

Accession

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Description

Keratin, type II cytoskeletal 2 epidermal OS=Homo sapiens OX=9606 GN=KRT2 PE=1 SV=2 - [K22E_HUMAN]

Keratin, type I cytoskeletal 9 OS=Homo sapiens OX=9606 GN=KRT9 PE=1 SV=3 - [K1C9_HUMAN]

Keratin, type I cytoskeletal 16 OS=Homo sapiens OX=9606 GN=KRT16 PE=1 SV=4 - [K1C16_HUMAN]

E3 ubiquitin-protein ligase SMURF2 OS=Homo sapiens OX=9606 GN=SMURF2 PE=1 SV=1 - [SMUF2_HUMAN]

Keratin 1 OS=Homo sapiens OX=9606 GN=KRT1 PE=3 SV=1 - [H6VRF8_HUMAN]

Keratin, type I cytoskeletal 14 OS=Homo sapiens OX=9606 GN=KRT14 PE=1 SV=4 - [K1C14_HUMAN]

Keratin, type I cytoskeletal 10 OS=Homo sapiens OX=9606 GN=KRT10 PE=1 SV=6 - [K1C10_HUMAN]

cDNA FLJ53910, highly similar to Keratin, type II cytoskeletal 6A OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DRRO_HUMAN]

Immunoglobulin lambda constant 7 OS=Homo sapiens OX=9606 GN=IGLC7 PE=1 SV=3 - [IGLC7_HUMAN]

Keratin, type I cytoskeletal 17 OS=Homo sapiens OX=9606 GN=KRT17 PE=1 SV=2 - [K1C17_HUMAN]

Vimentin OS=Homo sapiens OX=9606 GN=VIM PE=1 SV=4 - [VIME_HUMAN]

Keratin, type II cytoskeletal 6B OS=Homo sapiens OX=9606 GN=KRT6B PE=1 SV=5 - [K2C6B_HUMAN]

Dermcidin OS=Homo sapiens OX=9606 GN=DCD PE=1 SV=2 - [DCD_HUMAN]

Keratin, type II cytoskeletal 5 OS=Homo sapiens OX=9606 GN=KRT5 PE=1 SV=3 - [K2C5_HUMAN]

CSDA protein variant (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q59EB5_HUMAN]

Protein S100-A7 OS=Homo sapiens OX=9606 GN=S100A7 PE=1 SV=4 - [S10A7_HUMAN]

Myosin light polypeptide 6 OS=Homo sapiens OX=9606 GN=MYL6 PE=1 SV=1 - [F8W1R7_HUMAN]

cDNA FLJ55253, highly similar to Actin, cytoplasmic 1 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DW52_HUMAN]

Histone H4 OS=Homo sapiens OX=9606 GN=HIST1H4A PE=1 SV=2 - [H4_HUMAN]

Protein S100-A8 OS=Homo sapiens OX=9606 GN=S100A8 PE=1 SV=1 - [S10A8_HUMAN]

Immunoglobulin J chain (Fragment) OS=Homo sapiens OX=9606 GN=JCHAIN PE=1 SV=1 - [C9JA05_HUMAN]

Hornerin OS=Homo sapiens OX=9606 GN=HRNR PE=1 SV=2 - [HORN_HUMAN]

Prolactin-inducible protein OS=Homo sapiens OX=9606 GN=PIP PE=1 SV=1 - [PIP_HUMAN]

Albumin (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [F6KPG5_HUMAN]

cDNA FLJ60647, highly similar to Keratin, type II cytoskeletal 6B OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DKV4_HUMAN]

Zymogen granule protein 16 homolog B OS=Homo sapiens OX=9606 GN=ZG16B PE=1 SV=1 - [A0A0C4DGN4_HUMAN]

Polyubiquitin-C (Fragment) OS=Homo sapiens OX=9606 GN=UBC PE=1 SV=8 - [F5H6Q2_HUMAN]

Pentraxin-related protein PTX3 OS=Homo sapiens OX=9606 GN=PTX3 PE=1 SV=3 - [PTX3_HUMAN]

Serpin B3 OS=Homo sapiens OX=9606 GN=SERPINB3 PE=1 SV=2 - [SPB3_HUMAN]

Immunoglobulin light chain (Fragment) OS=Homo sapiens OX=9606 PE=1 SV=1 - [Q0KKI6_HUMAN]

Myosin-reactive immunoglobulin heavy chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q9UL72_HUMAN]

Myosin-reactive immunoglobulin light chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q9UL78_HUMAN]

40S ribosomal protein S3 OS=Homo sapiens OX=9606 GN=RPS3 PE=1 SV=1 - [E9PPU1_HUMAN]

Cystatin-A OS=Homo sapiens OX=9606 GN=CSTA PE=1 SV=1 - [CYTA_HUMAN]

SCCA1/SCCA2 fusion protein OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q5K684_HUMAN]

Similar to ribosomal protein S8 (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q9BS10_HUMAN]

IgG L chain OS=Homo sapiens OX=9606 PE=2 SV=1 - [S6BAN6_HUMAN]

cDNA FLJ60461, highly similar to Peroxiredoxin-2 (EC 1.11.1.15) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DF70_HUMAN]

cDNA FLJ53012, highly similar to Tubulin beta-7 chain OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DMJ5_HUMAN]

Annexin A2 OS=Homo sapiens OX=9606 GN=ANXA2 PE=1 SV=2 - [ANXA2_HUMAN]

Ig heavy chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [A0A068LN13_HUMAN]

C1 segment protein (Fragment) OS=Homo sapiens OX=9606 GN=C1 segment PE=4 SV=1 - [A0M8Q9_HUMAN]

10E8 heavy chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [A0A193CHR0_HUMAN]

cDNA FLJ53662, highly similar to Actin, alpha skeletal muscle OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7Z6P1_HUMAN]

Profilin-1 OS=Homo sapiens OX=9606 GN=PFN1 PE=1 SV=2 - [PROF1_HUMAN]

Immunoglobulin heavy variable 3-49 OS=Homo sapiens OX=9606 GN=IGHV3-49 PE=3 SV=1 - [HV349_HUMAN]

Heat shock protein beta-1 OS=Homo sapiens OX=9606 GN=HSPB1 PE=1 SV=2 - [HSPB1_HUMAN]

Annexin A1 OS=Homo sapiens OX=9606 GN=ANXA1 PE=1 SV=2 - [ANXA1_HUMAN]

Actin-like protein (Fragment) OS=Homo sapiens OX=9606 GN=ACT PE=4 SV=1 - [Q562Z4_HUMAN]

MS-D1 light chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [A0A0X9TD47_HUMAN]

Zinc-alpha-2-glycoprotein OS=Homo sapiens OX=9606 GN=AZGP1 PE=1 SV=2 - [ZA2G_HUMAN]

Nuclease-sensitive element-binding protein 1 OS=Homo sapiens OX=9606 GN=YBX1 PE=1 SV=3 - [YBOX1_HUMAN]

