proteins | |
| | PCSK9_HUMAN | Proprotein convertase subtilisin/kexin type 9 | 47 | 2 | Metabolism | |
| | UGDH_HUMAN | UDP-glucose 6-dehydrogenase | 46 | 2 | Metabolism | |
| 20 | FBX37_HUMAN | F-box only protein 37 | 37 | 1 | Transport/binding proteins | |
| | G6PI_HUMAN | Glucose-6-phosphate isomerase | 40 | 5 | Metabolism | |
| 21 | gi 181965 | Elongation factor 1-alpha | 37 | 1 | Protein synthesis and degradation | |
| 22 | TBB5_HUMAN | Tubulin beta chain | 49 | 4 | Cytoskeleton/mobility | |

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|----|--------------|--|--|-----|----|-----------------------------------|
| | | EF1G_HUMAN | Elongation factor 1-gamma | 46 | 1 | Protein synthesis and degradation |
| | | ENOA_HUMAN | Alpha-enolase | 39 | 4 | Metabolism |
| | | EF1G_HUMAN | Elongation factor 1-gamma | 36 | 1 | Protein synthesis and degradation |
| 23 | ENOA_HUMAN | Alpha-enolase | | 365 | 10 | Metabolism |
| | | ENOB_HUMAN | Beta-enolase | 41 | 1 | Metabolism |
| | | 6PGD_HUMAN | 6-phosphogluconate dehydrogenase, decarboxylating | 39 | 1 | Metabolism |
| | | SH3K1_HUMAN | SH3 domain-containing kinase-binding protein 1 | 44 | 3 | Signal transduction |
| | | ENOGL_HUMAN | Gamma-enolase | 177 | 3 | Metabolism |
| 24 | 6PGD_HUMAN | 6-phosphogluconate dehydrogenase, decarboxylating | | 42 | 2 | Metabolism |
| | | PDE7A_HUMAN | High affinity cAMP-specific 3', 5'-cyclic phosphodiesterase 7A | 40 | 1 | Signal transduction |
| | | IDHC_HUMAN | Isocitrate dehydrogenase [NADP] cytoplasmic | 52 | 3 | Metabolism |
| | | ES8L1_HUMAN | Epidermal growth factor receptor kinase substrate 8-like protein 1 | 42 | 2 | Cytoskeleton/mobility |
| 25 | ACTBL_HUMAN | Beta-actin-like protein 2 | | 35 | 1 | Cytoskeleton/mobility |
| | ANKS3_HUMAN | Ankyrin repeat and SAM domain-containing protein 3 | | 35 | 1 | Unknown |
| | ACTB_HUMAN | Actin, cytoplasmic 1 | | 139 | 5 | Cytoskeleton/mobility |
| | B2RPJ1_HUMAN | Actin, beta-like 2 | | 79 | 3 | Cytoskeleton/mobility |
| 26 | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | | 39 | 2 | Cytoskeleton/mobility |
| | GGH_HUMAN | Gamma-glutamyl hydrolase | | 40 | 1 | Metabolism |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | | 111 | 4 | Cytoskeleton/mobility |
| | Q53FF7_HUMAN | CD27-binding (Siva) protein isoform 1 variant | | 40 | 1 | Transport/binding proteins |
| 27 | MIPT3_HUMAN | TRAF3-interacting protein 1 | | 35 | 1 | Signal transduction |
| | ALDOA_HUMAN | Fructose-bisphosphate aldolase A | | 40 | 1 | Metabolism |
| | Q4W5M8_HUMAN | PET112-like (yeast) | | 39 | 1 | Protein synthesis and degradation |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | | 38 | 1 | Cytoskeleton/mobility |
| 28 | DDX53_HUMAN | Probable ATP-dependent RNA helicase DDX53 | | 37 | 1 | DNA replication/gene regulation |
| | ALDOA_HUMAN | Fructose-bisphosphate aldolase A | | 77 | 4 | Metabolism |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | | 41 | 2 | Cytoskeleton/mobility |
| | gi_338057 | Sulfated glycoprotein-2 | | 95 | 3 | Transport/binding proteins |
