| ID_1 | Name_1 | ID_2 | Name_2 | FDR | Count in human | Average count in random |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00001 | 7tm_1 | PF00001 | 7tm_1 | 0 | 14 | 2.21 |
| PF00001 | 7tm_1 | PF00048 | IL8 | 0 | 37 | 1.25 |
| PF00001 | 7tm_1 | PF00339 | Arrestin_N | 0 | 9 | 0.61 |
| PF00001 | 7tm_1 | PF00503 | G-alpha | 0 | 63 | 2.36 |
| PF00001 | 7tm_1 | PF00615 | RGS | 0 | 10 | 1.55 |
| PF00001 | 7tm_1 | PF03002 | Somatostatin | 0 | 12 | 0.18 |
| PF00008 | EGF | PF00008 | EGF | 0 | 22 | 3.65 |
| PF00008 | EGF | PF00040 | fn2 | 0 | 9 | 1.36 |
| PF00008 | EGF | PF00041 | fn3 | 0 | 20 | 7.06 |
| PF00008 | EGF | PF00047 | ig | 0 | 16 | 5.79 |
| PF00008 | EGF | PF00053 | Laminin_EGF | 0 | 8 | 0.84 |
| PF00008 | EGF | PF00079 | Serpin | 0 | 9 | 1.92 |
| PF00008 | EGF | PF00089 | Trypsin | 0 | 14 | 3.32 |
| PF00008 | EGF | PF00147 | Fibrinogen_C | 0 | 8 | 0.69 |
| PF00008 | EGF | PF00413 | Peptidase_M10 | 0 | 18 | 1.08 |
| PF00008 | EGF | PF00640 | PID | 0 | 16 | 2.55 |
| PF00008 | EGF | PF01391 | Collagen | 0 | 19 | 2.85 |
| PF00008 | EGF | PF01821 | ANATO | 0 | 9 | 0.39 |
| PF00008 | EGF | PF02210 | Laminin_G_2 | 0 | 10 | 1.25 |
| PF00008 | EGF | PF07645 | EGF_CA | 0 | 24 | 3.61 |
| PF00008 | EGF | PF07679 | I-set | 0 | 12 | 3.33 |
| PF00008 | EGF | PF07974 | EGF_2 | 0 | 14 | 2.32 |
| PF00010 | HLH | PF00010 | HLH | 0 | 34 | 1.09 |
| PF00010 | HLH | PF00104 | Hormone_recep | 0 | 25 | 3.58 |
| PF00010 | HLH | PF00439 | Bromodomain | 0 | 9 | 1.75 |
| PF00012 | HSP70 | PF00515 | TPR_1 | 0 | 7 | 0.34 |
| PF00013 | KH_1 | PF00013 | KH_1 | 0 | 6 | 0.1 |
| PF00013 | KH_1 | PF00076 | RRM_1 | 0 | 13 | 1.14 |
| PF00017 | SH2 | PB001482 | Pfam-B_1482 | 0 | 8 | 0.43 |
| PF00017 | SH2 | PB023385 | Pfam-B_23385 | 0 | 6 | 0.28 |
| PF00017 | SH2 | PF00017 | SH2 | 0 | 79 | 17.08 |
| PF00017 | SH2 | PF00018 | SH3_1 | 0 | 90 | 29.23 |
| PF00017 | SH2 | PF00041 | fn3 | 0 | 73 | 15.78 |
| PF00017 | SH2 | PF00047 | ig | 0 | 62 | 12.8 |
| PF00017 | SH2 | PF00097 | zf-C3HC4 | 0 | 25 | 13.47 |
| PF00017 | SH2 | PF00102 | Y_phosphatase | 0 | 28 | 6.08 |
| PF00017 | SH2 | PF00130 | C1_1 | 0 | 20 | 9.26 |
| PF00017 | SH2 | PF00169 | PH | 0 | 49 | 14.92 |
| PF00017 | SH2 | PF00620 | RhoGAP | 0 | 16 | 4.36 |
| PF00017 | SH2 | PF00621 | RhoGEF | 0 | 17 | 5.59 |
| PF00017 | SH2 | PF00627 | UBA | 0 | 18 | 3.76 |