Chaperone Hsp10, affects cell division OS=Homo sapiens OX=9606 GN=groES PE=4 SV=1 - [A0A2P9ALU8_HUMAN]
Keratin, type I cytoskeletal 10 (Fragment) OS=Homo sapiens OX=9606 GN=KRT10 PE=1 SV=1 - [A0A1B0GVI3_HUMAN]
Protein S100-P OS=Homo sapiens OX=9606 GN=S100P PE=1 SV=2 - [S100P_HUMAN]
40S ribosomal protein S20 OS=Homo sapiens OX=9606 GN=RPS20 PE=1 SV=1 - [E5RIP1_HUMAN]
Arginase-1 OS=Homo sapiens OX=9606 GN=ARG1 PE=1 SV=2 - [ARGI1_HUMAN]
cDNA FLJ55805, highly similar to Keratin, type II cytoskeletal 4 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DRW1_HUMAN]
Tubulin alpha chain OS=Homo sapiens OX=9606 PE=2 SV=1 - [B3KPS3_HUMAN]
SCCA2b OS=Homo sapiens OX=9606 GN=SCCA2 PE=2 SV=1 - [Q9BYF7_HUMAN]
Keratin, type I cytoskeletal 19 OS=Homo sapiens OX=9606 GN=KRT19 PE=1 SV=4 - [K1C19_HUMAN]
Cofilin-1 (Fragment) OS=Homo sapiens OX=9606 GN=CFL1 PE=1 SV=1 - [E9PQB7_HUMAN]
Myosin-9 OS=Homo sapiens OX=9606 GN=MYH9 PE=1 SV=4 - [MYH9_HUMAN]
Anti-(ED-B) scFV (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=2 - [A2KBC1_HUMAN]
Ras-related protein Rab-6B (Fragment) OS=Homo sapiens OX=9606 GN=RAB6B PE=4 SV=1 - [C9JB90_HUMAN]
Peroxiredoxin-1 (Fragment) OS=Homo sapiens OX=9606 GN=PRDX1 PE=1 SV=1 - [A0A0A0MSI0_HUMAN]
Keratin, type II cytoskeletal 78 (Fragment) OS=Homo sapiens OX=9606 GN=KRT78 PE=3 SV=2 - [H0YI54_HUMAN]
60S acidic ribosomal protein P0 (Fragment) OS=Homo sapiens OX=9606 GN=RPLP0 PE=1 SV=1 - [F8VPE8_HUMAN]
BPI fold-containing family A member 1 OS=Homo sapiens OX=9606 GN=BPIFA1 PE=1 SV=1 - [BPIA1_HUMAN]
Hemoglobin alpha 2 (Fragment) OS=Homo sapiens OX=9606 GN=HBA2 PE=3 SV=1 - [Q96T46_HUMAN]
cDNA FLJ53742, highly similar to Urokinase-type plasminogen activator (EC 3.4.21.73) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DNJ4_HUMAN]
cDNA FLJ54552, highly similar to Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DUQ1_HUMAN]
40S ribosomal protein S13 OS=Homo sapiens OX=9606 GN=RPS13 PE=1 SV=1 - [J3KMX5_HUMAN]
Pyruvate kinase (Fragment) OS=Homo sapiens OX=9606 GN=PKM PE=1 SV=1 - [H3BTN5_HUMAN]
Galactin-7 OS=Homo sapiens OX=9606 GN=LGALS7 PE=1 SV=2 - [LEG7_HUMAN]
Lysozyme OS=Homo sapiens OX=9606 GN=LYZ PE=1 SV=1 - [F8VV32_HUMAN]
40S ribosomal protein S16 (Fragment) OS=Homo sapiens OX=9606 GN=RPS16 PE=1 SV=1 - [M0QX76_HUMAN]
Immunoglobulin kappa variable 2D-29 OS=Homo sapiens OX=9606 GN=IGKV2D-29 PE=3 SV=1 - [KVD29_HUMAN]
Junction plakoglobin, isoform CRA_a OS=Homo sapiens OX=9606 GN=JUP PE=4 SV=1 - [A0A024R1X8_HUMAN]
Keratin 13 OS=Homo sapiens OX=9606 GN=KRT13 PE=1 SV=1 - [A1A4E9_HUMAN]
MS-C1 heavy chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [A0A125U0U7_HUMAN]
Uncharacterized protein OS=Homo sapiens OX=9606 PE=4 SV=1 - [M0QZK8_HUMAN]
Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens OX=9606 GN=GAPDH PE=1 SV=3 - [G3P_HUMAN]
Rheumatoid factor RF-IP9 (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [A2J1M2_HUMAN]
Chaperone Hsp10, affects cell division OS=Homo sapiens OX=9606 GN=groES PE=4 SV=1 - [A0A2P9ANF3_HUMAN]
Uncharacterized protein (Fragment) OS=Homo sapiens OX=9606 PE=3 SV=1 - [V9GZNO_HUMAN]
14-3-3 protein sigma OS=Homo sapiens OX=9606 GN=SFN PE=1 SV=1 - [1433S_HUMAN]
Caspase-14 OS=Homo sapiens OX=9606 GN=CASP14 PE=1 SV=2 - [CASPE_HUMAN]
Nucleophosmin OS=Homo sapiens OX=9606 GN=NPM1 PE=1 SV=2 - [NPM_HUMAN]
60S ribosomal protein L17 (Fragment) OS=Homo sapiens OX=9606 GN=RPL17 PE=3 SV=1 - [J3KSJ0_HUMAN]
Translation elongation factor 1 alpha 1-like 14 OS=Homo sapiens OX=9606 GN=EEF1A1L14 PE=2 SV=1 - [Q96RE1_HUMAN]
Tropomyosin alpha-3 chain OS=Homo sapiens OX=9606 GN=TPM3 PE=1 SV=1 - [A0A087WWU8_HUMAN]
Uncharacterized protein OS=Homo sapiens OX=9606 GN=DKFZp686K03196 PE=1 SV=1 - [Q6N095_HUMAN]
Tubulin alpha-4A chain OS=Homo sapiens OX=9606 GN=TUBA4A PE=1 SV=1 - [TBA4A_HUMAN]
L-lactate dehydrogenase A chain OS=Homo sapiens OX=9606 GN=LDHA PE=1 SV=2 - [LDHA_HUMAN]
Proteasome subunit beta type-3 OS=Homo sapiens OX=9606 GN=PSMB3 PE=1 SV=1 - [A0A087WXQ8_HUMAN]
Keratin, type II cytoskeletal 80 OS=Homo sapiens OX=9606 GN=KRT80 PE=1 SV=2 - [K2C80_HUMAN]
Myosin-reactive immunoglobulin light chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q9UL83_HUMAN]
Lactotransferrin (Fragment) OS=Homo sapiens OX=9606 GN=LTF PE=1 SV=1 - [E7EQB2_HUMAN]
cDNA FLJ14473 fis, clone MAMMA1001080, highly similar to Homo sapiens SNC73 protein (SNC73) mRNA OS=Homo sapiens OX=9606 PE=2 SV=1
40S ribosomal protein S14 OS=Homo sapiens OX=9606 GN=RPS14 PE=1 SV=3 - [RS14_HUMAN]
Histone H2B OS=Homo sapiens OX=9606 PE=2 SV=1 - [A8K9J7_HUMAN]
Haptoglobin OS=Homo sapiens OX=9606 GN=HP PE=1 SV=1 - [HPT_HUMAN]
14-3-3 protein zeta/delta OS=Homo sapiens OX=9606 GN=YWHAZ PE=1 SV=1 - [1433Z_HUMAN]
Protease serine 1 (Fragment) OS=Homo sapiens OX=9606 GN=PRSS1 PE=4 SV=1 - [Q86W20_HUMAN]
Triosephosphate isomerase (Fragment) OS=Homo sapiens OX=9606 GN=TPI1 PE=1 SV=1 - [U3KQF3_HUMAN]
Alpha-1-antitrypsin (Fragment) OS=Homo sapiens OX=9606 GN=SERPINA1 PE=1 SV=1 - [G3V5R8_HUMAN]
PNAS-20 OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q9BZU1_HUMAN]

Uncharacterized protein DKFZp686C15213 OS=Homo sapiens OX=9606 GN=DKFZp686C15213 PE=2 SV=1 - [Q6MZU6_HUMAN]
cDNA FLJ53691, highly similar to Serotransferrin OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4E1B2_HUMAN]
Rho GDP-dissociation inhibitor 2 (Fragment) OS=Homo sapiens OX=9606 GN=ARHGDI2 PE=1 SV=1 - [F5H2R5_HUMAN]
IgG L chain OS=Homo sapiens OX=9606 PE=2 SV=1 - [S6B2A1_HUMAN]
Tropomyosin alpha-4 chain (Fragment) OS=Homo sapiens OX=9606 GN=TPM4 PE=1 SV=1 - [K7ENT6_HUMAN]
Stress-70 protein, mitochondrial (Fragment) OS=Homo sapiens OX=9606 GN=HSPA9 PE=1 SV=1 - [D6RA73_HUMAN]
Desmoplakin OS=Homo sapiens OX=9606 GN=DSP PE=1 SV=3 - [DESP_HUMAN]
Cathepsin D (Fragment) OS=Homo sapiens OX=9606 GN=CTSD PE=1 SV=1 - [H7C1V0_HUMAN]
Keratin, type II cuticular Hb4 OS=Homo sapiens OX=9606 GN=KRT84 PE=2 SV=2 - [KRT84_HUMAN]
Keratin, type II cytoskeletal 79 OS=Homo sapiens OX=9606 GN=KRT79 PE=1 SV=2 - [K2C79_HUMAN]
Protein-glutamine gamma-glutamyltransferase E OS=Homo sapiens OX=9606 GN=TGM3 PE=1 SV=4 - [TGM3_HUMAN]
40S ribosomal protein S15a OS=Homo sapiens OX=9606 GN=RPS15A PE=1 SV=1 - [I3L3P7_HUMAN]
FGA protein (Fragment) OS=Homo sapiens OX=9606 GN=FGA PE=2 SV=1 - [Q3KRA7_HUMAN]
Mucin-7 (Fragment) OS=Homo sapiens OX=9606 GN=MUC7 PE=1 SV=1 - [D6RHX1_HUMAN]
40S ribosomal protein S3a (Fragment) OS=Homo sapiens OX=9606 GN=RPS3A PE=1 SV=1 - [D6RGE0_HUMAN]
Mucin (Fragment) OS=Homo sapiens OX=9606 GN=MUC5AC PE=2 SV=1 - [Q14887_HUMAN]
Keratin 77 OS=Homo sapiens OX=9606 GN=KRT77 PE=1 SV=1 - [Q0IIN1_HUMAN]
RPL27/NME2 fusion protein (Fragment) OS=Homo sapiens OX=9606 GN=RPL27 PE=2 SV=1 - [E4W6B6_HUMAN]
Fatty acid-binding protein, epidermal OS=Homo sapiens OX=9606 GN=FABP5 PE=1 SV=3 - [FABP5_HUMAN]
Cellular tumor antigen p53 (Fragment) OS=Homo sapiens OX=9606 GN=TP53 PE=3 SV=1 - [A0A0M4B4Y9_HUMAN]
60S acidic ribosomal protein P2 (Fragment) OS=Homo sapiens OX=9606 GN=RPLP2 PE=1 SV=1 - [HOYDD8_HUMAN]
Cell growth-inhibiting protein 34 OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q08ES8_HUMAN]
Caveolin-1 OS=Homo sapiens OX=9606 GN=CAV1 PE=1 SV=4 - [CAV1_HUMAN]
Cathepsin G OS=Homo sapiens OX=9606 GN=CTSG PE=1 SV=2 - [CATG_HUMAN]
Endoplasmic reticulum chaperone BiP OS=Homo sapiens OX=9606 GN=HSPA5 PE=1 SV=2 - [BIP_HUMAN]
40S ribosomal protein S18 OS=Homo sapiens OX=9606 GN=RPS18 PE=1 SV=3 - [RS18_HUMAN]
CAPZB protein (Fragment) OS=Homo sapiens OX=9606 GN=CAPZB PE=2 SV=1 - [Q7L4N0_HUMAN]
Putative nucleoside diphosphate kinase OS=Homo sapiens OX=9606 GN=NME2P1 PE=5 SV=1 - [NDK8_HUMAN]
Thioredoxin OS=Homo sapiens OX=9606 GN=TXN PE=1 SV=3 - [THIO_HUMAN]
60S ribosomal protein L28 OS=Homo sapiens OX=9606 GN=RPL28 PE=1 SV=1 - [HOYLP6_HUMAN]
cDNA FLJ78142, highly similar to Homo sapiens transglutaminase 1 (K polypeptide epidermal type I, protein-glutamine-gamma-glutamyltrans
60S ribosomal protein L23 OS=Homo sapiens OX=9606 GN=RPL23 PE=1 SV=1 - [J3KTJ3_HUMAN]
Ketosteroid isomerase OS=Homo sapiens OX=9606 GN=BQ8482_380124 PE=4 SV=1 - [A0A2P9ARY7_HUMAN]
Myeloperoxidase OS=Homo sapiens OX=9606 GN=MPO PE=1 SV=1 - [PERM_HUMAN]
Sulfated glycoprotein 2 (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q6LDQ3_HUMAN]
60S ribosomal protein L27a OS=Homo sapiens OX=9606 GN=RPL27A PE=1 SV=1 - [E9PJD9_HUMAN]
cDNA FLJ54625, highly similar to ATP synthase subunit alpha, mitochondrial (EC 3.6.3.14) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DGW
Annexin (Fragment) OS=Homo sapiens OX=9606 GN=ANXA3 PE=1 SV=1 - [D6RCA8_HUMAN]
Heat shock 70kDa protein 8 isoform 2 variant (Fragment) OS=Homo sapiens OX=9606 PE=1 SV=1 - [Q53HF2_HUMAN]
60S ribosomal protein L4 (Fragment) OS=Homo sapiens OX=9606 GN=RPL4 PE=1 SV=1 - [H3BU31_HUMAN]
ATP synthase subunit beta (Fragment) OS=Homo sapiens OX=9606 GN=ATP5F1B PE=1 SV=1 - [HOYH81_HUMAN]
Keratin, type II cytoskeletal 78 OS=Homo sapiens OX=9606 GN=KRT78 PE=1 SV=2 - [K2C78_HUMAN]
60S ribosomal protein L31 OS=Homo sapiens OX=9606 GN=RPL31 PE=1 SV=1 - [B8ZZK4_HUMAN]
RPS2 protein (Fragment) OS=Homo sapiens OX=9606 GN=RPS2 PE=2 SV=2 - [Q9BSW5_HUMAN]
40S ribosomal protein SA (Fragment) OS=Homo sapiens OX=9606 GN=RPSA PE=1 SV=2 - [F8WD59_HUMAN]
Uncharacterized protein OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7ZW15_HUMAN]
Protein S100 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B2R4M6_HUMAN]
Receptor of-activated protein C kinase 1 (Fragment) OS=Homo sapiens OX=9606 GN=RACK1 PE=1 SV=1 - [HOY9P0_HUMAN]
Fructose-bisphosphate aldolase OS=Homo sapiens OX=9606 GN=ALDOA PE=1 SV=1 - [H3BQN4_HUMAN]
Immunoglobulin alpha-2 heavy chain OS=Homo sapiens OX=9606 PE=1 SV=1 - [IGA2_HUMAN]
Gasdermin-A (Fragment) OS=Homo sapiens OX=9606 GN=GSDMA PE=1 SV=8 - [J3KRG2_HUMAN]
GTP-binding nuclear protein Ran (Fragment) OS=Homo sapiens OX=9606 GN=RAN PE=1 SV=1 - [HOYFC6_HUMAN]
cDNA FLJ54303, highly similar to Heat shock 70 kDa protein 1 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DFN9_HUMAN]
Alpha-actinin-1 (Fragment) OS=Homo sapiens OX=9606 GN=ACTN1 PE=1 SV=1 - [HOYJ11_HUMAN]
Protein S100-A14 OS=Homo sapiens OX=9606 GN=S100A14 PE=1 SV=1 - [S10AE_HUMAN]
Follicle-stimulating hormone beta subunit (Fragment) OS=Homo sapiens OX=9606 GN=FSHB PE=3 SV=1 - [Q1W658_HUMAN]