| | RLA0_HUMAN | 60S acidic ribosomal protein P0 | | 34 | 1 | Transport/binding proteins |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | | 65 | 4 | Cytoskeleton/mobility |
| 29 | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | | 40 | 2 | Cytoskeleton/mobility |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | | 73 | 7 | Cytoskeleton/mobility |
| | ANXA2_HUMAN | Annexin A2 | | 51 | 4 | Cytoskeleton/mobility |
| | NOSTN_HUMAN | Nostrin | | 35 | 1 | Transport/binding proteins |
| 30 | G3P_HUMAN | Glyceraldehyde-3-phosphate dehydrogenase | | 128 | 7 | Metabolism |
| | FGL1_HUMAN | Fibrinogen-like protein 1 | | 57 | 1 | Signal transduction |

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|----|--------------|---|-----|---|---------------------------------|
| | CHD8_HUMAN | Chromodomain-helicase-DNA-binding protein 8 | 38 | 1 | DNA replication/gene regulation |
| | K2C1B_HUMAN | Keratin, type II cytoskeletal 1b | 45 | 1 | Cytoskeleton/mobility |
| 31 | FGL1_HUMAN | Fibrinogen-like protein 1 | 83 | 6 | Signal transduction |
| 32 | K1C13_HUMAN | Keratin, type I cytoskeletal 13 | 48 | 1 | Cytoskeleton/mobility |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 193 | 9 | Cytoskeleton/mobility |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 82 | 7 | Cytoskeleton/mobility |
| | ENOA_HUMAN | Alpha-enolase | 48 | 1 | Metabolism |
| 33 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 109 | 7 | Cytoskeleton/mobility |
| | K1C10_HUMAN | Keratin, type I cytoskeletal 10 | 35 | 1 | Cytoskeleton/mobility |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 39 | 4 | Cytoskeleton/mobility |
| 34 | TPIS_HUMAN | Triosephosphate isomerase | 39 | 2 | Metabolism |
| | CHD1_HUMAN | Chromodomain-helicase-DNA-binding protein 1 | 38 | 1 | DNA replication/gene regulation |
| | gi 119577863 | hCG2041066 | 38 | 1 | Unknown |
| | NSF1C_HUMAN | NSFL1 cofactor p47 | 38 | 1 | Cytoskeleton/mobility |
| 35 | CP20A_HUMAN | Cytochrome P450 20A1 | 43 | 1 | Metabolism |
| | AGA11_HUMAN | Arf-GAP, GTPase, ANK repeat and PH domain-containing protein 11 | 40 | 1 | Metabolism |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 42 | 4 | Cytoskeleton/mobility |
| 36 | Q5VY93_HUMAN | Rho/rac guanine nucleotide exchange factor (GEF) 2 | 39 | 1 | Signal transduction |
| | CF223_HUMAN | Uncharacterized protein C6orf223 | 42 | 2 | Unknown |
| | gi 119622046 | hCG1646228 | 42 | 1 | Unknown |
| | Q8TEM7_HUMAN | FLJ00166 protein | 41 | 1 | DNA replication/gene regulation |
| 37 | K22E_HUMAN | Keratin, type II cytoskeletal 2 epidermal | 38 | 2 | Cytoskeleton/mobility |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 36 | 2 | Cytoskeleton/mobility |
| | FIGL1_HUMAN | Fidgetin-like protein 1 | 38 | 1 | Metabolism |
| 38 | GDF15_HUMAN | Growth/differentiation factor 15 | 40 | 1 | Signal transduction |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 36 | 2 | Cytoskeleton/mobility |
| | PRDX1_HUMAN | Peroxiredoxin-1 | 35 | 2 | Transport/binding proteins |