| PF00017 | SH2 | PF02174 | IRS | 0 | 13 | 1.82 |
| PF00017 | SH2 | PF02189 | ITAM | 0 | 9 | 1.28 |
| PF00017 | SH2 | PF07686 | V-set | 0 | 28 | 7.38 |
| PF00017 | SH2 | PF07714 | Pkinase_Tyr | 0 | 168 | 28.13 |
| PF00018 | SH3_1 | PF00018 | SH3_1 | 0 | 47 | 16.75 |
| PF00018 | SH3_1 | PF00041 | fn3 | 0 | 40 | 16.02 |
| PF00018 | SH3_1 | PF00047 | ig | 0 | 40 | 12.4 |
| PF00018 | SH3_1 | PF00097 | zf-C3HC4 | 0 | 24 | 13.38 |
| PF00018 | SH3_1 | PF00102 | Y_phosphatase | 0 | 18 | 6.14 |
| PF00018 | SH3_1 | PF00169 | PH | 0 | 42 | 14.85 |


| PF00018 | SH3_1 | PF00350 | Dynamin_N | 0 | 8 | 0.61 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00018 | SH3_1 | PF00412 | LIM | 0 | 16 | 7.34 |
| PF00018 | SH3_1 | PF00568 | WH1 | 0 | 15 | 1.85 |
| PF00018 | SH3_1 | PF00617 | RasGEF | 0 | 12 | 1.54 |
| PF00018 | SH3_1 | PF00620 | RhoGAP | 0 | 12 | 4.27 |
| PF00018 | SH3_1 | PF00621 | RhoGEF | 0 | 14 | 5.53 |
| PF00018 | SH3_1 | PF00627 | UBA | 0 | 21 | 3.79 |
| PF00018 | SH3_1 | PF00786 | PBD | 0 | 14 | 1.69 |
| PF00018 | SH3_1 | PF02189 | ITAM | 0 | 8 | 1.32 |
| PF00018 | SH3_1 | PF02205 | WH2 | 0 | 16 | 1.15 |
| PF00018 | SH3_1 | PF07686 | V-set | 0 | 16 | 7.35 |
| PF00018 | SH3_1 | PF07714 | Pkinase_Tyr | 0 | 113 | 29.19 |
| PF00019 | TGF_beta | PF00019 | TGF_beta | 0 | 7 | 0.21 |
| PF00019 | TGF_beta | PF00069 | Pkinase | 0 | 42 | 5.31 |
| PF00019 | TGF_beta | PF00100 | Zona_pellucida | 0 | 7 | 0.14 |
| PF00020 | TNFR_c6 | PF00097 | zf-C3HC4 | 0 | 13 | 1.32 |
| PF00020 | TNFR_c6 | PF00229 | TNF | 0 | 17 | 0.22 |
| PF00020 | TNFR_c6 | PF00917 | MATH | 0 | 14 | 0.3 |
| PF00022 | Actin | PF00241 | Cofilin_ADF | 0 | 9 | 0.12 |
| PF00022 | Actin | PF00435 | Spectrin | 0 | 7 | 0.51 |
| PF00023 | Ank | PF00023 | Ank | 0 | 9 | 1.03 |
| PF00023 | Ank | PF00069 | Pkinase | 0 | 22 | 10.88 |
| PF00023 | Ank | PF00169 | PH | 0 | 12 | 4.25 |
| PF00023 | Ank | PF01833 | TIG | 0 | 12 | 1.1 |
| PF00023 | Ank | PF07679 | I-set | 0 | 10 | 1.92 |
| PF00025 | Arf | PF06456 | Arfaptin | 0 | 6 | 0.05 |
| PF00028 | Cadherin | PF00514 | Arm | 0 | 15 | 0.7 |
| PF00031 | Cystatin | PF00112 | Peptidase_C1 | 0 | 7 | 0.02 |
| PF00036 | efhand | PF00036 | efhand | 0 | 20 | 1.82 |
| PF00036 | efhand | PF00520 | Ion_trans | 0 | 12 | 1.78 |
| PF00036 | efhand | PF00622 | SPRY | 0 | 7 | 0.77 |