GCT-A5 heavy chain variable region (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [A0A0X9T0H6_HUMAN]
Desmoglein-1 OS=Homo sapiens OX=9606 GN=DSG1 PE=1 SV=2 - [DSG1_HUMAN]
ADP/ATP translocase 3 OS=Homo sapiens OX=9606 GN=SLC25A6 PE=1 SV=4 - [ADT3_HUMAN]
Secretoglobin family 1D member 2 OS=Homo sapiens OX=9606 GN=SCGB1D2 PE=2 SV=1 - [SG1D2_HUMAN]
60S ribosomal protein L18 (Fragment) OS=Homo sapiens OX=9606 GN=RPL18 PE=1 SV=1 - [F8VUA6_HUMAN]
cDNA FLJ40872 fis, clone TUTER2000283, highly similar to Homo sapiens transformer-2-beta (SFRS10) gene OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7Z9V0_HUMAN]
Histone H1.2 OS=Homo sapiens OX=9606 GN=HIST1H1C PE=1 SV=2 - [H12_HUMAN]
Polymeric immunoglobulin receptor OS=Homo sapiens OX=9606 GN=PIGR PE=1 SV=4 - [PIGR_HUMAN]
Myosin regulatory light chain 12A (Fragment) OS=Homo sapiens OX=9606 GN=MYL12A PE=1 SV=1 - [J3KJTJ1_HUMAN]
HCG21173, isoform CRA_a OS=Homo sapiens OX=9606 GN=hCG_21173 PE=4 SV=1 - [D3DS95_HUMAN]
Neutrophil defensin 1 OS=Homo sapiens OX=9606 GN=DEFA1 PE=1 SV=1 - [DEF1_HUMAN]
Protein S100 (Fragment) OS=Homo sapiens OX=9606 GN=S100A6 PE=1 SV=1 - [R4GN98_HUMAN]
60S ribosomal protein L26 (Fragment) OS=Homo sapiens OX=9606 GN=RPL26 PE=4 SV=8 - [J3KJTJ8_HUMAN]
Katanin p60 ATPase-containing subunit A-like 2 (Fragment) OS=Homo sapiens OX=9606 GN=KATNAL2 PE=1 SV=1 - [K7EM02_HUMAN]
Ribosomal protein L7a (Fragment) OS=Homo sapiens OX=9606 GN=RP-L7a PE=2 SV=1 - [Q9BY74_HUMAN]
40S ribosomal protein S11 OS=Homo sapiens OX=9606 GN=RPS11 PE=1 SV=1 - [MOR1H6_HUMAN]
Bleomycin hydrolase (Fragment) OS=Homo sapiens OX=9606 GN=BLMH PE=1 SV=1 - [K7ESE8_HUMAN]
Poly(A) binding protein, cytoplasmic 1, isoform CRA_c OS=Homo sapiens OX=9606 GN=PABPC1 PE=4 SV=1 - [A0A024R9E2_HUMAN]
CLE OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q5RLJ0_HUMAN]
cDNA FLJ78614, highly similar to Homo sapiens eukaryotic translation initiation factor 4A, isoform 1 (EIF4A1), mRNA OS=Homo sapiens OX=9606 PE=2 SV=1 - [B2MUD5_HUMAN]
Proline-, glutamic acid- and leucine-rich protein 1 (Fragment) OS=Homo sapiens OX=9606 GN=PELP1 PE=1 SV=1 - [I3L1P4_HUMAN]
Fragile X mental retardation syndrome-related protein 1 (Fragment) OS=Homo sapiens OX=9606 GN=FXR1 PE=1 SV=1 - [C9JAJ4_HUMAN]
Protein S100 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B2R5H0_HUMAN]
Full-length cDNA clone CS0DD006YL02 of Neuroblastoma of Homo sapiens (human) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q86TT1_HUMAN]
Secretoglobin family 1C member 2 OS=Homo sapiens OX=9606 GN=SCGB1C2 PE=3 SV=1 - [SG1C2_HUMAN]
Keratin, type I cytoskeletal 28 OS=Homo sapiens OX=9606 GN=KRT28 PE=1 SV=2 - [K1C28_HUMAN]
Fibrinogen gamma chain, isoform CRA_a OS=Homo sapiens OX=9606 GN=FGG PE=4 SV=1 - [D3DP16_HUMAN]
RPS4X protein (Fragment) OS=Homo sapiens OX=9606 GN=RPS4X PE=2 SV=2 - [Q96IR1_HUMAN]
40S ribosomal protein S23 OS=Homo sapiens OX=9606 GN=RPS23 PE=1 SV=1 - [D6RD47_HUMAN]
Keratin, type I cuticular Ha6 OS=Homo sapiens OX=9606 GN=KRT36 PE=1 SV=1 - [KRT36_HUMAN]
IgG H chain OS=Homo sapiens OX=9606 PE=2 SV=1 - [S6BGE0_HUMAN]
Filaggrin-2 OS=Homo sapiens OX=9606 GN=FLG2 PE=1 SV=1 - [FILA2_HUMAN]
cDNA, FLJ93674 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B2R7Z6_HUMAN]
cDNA FLJ55803, highly similar to Gelsolin OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7Z4U6_HUMAN]
Protein disulfide-isomerase OS=Homo sapiens OX=9606 GN=P4HB PE=1 SV=1 - [I3L514_HUMAN]
ATP-citrate synthase (Fragment) OS=Homo sapiens OX=9606 GN=ACLY PE=1 SV=1 - [K7EIE7_HUMAN]
Peptidyl-prolyl cis-trans isomerase OS=Homo sapiens OX=9606 GN=PPIA PE=1 SV=1 - [F8WE65_HUMAN]
C-type lectin domain family 2 member B OS=Homo sapiens OX=9606 GN=CLEC2B PE=1 SV=1 - [F5H631_HUMAN]
60S ribosomal protein L8 OS=Homo sapiens OX=9606 GN=RPL8 PE=1 SV=1 - [E9PP36_HUMAN]
Alpha-lactalbumin OS=Homo sapiens OX=9606 GN=LALBA PE=1 SV=1 - [F8VWU1_HUMAN]
LMNA protein OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q8N519_HUMAN]
40S ribosomal protein S25 OS=Homo sapiens OX=9606 GN=RPS25 PE=1 SV=1 - [RS25_HUMAN]
cDNA FLJ51392, highly similar to Keratin, type II cuticular Hb5 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7Z7N3_HUMAN]
E3 ubiquitin-protein ligase KCMF1 OS=Homo sapiens OX=9606 GN=KCMF1 PE=1 SV=2 - [KCMF1_HUMAN]
Deleted in malignant brain tumors 1 protein OS=Homo sapiens OX=9606 GN=DMBT1 PE=1 SV=2 - [DMBT1_HUMAN]
Putative lipocalin 1-like protein 1 OS=Homo sapiens OX=9606 GN=LCN1P1 PE=5 SV=1 - [LC1L1_HUMAN]
40S ribosomal protein S5 OS=Homo sapiens OX=9606 GN=RPS5 PE=1 SV=1 - [M0QZN2_HUMAN]
Keratin, type I cuticular Ha3-II OS=Homo sapiens OX=9606 GN=KRT33B PE=1 SV=3 - [KT33B_HUMAN]
Histone H3.1 OS=Homo sapiens OX=9606 GN=HIST1H3A PE=1 SV=2 - [H31_HUMAN]
Alpha-2-macroglobulin OS=Homo sapiens OX=9606 GN=A2M PE=1 SV=3 - [A2MG_HUMAN]
Adenosylhomocysteinase OS=Homo sapiens OX=9606 GN=AHCY PE=2 SV=1 - [Q1RMG2_HUMAN]
L-plastin variant (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q59GX5_HUMAN]
VH6DJ protein (Fragment) OS=Homo sapiens OX=9606 GN=VH6DJ PE=2 SV=1 - [A2N0U5_HUMAN]
Cytovillin 2 (Fragment) OS=Homo sapiens OX=9606 GN=VIL2 PE=2 SV=1 - [Q9UJU1_HUMAN]