| 39 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 70 | 4 | Cytoskeleton/mobility |
| | EPHB1_HUMAN | Ephrin type-B receptor 1 | 41 | 2 | Transport/binding proteins |
| | gi 6643713 | Immunoglobulin V lambda/J lambda light chain | 42 | 1 | Immunological reaponse |
| | K1C9_HUMAN | Keratin, type I cytoskeletal 9 | 42 | 2 | Cytoskeleton/mobility |
| 40 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 46 | 2 | Cytoskeleton/mobility |
| | Q3SYD4_HUMAN | SSX4 protein | 35 | 1 | DNA replication/gene regulation |
| | AP2B_HUMAN | Transcription factor AP-2 beta | 35 | 1 | DNA replication/gene regulation |
| | ZN536_HUMAN | Zinc finger protein 536 | 35 | 1 | DNA replication/gene regulation |
| 41 | EMIL1_HUMAN | EMILIN-1 | 42 | 2 | Cytoskeleton/mobility |
| | ZN511_HUMAN | Zinc finger protein 511 | 41 | 1 | DNA replication/gene regulation |

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|----|--------------|---|-----|---|-----------------------------------|
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 41 | 2 | Cytoskeleton/mobility |
| 42 | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 41 | 4 | Cytoskeleton/mobility |
| | Q9H2G3_HUMAN | CTCL tumor antigen se20-7 | 42 | 2 | Unknown |
| | RGL1_HUMAN | Ral guanine nucleotide dissociation stimulator-like 1 | 42 | 2 | Signal transduction |
| | BAI1_HUMAN | Brain-specific angiogenesis inhibitor 1 | 37 | 2 | Signal transduction |
| | ZN787_HUMAN | Zinc finger protein 787 | 43 | 1 | DNA replication/gene regulation |
| 43 | E41L2_HUMAN | Band 4.1-like protein 2 | 39 | 2 | Membrane proteins |
| | ZPR1_HUMAN | Zinc finger protein ZPR1 | 38 | 2 | DNA replication/gene regulation |
| | HERC2_HUMAN | Probable E3 ubiquitin-protein ligase HERC2 | 38 | 1 | Transport/binding proteins |
| | K2C1B_HUMAN | Keratin, type II cytoskeletal 1b | 47 | 1 | Cytoskeleton/mobility |
| 44 | MCES_HUMAN | mRNA cap guanine-N7 methyltransferase | 54 | 1 | DNA replication/gene regulation |
| | PPIA_HUMAN | Peptidyl-prolyl cis-trans isomerase A | 52 | 2 | Protein synthesis and degradation |
| | ZDBF2_HUMAN | DBF4-type zinc finger-containing protein 2 | 35 | 2 | Unknown |
| 45 | PERI_HUMAN | Peripherin | 41 | 1 | Cytoskeleton/mobility |
| | PPIA_HUMAN | Peptidyl-prolyl cis-trans isomerase A | 39 | 2 | Protein synthesis and degradation |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 37 | 1 | Cytoskeleton/mobility |
| 47 | THIO_HUMAN | Thioredoxin | 38 | 3 | Transport/binding proteins |
| | gi 169176438 | PREDICTED: hypothetical protein | 35 | 1 | Unknown |
| | H4_HUMAN | Histone H4 | 38 | 2 | DNA replication/gene regulation |
| | THIO_HUMAN | Thioredoxin | 39 | 1 | Transport/binding proteins |
| 48 | gi 171853445 | Immunoglobulin heavy chain variable region | 38 | 1 | Immunological response |
| | CO5A1_HUMAN | Collagen alpha-1(V) chain | 37 | 1 | Protein synthesis and degradation |
| | K2C1_HUMAN | Keratin, type II cytoskeletal 1 | 119 | 6 | Cytoskeleton/mobility |
| 49 | UBIQ_HUMAN | Ubiquitin | 52 | 4 | Transport/binding proteins |