| PF00036 | efhand | PF01023 | S_100 | 0 | 8 | 0.54 |
| PF00038 | Filament | PF00038 | Filament | 0 | 16 | 0.56 |
| PF00038 | Filament | PF00244 | 14-3-3 | 0 | 7 | 0.82 |
| PF00038 | Filament | PF00681 | Plectin | 0 | 7 | 0.25 |
| PF00040 | fn2 | PF01391 | Collagen | 0 | 9 | 0.48 |
| PF00041 | fn3 | PF00041 | fn3 | 0 | 20 | 3.6 |
| PF00041 | fn3 | PF00047 | ig | 0 | 21 | 6.05 |
| PF00041 | fn3 | PF00102 | Y_phosphatase | 0 | 19 | 2.62 |
| PF00041 | fn3 | PF00169 | PH | 0 | 23 | 7.59 |
| PF00041 | fn3 | PF00812 | Ephrin | 0 | 9 | 0.37 |
| PF00041 | fn3 | PF02174 | IRS | 0 | 8 | 0.88 |
| PF00041 | fn3 | PF07679 | I-set | 0 | 15 | 3.3 |
| PF00041 | fn3 | PF07686 | V-set | 0 | 11 | 3.28 |
| PF00041 | fn3 | PF07714 | Pkinase_Tyr | 0 | 33 | 13.98 |
| PF00045 | Hemopexin | PF00965 | TIMP | 0 | 8 | 0.06 |
| PF00045 | Hemopexin | PF07686 | V-set | 0 | 7 | 0.6 |
| PF00046 | Homeobox | PF00046 | Homeobox | 0 | 28 | 1.23 |
| PF00046 | Homeobox | PF00104 | Hormone_recep | 0 | 15 | 3.31 |
| PF00046 | Homeobox | PF00292 | PAX | 0 | 6 | 0.23 |
| PF00047 | ig | PF00047 | ig | 0 | 15 | 2.5 |
| PF00047 | ig | PF00102 | Y_phosphatase | 0 | 12 | 2.29 |
| PF00047 | ig | PF00167 | FGF | 0 | 9 | 0.35 |


| PF00047 | ig | PF00640 | PID | 0 | 9 | 1.92 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00047 | ig | PF07645 | EGF_CA | 0 | 11 | 3.18 |
| PF00047 | ig | PF07679 | I-set | 0 | 13 | 2.55 |
| PF00047 | ig | PF07686 | V-set | 0 | 15 | 2.83 |
| PF00047 | ig | PF07714 | Pkinase_Tyr | 0 | 30 | 11.29 |
| PF00048 | IL8 | PF07686 | V-set | 0 | 7 | 0.71 |
| PF00059 | Lectin_C | PF00059 | Lectin_C | 0 | 8 | 0.22 |
| PF00059 | Lectin_C | PF00084 | Sushi | 0 | 7 | 0.54 |
| PF00060 | Lig_chan | PF00595 | PDZ | 0 | 21 | 1.28 |
| PF00069 | Pkinase | PF00069 | Pkinase | 0 | 98 | 24.58 |
| PF00069 | Pkinase | PF00130 | C1_1 | 0 | 34 | 10.61 |
| PF00069 | Pkinase | PF00134 | Cyclin_N | 0 | 17 | 4.74 |
| PF00069 | Pkinase | PF00169 | PH | 0 | 31 | 19.32 |
| PF00069 | Pkinase | PF00244 | 14-3-3 | 0 | 20 | 5.85 |
| PF00069 | Pkinase | PF00531 | Death | 0 | 20 | 7.78 |
| PF00069 | Pkinase | PF00564 | PB1 | 0 | 16 | 3.28 |
| PF00069 | Pkinase | PF00581 | Rhodanese | 0 | 15 | 1.93 |
| PF00069 | Pkinase | PF00917 | MATH | 0 | 13 | 4.3 |
| PF00069 | Pkinase | PF01412 | ArfGap | 0 | 10 | 1.83 |
| PF00071 | Ras | PB000002 | Pfam-B_2 | 0 | 21 | 3.72 |