cDNA FLJ54533, highly similar to Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DP51_HUMAN]
Peptidyl-prolyl cis-trans isomerase OS=Homo sapiens OX=9606 GN=HEL-S-39 PE=2 SV=1 - [V9HWC6_HUMAN]
60S ribosomal protein L7 OS=Homo sapiens OX=9606 GN=RPL7 PE=1 SV=1 - [A8MUD9_HUMAN]
Transcobalamin-1 OS=Homo sapiens OX=9606 GN=TCN1 PE=1 SV=2 - [TCO1_HUMAN]
26S proteasome regulatory subunit 6A (Fragment) OS=Homo sapiens OX=9606 GN=PSMC3 PE=1 SV=9 - [E9PN50_HUMAN]
Plakophilin 1 (Ectodermal dysplasia/skin fragility syndrome), isoform CRA_a OS=Homo sapiens OX=9606 GN=PKP1 PE=4 SV=1 - [A0A024R9]
cDNA FLJ32530 fis, clone SMINT2000185, highly similar to TRANSKETOLASE (EC 2.2.1.1) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B3KPZ8_HUMAN]
Enolase 1 (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [A4UCS8_HUMAN]
cDNA FLJ59103, highly similar to T-complex protein 1 subunit epsilon OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DZY9_HUMAN]
cDNA FLJ35376 fis, clone SKMUS2004044, highly similar to Homo sapiens ribosomal protein L3 (RPL3), transcript variant 2, mRNA OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q6MZX7_HUMAN]
Uncharacterized protein DKFZp686M24218 OS=Homo sapiens OX=9606 GN=DKFZp686M24218 PE=2 SV=1 - [Q6MZX7_HUMAN]
Adenylyl cyclase-associated protein 1 (Fragment) OS=Homo sapiens OX=9606 GN=CAP1 PE=1 SV=1 - [Q5T0R7_HUMAN]
Heat shock protein HSP 90-alpha (Fragment) OS=Homo sapiens OX=9606 GN=HSP90AA1 PE=1 SV=1 - [G3V2J8_HUMAN]
Chaperone Hsp60, peptide-dependent ATPase, heat shock protein OS=Homo sapiens OX=9606 GN=groEL PE=4 SV=1 - [A0A2P9AGW9_HUMAN]
Coronin-1A OS=Homo sapiens OX=9606 GN=CORO1A PE=1 SV=4 - [COR1A_HUMAN]
B-cell receptor-associated protein 31 (Fragment) OS=Homo sapiens OX=9606 GN=BCAP31 PE=1 SV=1 - [C9JMD7_HUMAN]
Glucose-6-phosphate isomerase (Fragment) OS=Homo sapiens OX=9606 GN=GPI PE=1 SV=1 - [A0A0J9YXP8_HUMAN]
Heterogeneous nuclear ribonucleoproteins C1/C2 (Fragment) OS=Homo sapiens OX=9606 GN=HNRNPC PE=1 SV=1 - [G3V5V7_HUMAN]
Extracellular matrix protein 1 OS=Homo sapiens OX=9606 GN=ECM1 PE=1 SV=2 - [ECM1_HUMAN]
Calmodulin-like skin protein variant (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q53H37_HUMAN]
Prohibitin (Fragment) OS=Homo sapiens OX=9606 GN=PHB PE=1 SV=1 - [C9JZ20_HUMAN]
cDNA FLJ52524, moderately similar to Matrix metalloproteinase-9 (EC3.4.24.35) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7Z7V0_HUMAN]
Dynactin subunit 3 (Fragment) OS=Homo sapiens OX=9606 GN=DCTN3 PE=1 SV=1 - [X6RCK5_HUMAN]
Ribosomal protein L19 (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q8IWR8_HUMAN]
Ganglioside GM2 activator OS=Homo sapiens OX=9606 GN=GM2A PE=1 SV=4 - [SAP3_HUMAN]
Keratin 23 (Histone deacetylase inducible) OS=Homo sapiens OX=9606 GN=KRT23 PE=2 SV=1 - [Q8TC04_HUMAN]
Phosphoglycerate kinase OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DHM5_HUMAN]
Ribosome-binding protein 1 OS=Homo sapiens OX=9606 GN=RRBP1 PE=1 SV=1 - [A0A087WVV2_HUMAN]
SERPINB12 protein OS=Homo sapiens OX=9606 GN=SERPINB12 PE=2 SV=1 - [Q3SYB5_HUMAN]
Alpha-1-antichymotrypsin OS=Homo sapiens OX=9606 GN=SERPINA3 PE=1 SV=1 - [G3V3A0_HUMAN]
ALDH2 (Fragment) OS=Homo sapiens OX=9606 GN=ALDH2 PE=3 SV=1 - [B4YAH7_HUMAN]
cDNA FLJ51046, highly similar to 60 kDa heat shock protein, mitochondrial OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7Z5E7_HUMAN]
Vitamin D-binding protein (Fragment) OS=Homo sapiens OX=9606 GN=GC PE=1 SV=1 - [D6RF20_HUMAN]
Malate dehydrogenase OS=Homo sapiens OX=9606 PE=2 SV=1 - [B3KTM1_HUMAN]
KRT83 protein OS=Homo sapiens OX=9606 GN=KRT83 PE=2 SV=1 - [A0JNT2_HUMAN]
Neuroblast differentiation-associated protein AHNAK OS=Homo sapiens OX=9606 GN=AHNAK PE=1 SV=2 - [AHNK_HUMAN]
Catalase OS=Homo sapiens OX=9606 GN=CAT PE=1 SV=3 - [CATA_HUMAN]
Neutrophil gelatinase-associated lipocalin OS=Homo sapiens OX=9606 GN=LCN2 PE=1 SV=2 - [NGAL_HUMAN]
cDNA FLJ60780, highly similar to Carboxypeptidase A4 (EC 3.4.17.-) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B7Z577_HUMAN]
cDNA FLJ56576, highly similar to Collagen alpha-2(I) chain OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DN66_HUMAN]
40S ribosomal protein S7 OS=Homo sapiens OX=9606 GN=RPS7 PE=1 SV=1 - [B5MCP9_HUMAN]
Keratin, type I cytoskeletal 20 OS=Homo sapiens OX=9606 GN=KRT20 PE=1 SV=1 - [K1C20_HUMAN]
cDNA FLJ53009, highly similar to Calreticulin OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DHR1_HUMAN]
cDNA FLJ52062, highly similar to Erythrocyte band 7 integral membrane protein OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4E2V5_HUMAN]
Proteinase 3 OS=Homo sapiens OX=9606 GN=PRTN3 PE=2 SV=1 - [D6CHE9_HUMAN]
HNRNPA2B1 protein OS=Homo sapiens OX=9606 GN=HNRNPA2B1 PE=2 SV=1 - [I6L957_HUMAN]
F-box only protein 50 OS=Homo sapiens OX=9606 GN=NCCRP1 PE=1 SV=1 - [FBX50_HUMAN]
Elongation factor Tu, mitochondrial OS=Homo sapiens OX=9606 GN=TUFM PE=1 SV=2 - [EFTU_HUMAN]
NADH-ubiquinone oxidoreductase chain 5 OS=Homo sapiens OX=9606 GN=ND5 PE=3 SV=1 - [A0A096WP35_HUMAN]
Proteasome subunit beta type-4 OS=Homo sapiens OX=9606 GN=PSMB4 PE=1 SV=4 - [PSB4_HUMAN]
Retroviral-like aspartic protease 1 OS=Homo sapiens OX=9606 GN=ASPRV1 PE=1 SV=1 - [APRV1_HUMAN]
Desmocollin 1, isoform CRA_b OS=Homo sapiens OX=9606 GN=DSC1 PE=4 SV=1 - [Q9HB00_HUMAN]
Eukaryotic translation initiation factor 4E OS=Homo sapiens OX=9606 GN=EIF4E PE=1 SV=2 - [IF4E_HUMAN]
FN1 protein OS=Homo sapiens OX=9606 GN=FN1 PE=2 SV=1 - [B7ZLE5_HUMAN]
Elongation factor 2 OS=Homo sapiens OX=9606 GN=EEF2 PE=1 SV=4 - [EF2_HUMAN]
ATP-dependent RNA helicase DDX3X OS=Homo sapiens OX=9606 GN=DDX3X PE=1 SV=1 - [A0A0D9SFB3_HUMAN]