| PF00071 | Ras | PF00130 | C1_1 | 0 | 13 | 3.46 |
| PF00071 | Ras | PF00168 | C2 | 0 | 16 | 4.17 |
| PF00071 | Ras | PF00169 | PH | 0 | 25 | 6.4 |
| PF00071 | Ras | PF00595 | PDZ | 0 | 19 | 8.08 |
| PF00071 | Ras | PF00617 | RasGEF | 0 | 12 | 0.58 |
| PF00071 | Ras | PF00620 | RhoGAP | 0 | 11 | 1.74 |
| PF00071 | Ras | PF00621 | RhoGEF | 0 | 14 | 1.97 |
| PF00071 | Ras | PF00786 | PBD | 0 | 12 | 0.72 |
| PF00071 | Ras | PF00788 | RA | 0 | 26 | 1.2 |
| PF00071 | Ras | PF01843 | DIL | 0 | 12 | 0.59 |
| PF00071 | Ras | PF02115 | Rho_GDI | 0 | 8 | 0.17 |
| PF00071 | Ras | PF02204 | VPS9 | 0 | 7 | 0.13 |
| PF00071 | Ras | PF06920 | Ded_cyto | 0 | 6 | 0.2 |
| PF00076 | RRM_1 | PF00076 | RRM_1 | 0 | 40 | 1.77 |
| PF00076 | RRM_1 | PF00397 | WW | 0 | 9 | 1.64 |
| PF00079 | Serpin | PF00089 | Trypsin | 0 | 47 | 0.79 |
| PF00084 | Sushi | PF00084 | Sushi | 0 | 6 | 0.27 |
| PF00084 | Sushi | PF00147 | Fibrinogen_C | 0 | 6 | 0.25 |
| PF00084 | Sushi | PF01821 | ANATO | 0 | 9 | 0.18 |
| PF00089 | Trypsin | PF00089 | Trypsin | 0 | 7 | 0.7 |
| PF00089 | Trypsin | PF01391 | Collagen | 0 | 10 | 1.5 |
| PF00092 | VWA | PF01391 | Collagen | 0 | 9 | 0.51 |
| PF00096 | zf-C2H2 | PF00096 | zf-C2H2 | 0 | 28 | 2.1 |
| PF00096 | zf-C2H2 | PF00097 | zf-C3HC4 | 0 | 14 | 4.35 |
| PF00096 | zf-C2H2 | PF00643 | zf-B_box | 0 | 11 | 1.44 |
| PF00096 | zf-C2H2 | PF00651 | BTB | 0 | 11 | 1.03 |
| PF00096 | zf-C2H2 | PF02023 | SCAN | 0 | 20 | 0.78 |
| PF00096 | zf-C2H2 | PF03165 | MH1 | 0 | 10 | 1.14 |
| PF00097 | zf-C3HC4 | PF00179 | UQ_con | 0 | 18 | 1.08 |
| PF00097 | zf-C3HC4 | PF00888 | Cullin | 0 | 6 | 0.28 |
| PF00097 | zf-C3HC4 | PF01454 | MAGE | 0 | 8 | 0.92 |
| PF00102 | Y_phosphatase | PF00169 | PH | 0 | 12 | 2.61 |
| PF00102 | Y_phosphatase | PF07714 | Pkinase_Tyr | 0 | 28 | 5.25 |


| PF00104 | Hormone_recep | PF00104 | Hormone_recep | 0 | 31 | 2.19 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00104 | Hormone_recep | PF00249 | Myb_DNA-binding | 0 | 9 | 1.42 |
| PF00104 | Hormone_recep | PF00439 | Bromodomain | 0 | 17 | 1.97 |
| PF00104 | Hormone_recep | PF00628 | PHD | 0 | 9 | 0.96 |
| PF00104 | Hormone_recep | PF00989 | PAS | 0 | 32 | 1.92 |
| PF00110 | wnt | PF01392 | Fz | 0 | 7 | 0 |
| PF00128 | Alpha-amylase | PF00324 | AA_permease | 0 | 6 | 0 |
| PF00130 | C1_1 | PF00169 | PH | 0 | 19 | 4.24 |