Acylamino-acid-releasing enzyme (Fragment) OS=Homo sapiens OX=9606 GN=APEH PE=1 SV=1 - [H7C1U0_HUMAN]
Protein POF1B OS=Homo sapiens OX=9606 GN=POF1B PE=1 SV=3 - [POF1B_HUMAN]
Collagen alpha-1(I) chain OS=Homo sapiens OX=9606 GN=COL1A1 PE=1 SV=5 - [CO1A1_HUMAN]
S protein OS=Homo sapiens OX=9606 GN=S PE=4 SV=1 - [Q0EFA5_HUMAN]
PURA protein (Fragment) OS=Homo sapiens OX=9606 GN=PURA PE=2 SV=1 - [Q2NLD4_HUMAN]
Creatine kinase U-type, mitochondrial (Fragment) OS=Homo sapiens OX=9606 GN=CKMT1B PE=1 SV=1 - [C9J6W7_HUMAN]
Tigger transposable element-derived protein 6 OS=Homo sapiens OX=9606 GN=TIGD6 PE=2 SV=2 - [TIGD6_HUMAN]
Glucose-6-phosphate 1-dehydrogenase (Fragment) OS=Homo sapiens OX=9606 GN=G6PD PE=3 SV=1 - [Q0PHS1_HUMAN]
ELAV-like protein 1 OS=Homo sapiens OX=9606 GN=ELAVL1 PE=1 SV=2 - [ELAV1_HUMAN]
PAI-1 mRNA-binding protein variant OS=Homo sapiens OX=9606 PE=2 SV=1 - [Q5VU21_HUMAN]
Truncated profilaggrin OS=Homo sapiens OX=9606 GN=FLG PE=4 SV=1 - [I0B0K8_HUMAN]
Fibrinogen beta chain OS=Homo sapiens OX=9606 GN=FGB PE=1 SV=1 - [D6REL8_HUMAN]
cDNA FLJ51265, moderately similar to Beta-2-glycoprotein 1 (Beta-2-glycoprotein I) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DPN0_HUMAN]
Ribosomal protein L5, isoform CRA_b OS=Homo sapiens OX=9606 GN=RPL5 PE=2 SV=1 - [B3KTM6_HUMAN]
Skin-specific protein 32 OS=Homo sapiens OX=9606 GN=XP32 PE=1 SV=1 - [XP32_HUMAN]
cDNA FLJ35087 fis, clone PLACE6005546, highly similar to Polymerase I and transcript release factor OS=Homo sapiens OX=9606 PE=2 SV=1 - [PLACE6005546_HUMAN]
Sulfotransferase family cytosolic 2B member 1 OS=Homo sapiens OX=9606 GN=SULT2B1 PE=1 SV=2 - [ST2B1_HUMAN]
Ubiquitin thioesterase OTU1 OS=Homo sapiens OX=9606 GN=YOD1 PE=1 SV=1 - [OTU1_HUMAN]
cDNA FLJ41699 fis, clone HCHON2004776, highly similar to Homo sapiens cytoskeleton-associated protein 4 (CKAP4), mRNA OS=Homo sapiens OX=9606 PE=2 SV=1 - [HCHON2004776_HUMAN]
Thymidine phosphorylase (Fragment) OS=Homo sapiens OX=9606 GN=TYMP PE=1 SV=1 - [C9JGI3_HUMAN]
cDNA FLJ51017, highly similar to ATP-dependent RNA helicase DDX1 (EC 3.6.1.-) (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DFI1_HUMAN]
Sulfotransferase OS=Homo sapiens OX=9606 GN=BQ8482_180419 PE=4 SV=1 - [A0A2P9AJ95_HUMAN]
cDNA FLJ51358, highly similar to Involucrin OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DWR5_HUMAN]
cDNA FLJ35968 fis, clone TESTI2013053, highly similar to INSULIN-DEGRADING ENZYME (EC 3.4.24.56) OS=Homo sapiens OX=9606 PE=2 SV=1 - [TESTI2013053_HUMAN]
E3 ubiquitin-protein ligase TRIM21 OS=Homo sapiens OX=9606 GN=TRIM21 PE=1 SV=1 - [RO52_HUMAN]
Constitutive coactivator of PPAR-gamma-like protein 1 (Fragment) OS=Homo sapiens OX=9606 GN=FAM120A PE=1 SV=1 - [A0A0C4DH52_HUMAN]
Kinesin-like protein KIF15 (Fragment) OS=Homo sapiens OX=9606 GN=KIF15 PE=1 SV=1 - [D6RCT7_HUMAN]
Serpin A12 OS=Homo sapiens OX=9606 GN=SERPINA12 PE=1 SV=1 - [SPA12_HUMAN]
Granulins (Fragment) OS=Homo sapiens OX=9606 GN=GRN PE=1 SV=1 - [K7EKL3_HUMAN]
Chaperone Hsp60, peptide-dependent ATPase, heat shock protein OS=Homo sapiens OX=9606 GN=groEL PE=4 SV=1 - [A0A2P9ALX2_HUMAN]
Synaptosomal-associated protein 47 OS=Homo sapiens OX=9606 GN=SNAP47 PE=1 SV=1 - [A0A087X0B7_HUMAN]
Insulin-like growth factor 2 mRNA-binding protein 3 OS=Homo sapiens OX=9606 GN=IGF2BP3 PE=1 SV=2 - [IF2B3_HUMAN]
Plectin OS=Homo sapiens OX=9606 GN=PLEC PE=1 SV=3 - [PLEC_HUMAN]
cDNA FLJ32635 fis, clone SYNOV2000178, highly similar to Proteoglycan-4 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B3KQ20_HUMAN]
Kinesin-like protein KIF11 OS=Homo sapiens OX=9606 GN=KIF11 PE=1 SV=2 - [KIF11_HUMAN]
cDNA FLJ57339, highly similar to Complement C3 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4E216_HUMAN]
cDNA FLJ59294, highly similar to HECT, UBA and WWE domain-containing protein 1 (EC 6.3.2.-) (Fragment) OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4E216_HUMAN]
cDNA FLJ51803, highly similar to Ryanodine receptor 1 OS=Homo sapiens OX=9606 PE=2 SV=1 - [B4DET7_HUMAN]
Desmocollin 3, isoform CRA_b OS=Homo sapiens OX=9606 GN=DSC3 PE=4 SV=1 - [A0A024RC29_HUMAN]
Putative uncharacterized protein MYH16 OS=Homo sapiens OX=9606 GN=MYH16 PE=1 SV=2 - [MYH16_HUMAN]
Periplakin (Fragment) OS=Homo sapiens OX=9606 GN=PPL PE=1 SV=1 - [K7EQ71_HUMAN]
Mucin-5B OS=Homo sapiens OX=9606 GN=MUC5B PE=1 SV=3 - [MUC5B_HUMAN]
Collagen alpha-1(III) chain OS=Homo sapiens OX=9606 GN=COL3A1 PE=1 SV=4 - [CO3A1_HUMAN]
Fatty acid synthase OS=Homo sapiens OX=9606 GN=FASN PE=1 SV=1 - [A0A0U1RQF0_HUMAN]
Filamin-A OS=Homo sapiens OX=9606 GN=FLNA PE=1 SV=1 - [A0A087WWY3_HUMAN]
Cardiomyopathy-associated protein 5 OS=Homo sapiens OX=9606 GN=CMYA5 PE=1 SV=3 - [CMYA5_HUMAN]

Σ Coverage	Σ # Proteins	Σ # Unique Peptides	Σ # Peptides	Σ # PSMs	Score R20180600747_A SMSCs	Coverage R20180600747_ ASMSCs	# Peptides R20180600747_ ASMSCs
71.36	4	38	49	211	1331.49	54.62	32
67.09	2	45	46	231	1360.13	52.49	28
67.02	12	18	31	98	666.26	43.97	21
63.16	2	6	6	16	112.21	33.33	3
61.49	19	47	54	304	1768.91	44.10	37
58.90	10	11	26	108	833.69	41.31	20
58.73	6	29	40	202	1499.37	52.23	28
48.97	14	7	39	135	563.47	33.83	21
46.23	3	1	4	12	155.86	46.23	4
44.21	13	10	19	50	242.71	21.53	10
42.92	34	18	19	55	262.71	18.24	9
42.73	8	1	33	129	538.51	28.90	20
42.73	1	5	5	12	94.93	30.00	3
42.71	13	22	37	145	713.66	35.42	22
41.04	8	1	4	8	29.61	8.21	1
40.59	2	3	3	8	57.96	10.89	1
40.00	19	5	5	12	0.00	8.97	1
38.90	72	4	10	42	252.52	30.84	8
38.83	3	4	4	11	121.76	38.83	4
38.71	1	4	4	8	62.69	31.18	3
38.57	4	3	3	4	72.01	38.57	3
38.32	1	44	44	134	413.63	17.16	13
37.67	1	4	4	9	94.88	24.66	3
36.75	14	30	30	133	674.83	32.31	24
36.50	3	1	27	89	436.15	23.38	15
34.83	5	5	5	6	90.22	34.83	5
33.61	39	2	2	6	28.25	26.23	1
33.04	7	15	15	64	254.06	28.70	9
32.56	2	2	11	22	105.21	20.00	7
31.51	29	4	6	13	247.38	23.74	3
31.36	139	1	3	4	49.72	25.42	2
31.19	10	1	2	4	151.76	31.19	2
31.01	14	4	4	11	85.55	13.92	2
30.61	2	2	2	4	53.88	30.61	2
30.26	2	0	10	19	105.21	17.44	6
30.00	3	2	2	2	29.04	16.25	1
29.63	33	3	6	19	181.87	22.69	4
29.51	4	4	5	8	59.59	11.48	2
29.34	49	4	4	6	42.32	5.79	1
29.20	27	9	9	18	65.39	10.32	3
28.81	136	1	3	4	47.65	22.88	2
28.57	15	1	3	12	117.35	19.05	2
28.24	134	1	3	5	0.00	14.50	1
27.41	58	3	8	36	163.93	24.20	7
27.14	3	3	3	5	66.05	27.14	3
26.05	84	2	3	4	40.97	9.24	1
25.85	4	4	4	7	24.77	7.80	1
25.72	5	7	7	11	330.66	19.65	5
25.24	5	1	2	5	51.27	17.48	1
25.23	5	2	2	3	56.76	16.82	1
25.17	4	7	7	13	149.07	18.79	4
25.00	10	3	6	12	29.61	3.40	1

25.00	1	1	2	2	0.00	15.38	1
24.30	1	1	2	3	67.33	24.30	2
24.21	1	2	2	3	59.36	13.68	1
23.91	3	1	1	2	41.85	23.91	1
23.91	1	7	7	10	44.33	3.42	1
23.84	19	7	11	31	181.01	9.28	4
23.32	35	2	6	9	0.00	2.88	1
23.31	8	2	8	15	74.81	12.47	4
23.25	11	3	11	34	270.10	19.00	9
22.95	8	2	2	2	41.43	13.93	1
22.60	19	35	35	46	82.99	3.42	5
22.46	149	1	4	7	145.96	14.83	2
22.45	75	1	1	1	43.96	22.45	1
22.22	4	3	4	7	46.67	6.43	1
21.95	1	1	1	8	44.55	21.95	1
21.57	17	3	3	5	43.19	8.50	1
21.48	1	4	4	5	205.14	21.48	4
21.05	23	2	2	2	47.65	21.05	2
21.00	1	7	7	12	165.09	15.49	5
20.96	9	7	7	13	140.43	12.76	4
20.95	3	3	3	4	59.71	14.86	2
20.82	20	9	9	17	168.10	12.37	5
20.59	1	2	2	2	82.55	20.59	2
20.19	3	2	2	2	76.41	20.19	2
20.00	6	1	1	2	58.87	20.00	1
20.00	16	1	2	3	0.00	10.83	1
19.73	12	12	12	25	129.04	7.25	4
19.65	4	6	10	26	253.14	10.70	6
19.49	85	1	2	3	32.63	13.56	1
19.42	3	2	2	3	37.25	19.42	2
19.40	7	5	5	9	82.48	8.96	3
19.39	12	2	2	4	40.52	12.24	1
19.23	1	1	2	2	0.00	9.62	1
19.15	21	1	1	1	53.93	19.15	1
18.95	31	3	5	8	76.12	11.69	3
18.18	2	4	4	6	33.56	8.68	2
18.03	10	4	4	5	27.76	7.14	1
17.86	10	1	1	1	46.65	17.86	1
17.84	37	6	6	20	86.42	14.07	4
17.62	46	2	4	5	32.10	4.41	1
17.47	20	6	7	27	266.93	14.95	6
17.19	17	1	5	7	0.00	2.68	1
16.87	8	5	5	7	187.08	16.87	5
16.82	4	1	1	1	52.85	16.82	1
16.81	13	6	7	22	144.09	7.30	3
16.67	2	1	1	1	37.12	16.67	1
16.24	16	10	10	13	305.02	13.51	8
15.99	93	5	7	21	237.26	10.73	4
15.89	3	2	2	3	48.03	15.89	2
15.87	21	2	2	4	23.27	8.73	1
15.52	20	6	7	8	29.91	2.46	1
15.51	42	2	4	5	24.29	3.67	1
15.48	6	1	1	3	24.07	15.48	1
15.31	7	1	1	2	32.68	15.31	1
15.22	7	1	1	1	39.74	15.22	1
15.19	4	1	1	1	35.24	15.19	1

15.09	87	3	5	7	41.43	5.17	2
14.90	11	9	9	21	66.38	4.87	3
14.77	6	1	1	1	32.35	14.77	1
14.67	14	1	2	7	45.45	9.78	1
14.53	38	1	3	4	32.10	5.59	1
14.52	8	1	1	1	20.11	14.52	1
14.45	7	42	42	73	269.31	5.75	14
14.29	12	3	3	4	43.58	4.23	1
14.17	9	3	8	22	64.26	2.33	2
14.02	3	2	10	41	204.16	8.60	5
14.00	3	8	8	10	44.33	1.30	1
14.00	5	1	1	1	35.22	14.00	1
14.00	7	2	2	3	53.19	14.00	2
13.97	2	2	2	3	76.00	13.97	2
13.85	12	1	1	2	38.17	13.85	1
13.84	8	3	3	3	59.61	13.84	3
13.49	5	5	9	35	242.92	8.65	5
13.49	6	2	2	2	64.71	13.49	2
13.33	2	2	2	5	50.71	6.67	1
13.08	47	2	2	9	51.57	12.15	1
13.04	2	1	1	1	44.10	13.04	1
12.99	4	2	2	3	62.24	12.99	2
12.92	4	2	2	2	42.16	12.92	2
12.55	1	3	3	4	87.66	12.55	3
12.54	5	7	8	11	60.69	1.68	1
12.50	2	2	2	4	22.50	7.24	1
12.50	6	1	1	1	36.76	12.50	1
12.41	6	1	1	1	0.00	12.41	1
12.38	1	1	1	3	67.75	12.38	1
12.36	5	1	1	1	31.59	12.36	1
12.36	7	9	9	12	140.24	9.18	6
12.35	3	1	1	1	39.35	12.35	1
12.25	1	1	1	1			
12.21	3	8	8	12	173.13	9.53	6
12.12	6	1	1	1	64.76	12.12	1
12.09	6	1	1	1	30.30	12.09	1
11.98	7	2	2	3	46.68	11.98	2
11.94	4	1	1	1	72.11	11.94	1
11.76	28	3	5	12	95.14	7.71	3
11.70	9	2	2	2	0.00	6.43	1
11.60	5	3	3	3	33.21	3.04	1
11.54	3	5	6	11	85.06	6.54	3
11.39	7	1	1	3	31.40	11.39	1
11.34	10	1	1	3	37.27	11.34	1
11.21	6	1	1	1	0.00	11.21	1
11.20	12	1	1	1	39.18	11.20	1
10.96	6	8	8	13	154.89	6.55	5
10.84	10	1	1	1	49.17	10.84	1
10.80	10	4	4	5	44.26	4.43	2
10.77	83	2	4	9	121.38	7.03	2
10.76	2	2	2	2	34.36	5.06	1
10.68	5	1	1	1	32.69	10.68	1
10.66	32	2	4	8	121.98	6.64	3
10.63	17	2	2	4	49.47	5.80	1
10.58	1	1	1	1	54.55	10.58	1
10.53	2	1	1	2	26.97	10.53	1