| PF00130 | C1_1 | PF00244 | 14-3-3 | 0 | 11 | 1.22 |
| PF00130 | C1_1 | PF07714 | Pkinase_Tyr | 0 | 18 | 7.95 |
| PF00168 | C2 | PF00169 | PH | 0 | 15 | 4.71 |
| PF00168 | C2 | PF07714 | Pkinase_Tyr | 0 | 23 | 9.8 |
| PF00169 | PH | PF00169 | PH | 0 | 12 | 3.5 |
| PF00169 | PH | PF01412 | ArfGap | 0 | 7 | 0.63 |
| PF00169 | PH | PF07714 | Pkinase_Tyr | 0 | 60 | 13.49 |
| PF00170 | bZIP_1 | PB001741 | Pfam-B_1741 | 0 | 8 | 0.16 |
| PF00170 | bZIP_1 | PF00170 | bZIP_1 | 0 | 8 | 0.21 |
| PF00170 | bZIP_1 | PF07716 | bZIP_2 | 0 | 16 | 0.36 |
| PF00179 | UQ_con | PF00632 | HECT | 0 | 8 | 0.19 |
| PF00179 | UQ_con | PF01485 | IBR | 0 | 8 | 0.07 |
| PF00193 | Xlink | PF00413 | Peptidase_M10 | 0 | 8 | 0.16 |
| PF00307 | CH | PF00412 | LIM | 0 | 12 | 1.67 |
| PF00341 | PDGF | PF07679 | I-set | 0 | 7 | 0.19 |
| PF00373 | Band_41 | PB000002 | Pfam-B_2 | 0 | 8 | 1.14 |
| PF00373 | Band_41 | PF00595 | PDZ | 0 | 10 | 2.14 |
| PF00400 | WD40 | PF00631 | G-gamma | 0 | 15 | 0.24 |
| PF00452 | Bcl-2 | PF00452 | Bcl-2 | 0 | 13 | 0.16 |
| PF00503 | G-alpha | PF00615 | RGS | 0 | 24 | 0.8 |
| PF00503 | G-alpha | PF02188 | GoLoco | 0 | 8 | 0.21 |
| PF00520 | Ion_trans | PF00520 | Ion_trans | 0 | 30 | 0.46 |
| PF00531 | Death | PF00531 | Death | 0 | 12 | 0.38 |
| PF00531 | Death | PF00656 | Peptidase_C14 | 0 | 8 | 0.49 |
| PF00531 | Death | PF00917 | MATH | 0 | 8 | 0.63 |
| PF00531 | Death | PF01335 | DED | 0 | 13 | 0.31 |
| PF00531 | Death | PF01582 | TIR | 0 | 7 | 0.22 |
| PF00560 | LRR_1 | PF00560 | LRR_1 | 0 | 7 | 0.77 |
| PF00560 | LRR_1 | PF00619 | CARD | 0 | 7 | 0.8 |
| PF00564 | PB1 | PF00564 | PB1 | 0 | 7 | 0.05 |
| PF00595 | PDZ | PF00595 | PDZ | 0 | 16 | 5.38 |
| PF00619 | CARD | PF00619 | CARD | 0 | 14 | 0.18 |
| PF00619 | CARD | PF00656 | Peptidase_C14 | 0 | 17 | 0.48 |
| PF00619 | CARD | PF05729 | NACHT | 0 | 7 | 0.03 |
| PF00620 | RhoGAP | PF07714 | Pkinase_Tyr | 0 | 19 | 4.25 |
| PF00621 | RhoGEF | PF07714 | Pkinase_Tyr | 0 | 14 | 5.09 |
| PF00627 | UBA | PF07714 | Pkinase_Tyr | 0 | 13 | 3.18 |
| PF00640 | PID | PF02177 | A4_EXTRA | 0 | 9 | 0.2 |
| PF00640 | PID | PF07714 | Pkinase_Tyr | 0 | 15 | 4.66 |
| PF00653 | BIR | PF00656 | Peptidase_C14 | 0 | 7 | 0.22 |
| PF00656 | Peptidase_C14 | PF00656 | Peptidase_C14 | 0 | 6 | 0.23 |
| PF00656 | Peptidase_C14 | PF01335 | DED | 0 | 9 | 0.23 |
| PF00735 | GTP_CDC | PF00735 | GTP_CDC | 0 | 6 | 0.01 |
| PF00754 | F5_F8_type_C | PF01391 | Collagen | 0 | 6 | 0.2 |
| PF00787 | PX | PF07714 | Pkinase_Tyr | 0 | 10 | 2.1 |


| PF00788 | RA | PF07714 | Pkinase_Tyr | 0 | 15 | 2.69 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00822 | PMP22_Claudin | PF07653 | SH3_2 | 0 | 6 | 0.12 |
| PF00850 | Hist_deacetyl | PF00850 | Hist_deacetyl | 0 | 7 | 0.19 |
| PF00957 | Synaptobrevin | PF05739 | SNARE | 0 | 12 | 0.15 |
| PF00989 | PAS | PB000064 | Pfam-B_64 | 0 | 8 | 0.11 |
| PF00989 | PAS | PB003013 | Pfam-B_3013 | 0 | 7 | 0.06 |
| PF00989 | PAS | PF00989 | PAS | 0 | 14 | 0.3 |
| PF01007 | IRK | PF00595 | PDZ | 0 | 11 | 0.4 |
| PF01034 | Syndecan | PF00595 | PDZ | 0 | 7 | 0.52 |
| PF01094 | ANF_receptor | PF00595 | PDZ | 0 | 20 | 1.62 |
| PF01094 | ANF_receptor | PF01094 | ANF_receptor | 0 | 6 | 0.16 |
| PF01217 | Clat_adaptor_s | PF01602 | Adaptin_N | 0 | 7 | 0.01 |
| PF01335 | DED | PF01335 | DED | 0 | 8 | 0.06 |
| PF01391 | Collagen | PF01391 | Collagen | 0 | 7 | 0.67 |
| PF01391 | Collagen | PF01462 | LRRNT | 0 | 11 | 0.54 |
| PF01391 | Collagen | PF07645 | EGF_CA | 0 | 13 | 1.56 |
| PF01423 | LSM | PF01423 | LSM | 0 | 20 | 0.19 |
| PF01839 | FG-GAP | PF03921 | ICAM_N | 0 | 7 | 0.14 |
| PF01839 | FG-GAP | PF07974 | EGF_2 | 0 | 16 | 0.65 |
| PF02071 | NSF | PF05739 | SNARE | 0 | 8 | 0.09 |
| PF02174 | IRS | PF07714 | Pkinase_Tyr | 0 | 15 | 1.47 |
| PF02197 | RIIa | PF05716 | AKAP_110 | 0 | 7 | 0.03 |
| PF02210 | Laminin_G_2 | PF07714 | Pkinase_Tyr | 0 | 10 | 2.33 |
| PF02319 | E2F_TDP | PF02319 | E2F_TDP | 0 | 7 | 0.03 |
| PF04857 | CAF1 | PF07742 | BTG | 0 | 6 | 0 |
| PF05739 | SNARE | PF05739 | SNARE | 0 | 12 | 0.17 |
| PF07645 | EGF_CA | PF07679 | I-set | 0 | 9 | 1.94 |
| PF07654 | C1-set | PF07654 | C1-set | 0 | 11 | 0.15 |
| PF07679 | I-set | PF07679 | I-set | 0 | 12 | 0.61 |
| PF07686 | V-set | PF07686 | V-set | 0 | 12 | 0.77 |
| PF07714 | Pkinase_Tyr | PF07714 | Pkinase_Tyr | 0 | 50 | 13.37 |
| PF00002 | 7tm_2 | PF04901 | RAMP | 0.01 | 5 | 0.03 |
| PF00005 | ABC_tran | PF00595 | PDZ | 0.01 | 7 | 1.26 |
| PF00008 | EGF | PF00048 | IL8 | 0.01 | 7 | 1.34 |
| PF00008 | EGF | PF07714 | Pkinase_Tyr | 0.01 | 23 | 13.85 |
| PF00008 | EGF | PF07732 | Cu-oxidase_3 | 0.01 | 5 | 0.26 |
| PF00017 | SH2 | PF00028 | Cadherin | 0.01 | 10 | 3.57 |