10.53	4	1	1	1	40.17	10.53	1
10.20	1	7	7	19	95.14	4.67	3
10.07	10	3	3	4	57.75	2.68	1
10.00	1	1	1	3	51.24	10.00	1
10.00	9	1	1	1	0.00	10.00	1
9.92	7	2	2	3	38.92	3.57	1
9.90	9	3	3	3	85.86	9.90	3
9.86	10	2	2	4	58.09	9.86	2
9.69	1	6	6	7	232.87	9.69	6
9.65	9	1	1	1	51.16	9.65	1
9.57	4	1	1	4	45.30	9.57	1
9.57	2	1	1	1	0.00	9.57	1
9.41	2	1	1	2	48.81	9.41	1
9.38	10	1	1	3	42.84	9.38	1
9.38	6	1	1	1	0.00	9.38	1
9.15	4	1	1	1	0.00	9.15	1
9.09	3	1	1	1	36.54	9.09	1
9.09	5	2	2	2	16.82	9.09	2
9.07	27	4	4	4	38.51	2.63	1
9.02	4	2	2	3	42.46	9.02	2
8.87	23	3	3	3	44.70	2.22	1
8.85	2	2	2	2	44.82	8.85	2
8.78	8	1	1	1	0.00	8.78	1
8.61	6	1	1	1	42.26	8.61	1
8.57	2	1	1	3	51.38	8.57	1
8.53	4	3	3	6	49.93	4.27	2
8.42	2	1	1	1	54.85	8.42	1
8.41	6	1	6	20	132.25	4.09	3
8.38	6	3	3	3	68.39	8.38	3
8.23	4	2	2	2	43.25	2.88	1
8.21	2	1	1	1	0.00	8.21	1
8.14	1	3	4	4	38.51	3.43	2
8.00	14	1	2	3	66.72	4.00	1
7.95	1	7	7	18	99.13	2.97	2
7.85	2	3	3	3	116.07	7.85	3
7.55	13	4	4	4	53.61	7.55	4
7.55	11	1	1	1	33.20	7.55	1
7.52	4	1	1	1	36.47	7.52	1
7.50	7	1	1	1	49.19	7.50	1
7.50	11	1	1	1	41.34	7.50	1
7.43	4	1	1	1	28.80	7.43	1
7.32	2	1	1	5	45.65	7.32	1
7.31	8	3	3	3	70.91	7.31	3
7.20	1	1	1	1	58.39	7.20	1
7.12	10	1	2	2	29.95	4.07	1
7.09	1	1	1	1	0.00	7.09	1
6.96	1	1	1	2	79.89	6.96	1
6.79	2	1	1	2	69.34	6.79	1
6.72	5	1	1	3	50.31	6.72	1
6.68	11	2	3	3	76.17	4.95	2
6.62	4	1	1	1	29.39	6.62	1
6.58	1	8	8	14	35.80	0.68	1
6.54	2	2	2	2	39.51	2.61	1
6.43	10	3	3	3	117.51	6.43	3
6.42	25	2	2	2	40.53	6.42	1
6.38	19	1	1	3	47.02	6.38	1

6.29	13	1	1	3	36.35	6.29	1
6.25	2	1	1	1	37.87	6.25	1
6.25	4	1	1	1	38.77	6.25	1
6.24	1	3	3	3	103.54	6.24	3
6.23	7	1	1	1	0.00	6.23	1
6.20	4	4	4	6	115.83	6.20	4
6.15	8	1	1	1	38.04	6.15	1
6.02	4	1	1	1	40.90	6.02	1
6.01	13	1	1	1	24.84	6.01	1
5.93	13	2	2	2	23.21	2.54	1
5.88	1	2	3	3	45.30	5.04	2
5.75	13	1	1	1	28.68	5.75	1
5.75	11	1	1	1	53.67	5.75	1
5.71	3	3	3	3	55.25	3.50	2
5.64	10	3	3	3	65.64	5.64	3
5.62	6	1	1	1	59.99	5.62	1
5.60	9	1	1	1	68.76	5.60	1
5.56	18	1	1	1	27.71	5.56	1
5.56	3	3	3	3	61.58	5.56	3
5.48	2	1	1	1	48.77	5.48	1
5.47	8	1	1	1	36.39	5.47	1
5.43	5	1	1	1	29.07	5.43	1
5.30	2	1	1	1	42.32	5.30	1
5.20	5	1	1	3	61.99	5.20	1
5.18	1	1	1	1	63.17	5.18	1
4.98	4	1	2	3	75.66	4.98	2
4.96	4	1	1	1	38.67	4.96	1
4.93	9	4	4	5	54.03	2.25	2
4.92	2	1	1	1	34.57	4.92	1
4.88	3	1	1	1	44.06	4.88	1
4.86	3	1	1	1	0.00	4.86	1
4.84	5	1	1	2	132.20	4.84	1
4.83	6	1	1	1	51.99	4.83	1
4.76	6	1	1	1	63.81	4.76	1
4.70	15	1	2	2	33.01	2.01	1
4.70	5	5	5	11	68.40	1.99	2
4.55	1	2	2	2	41.06	2.85	1
4.55	3	1	1	1	28.71	4.55	1
4.42	3	1	1	1	55.09	4.42	1
4.41	4	1	1	1	0.00	4.41	1
4.28	2	1	1	1	41.84	4.28	1
4.25	2	1	2	8	61.10	4.25	2
4.25	3	1	1	1	0.00	4.25	1
4.22	3	1	1	1	49.03	4.22	1
4.19	3	1	1	2	55.28	4.19	1
4.02	4	1	1	3	43.51	4.02	1
4.00	1	1	1	1	47.18	4.00	1
3.98	1	1	1	1	45.57	3.98	1
3.81	1	1	1	1	0.00	3.81	1
3.79	1	1	1	1	43.61	3.79	1
3.79	1	1	1	1	90.88	3.79	1
3.69	2	3	3	4	21.08	1.67	1
3.69	4	1	1	1	39.12	3.69	1
3.62	23	7	7	10	110.00	2.99	6
3.61	7	3	3	3	34.06	1.05	1
3.59	12	2	2	2	42.01	3.59	2

3.57	3	1	1	1	25.74	3.57	1
3.57	1	2	2	2	55.62	3.57	2
3.55	5	5	5	7	60.53	1.50	2
3.52	13	1	1	1	35.75	3.52	1
3.51	4	1	1	1	20.42	3.51	1
3.45	9	1	1	1	40.34	3.45	1
3.45	1	1	1	1	0.00	3.45	1
3.37	8	1	1	1	32.59	3.37	1
3.37	1	1	1	1	37.43	3.37	1
3.36	5	1	1	2	64.09	3.36	1
3.33	11	7	7	11	50.17	1.77	3
3.31	4	1	1	2	62.35	3.31	1
3.28	3	1	1	1	44.71	3.28	1
3.24	3	1	1	1	39.00	3.24	1
3.20	1	1	1	1	35.27	3.20	1
3.19	2	1	1	1	22.91	3.19	1
3.01	1	1	1	1	71.98	3.01	1
2.59	1	1	1	1	24.90	2.59	1
2.50	4	1	1	1	31.80	2.50	1
2.47	3	1	1	1	33.86	2.47	1
2.46	4	1	1	1	31.02	2.46	1
2.28	1	1	1	3	42.54	2.28	1
2.23	3	1	1	1	27.55	2.23	1
2.16	4	1	1	1	50.20	2.16	1
2.11	1	1	1	1			
2.02	2	1	1	1	24.04	2.02	1
1.97	2	1	1	1	25.47	1.97	1
1.93	1	1	1	1	37.11	1.93	1
1.80	5	1	1	1	41.64	1.80	1
1.67	1	1	1	3	59.62	1.67	1
1.67	3	1	1	1			
1.55	1	1	1	1	0.00	1.55	1
1.47	4	6	6	6	28.55	0.17	1
1.42	4	1	1	1	50.11	1.42	1
1.33	1	1	1	1	66.49	1.33	1
1.28	2	1	1	1	26.44	1.28	1
1.12	6	1	1	1	23.42	1.12	1
1.04	4	1	1	1	48.56	1.04	1
1.00	3	1	1	1	42.53	1.00	1
1.00	1	1	1	1	35.36	1.00	1
0.99	3	1	1	2	47.43	0.99	1
0.89	2	4	4	4	152.60	0.89	4
0.75	1	1	1	3	42.61	0.75	1
0.44	2	1	1	1	43.80	0.44	1
0.35	6	1	1	1	28.33	0.35	1
0.17	1	1	1	1	0.00	0.17	1

# PSM R20180600747_A SMSCs	Score R20180600747_I GG	Coverage R20180600747_I GG	# Peptides R20180600747_I GG	# PSM R20180600747_I GG	# AAs	MW [kDa]	calc. pI
46	864.66	37.87	22	27	639	65.4	8.00
43	1096.41	52.33	27	34	623	62.0	5.24
29	271.81	29.60	11	14	473	51.2	5.05
4					114	13.2	6.13
58	1489.81	44.57	40	54	644	66.0	8.12
30	456.28	30.72	13	17	472	51.5	5.16
45	769.09	39.73	19	24	584	58.8	5.21
28	554.66	31.21	19	22	535	57.8	8.00
5	56.34	18.87	1	2	106	11.2	8.29
11	105.44	11.57	4	4	432	48.1	5.02
10	301.26	19.31	9	9	466	53.6	5.12
27	502.10	26.06	17	20	564	60.0	8.00
3	65.24	22.73	2	2	110	11.3	6.54
28	575.39	26.44	19	22	590	62.3	7.74
1					134	14.8	9.54
1					101	11.5	6.77
1	67.46	15.17	2	2	145	16.3	4.65
10	214.15	23.92	6	8	347	38.6	5.35
5					103	11.4	11.36
3	41.59	7.53	1	1	93	10.8	7.03
3					70	8.2	8.56
14	288.23	19.89	15	17	2850	282.2	10.04
3	60.36	8.22	1	1	146	16.6	8.05
27	536.33	20.51	15	22	585	66.5	6.04
19	365.26	21.48	11	13	526	55.7	8.43
6					178	19.6	5.95
1	55.60	26.23	1	1	122	13.7	6.14
12					345	39.0	8.48
8					390	44.5	6.81
4	124.93	14.16	2	2	219	24.0	8.06
2					118	12.9	6.58
2	47.40	14.68	1	1	109	11.6	8.29
2					158	17.4	9.50
2	50.18	18.37	1	1	98	11.0	5.50
7					390	44.7	6.64
1					80	9.3	9.72
4	66.20	17.13	2	3	216	22.9	8.31
2	38.87	6.01	1	1	183	20.1	8.78
1					242	27.3	4.50
3					339	38.6	7.75
2					118	13.3	7.12
2	56.34	19.05	1	2	105	11.3	7.87
1					131	14.4	7.96
8	96.70	13.99	4	5	343	38.6	5.38
3					140	15.0	8.27
1					119	13.0	8.62
1					205	22.8	6.40
5					346	38.7	7.02
2	45.97	17.48	1	1	103	11.5	7.58
1					107	11.6	8.51
4	49.03	3.36	1	1	298	34.2	6.05
1					324	35.9	9.88

1	57.45	9.62	1	1	104	11.2	6.15
2					107	10.0	8.44
1					95	10.4	4.88
1					46	5.4	9.61
1	33.18	3.42	1	1	322	34.7	7.21
5	138.27	9.28	4	5	474	51.7	6.81
1					416	46.2	5.12
5					369	42.3	6.05
12	70.50	5.00	2	2	400	44.1	5.14
1					122	13.9	8.72
5	54.73	1.33	2	2	1960	226.4	5.60
2	47.40	6.78	1	1	236	25.1	8.12
1					49	5.9	4.82
1	38.87	6.43	1	1	171	19.0	6.92
2	33.80	21.95	1	2	41	5.0	8.05
1					153	16.7	9.33
5					256	26.7	5.76
2					76	8.4	7.14
5					381	41.9	5.01
4					439	48.5	5.92
2					148	16.7	10.51
5					485	53.0	6.84
2					136	15.1	7.62
2					104	11.5	9.07
1					50	5.6	9.63
1	112.59	20.00	2	2	120	13.1	7.12
4	77.94	5.91	3	4	745	81.7	6.14
9	86.32	2.62	2	2	458	49.6	4.96
1					118	13.1	6.07
2					103	11.6	4.92
3	50.25	2.39	1	1	335	36.0	8.46
1					98	10.5	9.17
1	57.45	9.62	1	1	104	11.3	8.37
1					47	5.0	11.90
3					248	27.8	4.74
2					242	27.7	5.58
1					294	32.6	4.78
1					56	6.5	10.35
4	83.66	5.78	2	2	398	43.0	8.81
1					227	26.4	4.78
8	75.41	13.05	5	5	475	52.3	8.57
1					448	49.9	5.06
5					332	36.7	8.27
1					107	12.1	9.85
4	93.08	2.43	1	2	452	50.5	5.67
1					108	11.8	8.68
8					696	76.6	8.02
5	133.40	8.50	3	4	494	53.1	6.86
2					151	16.3	10.05
1	31.94	8.73	1	1	126	14.0	10.32
1					406	45.2	6.58
1					245	27.7	4.79
1	55.55	15.48	1	1	84	9.2	10.27
1					98	10.6	5.45
1					92	9.9	5.58
1					79	9.0	11.53

2	27.80	3.66	1	2	464	51.1	7.71
3	62.13	1.47	2	2	678	74.8	7.12
1					88	9.8	4.59
1	95.22	14.67	2	2	184	20.4	5.52
1					179	20.6	4.61
1					62	6.8	9.70
14	174.65	2.61	7	7	2871	331.6	6.81
1					189	20.4	8.44
2	81.25	5.00	2	4	600	64.8	7.56
8	165.80	8.60	4	6	535	57.8	7.20
1	55.80	2.89	1	1	693	76.6	5.86
1					100	11.5	10.15
2	17.27	6.00	1	1	250	28.8	9.44
3					136	15.5	10.07
1					65	7.3	9.38
3					477	50.6	8.47
8	122.41	3.46	2	3	578	61.8	5.85
2					126	14.2	10.46
1					135	15.2	7.01
1	69.84	12.15	1	1	107	11.9	8.37
1					92	9.1	4.46
2					177	20.1	9.60
2					178	20.5	6.02
3					255	28.8	11.19
1					654	72.3	5.16
1	33.90	7.24	1	1	152	17.7	10.99
1					80	9.2	7.97
1					137	15.5	8.57
1					105	11.7	4.92
1					89	9.7	10.36
6					817	89.7	6.04
1					81	8.9	10.14
	0.00	12.25	1	1	204	22.4	7.58
6					745	83.8	8.97
1					66	7.8	5.31
1					91	10.1	10.21
2					167	17.7	5.71
1					134	14.9	7.50
3					493	53.5	5.86
1					171	19.0	11.72
1					362	38.2	5.55
3	43.89	4.23	2	2	520	56.8	6.02
1					79	9.0	10.89
1					97	10.5	9.17
1					116	12.6	9.41
1					125	13.9	5.31
5					748	86.1	7.96
1					83	8.8	4.42
2					361	39.3	8.40
2	30.27	2.20	1	1	455	48.9	6.81
1					158	17.8	6.02
1					103	11.7	10.33
4					572	62.4	5.69
1					207	24.4	5.59
1					104	11.7	5.24
1					76	8.4	6.73

1					114	12.7	7.96
3	73.70	2.19	2	2	1049	113.7	5.03
1					298	32.8	9.74
1					90	9.9	8.25
1					130	14.5	11.75
1					252	29.2	10.65
3					303	34.5	7.71
2					213	21.4	10.93
7					764	83.2	5.74
1					114	13.0	4.88
1					94	10.1	10.07
1					94	10.2	6.99
1					85	9.7	5.45
1					96	11.5	10.90
1					128	14.0	7.34
1					142	15.6	10.04
1					88	10.1	6.93
2					231	26.7	6.79
1					419	47.3	8.92
2					244	28.0	6.44
1					406	46.1	5.48
2					192	20.6	8.90
1					148	15.1	10.68
1					151	17.9	4.98
1					105	11.7	7.18
2					375	41.2	6.79
1					95	10.5	4.81
4	93.24	1.94	1	2	464	50.5	5.47
3					334	37.7	6.29
1					243	27.2	9.94
1					134	14.8	10.23
2					467	52.2	4.94
1	30.71	4.00	1	1	300	32.2	7.85
2	93.89	5.06	3	3	2391	247.9	8.31
3					484	52.5	7.55
4					689	75.7	5.81
1					106	11.6	9.00
1					133	14.2	9.86
1					120	13.0	6.77
1					80	9.2	9.94
1					148	16.2	11.90
1	32.02	7.32	1	1	123	14.0	4.78
3					465	53.2	6.40
1					125	13.7	10.11
1					295	32.4	5.48
1					381	41.9	5.66
1					2413	260.6	5.44
1	69.76	6.79	1	1	162	17.9	5.00
1	38.69	6.72	1	1	134	14.8	8.48
2					404	46.2	4.84
1					136	15.4	11.12
1	31.60	1.22	1	1	1474	163.2	6.46
1					306	33.8	6.61
3					498	55.7	5.34
1					109	12.2	5.36
1					141	16.2	9.31

1					159	16.8	5.19	
1					208	22.7	9.32	
1					208	24.4	10.67	
3					433	48.2	5.03	
1					257	28.6	5.05	
4	35.24	1.10		1	1	726	80.4	8.97
1					358	38.9	7.72	
1					166	17.8	8.88	
1					283	32.0	8.50	
1					354	40.1	10.30	
2					476	52.4	7.77	
1					174	19.2	7.05	
1					174	20.1	4.63	
2					543	58.0	5.22	
3					461	51.0	6.68	
1					178	20.2	9.13	
1					268	30.4	9.01	
1					180	20.2	4.70	
3					540	60.6	6.71	
1					146	15.9	4.44	
1					201	22.3	5.96	
1					276	31.1	7.14	
1					151	17.3	6.58	
1					173	20.8	11.44	
1					193	20.8	5.31	
2					422	48.1	6.54	
1					242	25.7	5.78	
2					934	102.7	5.34	
1					183	20.9	5.87	
1					205	23.4	5.64	
1					247	26.6	5.90	
2					517	55.0	5.60	
1					145	16.1	5.83	
1					231	24.6	7.99	
1					447	49.6	5.39	
3					5890	628.7	6.15	
1					527	59.7	7.39	
1					198	22.6	8.91	
1					226	25.2	6.39	
1					204	19.8	10.10	
1					187	21.3	10.27	
2					424	48.5	5.69	
1					212	24.3	5.11	
1					237	25.9	7.39	
1					215	23.6	8.16	
1					249	28.4	4.86	
1					275	30.8	6.62	
1					452	49.5	7.61	
1					603	67.0	9.03	
1					264	29.2	5.97	
1					343	37.0	5.44	
1					840	93.8	5.53	
1					217	25.1	6.15	
6	30.95	1.29		2	2	2240	246.5	6.06
1					858	95.3	6.83	
2					640	70.8	7.36	

1					308	34.6	6.54
2					589	68.0	6.32
2					1464	138.9	5.80
1					512	49.9	8.13
1					285	32.0	7.05
1					203	22.5	9.22
1					521	58.6	6.80
1					297	33.9	5.90
1					326	36.1	9.17
1					387	42.4	8.44
3	34.25	0.20	1	1	4021	430.2	9.29
1					272	31.2	7.25
1					274	30.3	7.85
1					247	28.0	9.13
1					250	26.2	7.97
1					345	39.0	6.51
1					365	41.3	5.40
1					348	38.3	6.11
1					521	58.1	5.25
1					445	46.1	5.52
1					529	59.2	8.25
1	54.21	2.28	1	1	527	58.5	7.37
1					449	52.5	4.49
1					464	54.2	7.72
	40.76	2.11	1	1	475	54.1	6.38
1					495	54.0	7.52
1					711	82.4	6.48
1					414	47.1	9.29
1					388	41.3	6.52
1					539	56.8	5.39
	48.01	1.67	1	1	419	47.3	6.96
1					579	63.7	8.87
1					4684	531.5	5.96
1					494	55.5	9.92
1					1056	119.1	5.64
1					1098	122.5	6.93
1					1343	148.8	5.53
1					675	76.6	5.68
1					896	99.9	6.10
1					1097	128.2	5.49
2					1115	131.3	5.58
4					5762	596.0	6.64
1					1466	138.5	6.61
1					2509	273.0	6.47
1					2315	245.7	5.97
1					4069	448.9	4.78