| PF00017 | SH2 | PF01146 | Caveolin | 0.01 | 7 | 1.49 |
| PF00018 | SH3_1 | PF00536 | SAM_1 | 0.01 | 9 | 2.77 |
| PF00022 | Actin | PF00307 | CH | 0.01 | 7 | 0.99 |
| PF00022 | Actin | PF00626 | Gelsolin | 0.01 | 5 | 0.11 |
| PF00023 | Ank | PF00041 | fn3 | 0.01 | 10 | 3.7 |
| PF00023 | Ank | PF00615 | RGS | 0.01 | 6 | 0.93 |
| PF00025 | Arf | PF00169 | PH | 0.01 | 6 | 0.67 |
| PF00036 | efhand | PF01080 | Presenilin | 0.01 | 6 | 0.55 |
| PF00041 | fn3 | PF00049 | Insulin | 0.01 | 5 | 0.29 |
| PF00041 | fn3 | PF00059 | Lectin_C | 0.01 | 8 | 1.97 |
| PF00041 | fn3 | PF00193 | Xlink | 0.01 | 7 | 0.88 |
| PF00041 | fn3 | PF07645 | EGF_CA | 0.01 | 11 | 3.99 |
| PF00047 | ig | PF00791 | ZU5 | 0.01 | 6 | 0.41 |
| PF00047 | ig | PF07654 | C1-set | 0.01 | 7 | 1.31 |
| PF00047 | ig | PF07974 | EGF_2 | 0.01 | 8 | 2.18 |
| PF00050 | Kazal_1 | PF00089 | Trypsin | 0.01 | 5 | 0.12 |


| PF00057 | Ldl_recept_a | PF01821 | ANATO | 0.01 | 5 | 0.08 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00060 | Lig_chan | PF00060 | Lig_chan | 0.01 | 5 | 0.07 |
| PF00069 | Pkinase | PF01248 | Ribosomal_L7Ae | 0.01 | 6 | 0.58 |
| PF00069 | Pkinase | PF01387 | Synuclein | 0.01 | 6 | 0.95 |
| PF00071 | Ras | PF00996 | GDI | 0.01 | 5 | 0.09 |
| PF00071 | Ras | PF02196 | RBD | 0.01 | 7 | 0.93 |
| PF00076 | RRM_1 | PF00641 | zf-RanBP | 0.01 | 6 | 0.79 |
| PF00084 | Sushi | PF07974 | EGF_2 | 0.01 | 6 | 0.78 |
| PF00089 | Trypsin | PF00594 | Gla | 0.01 | 5 | 0.34 |
| PF00090 | TSP_1 | PF00413 | Peptidase_M10 | 0.01 | 5 | 0.12 |
| PF00096 | zf-C2H2 | PF00389 | 2-Hacid_dh | 0.01 | 6 | 0.34 |
| PF00096 | zf-C2H2 | PF00628 | PHD | 0.01 | 6 | 0.91 |
| PF00097 | zf-C3HC4 | PF00097 | zf-C3HC4 | 0.01 | 8 | 2.25 |
| PF00097 | zf-C3HC4 | PF00385 | Chromo | 0.01 | 6 | 0.71 |
| PF00102 | Y_phosphatase | PF02174 | IRS | 0.01 | 6 | 0.41 |
| PF00134 | Cyclin_N | PF01857 | RB_B | 0.01 | 5 | 0.31 |
| PF00179 | UQ_con | PF00240 | ubiquitin | 0.01 | 5 | 0.32 |
| PF00194 | Carb_anhydrase | PF00955 | HCO3_cotransp | 0.01 | 5 | 0.02 |
| PF00240 | ubiquitin | PF02809 | UIM | 0.01 | 5 | 0.23 |
| PF00170 | bZIP_1 | PF03131 | bZIP_Maf | 0.01 | 5 | 0.03 |
| PF00249 | Myb_DNA-binding | PF00850 | Hist_deacetyl | 0.01 | 6 | 0.49 |
| PF00307 | CH | PF05556 | Calsarcin | 0.01 | 5 | 0.1 |
| PF00307 | CH | PF07679 | I-set | 0.01 | 8 | 1.53 |
| PF00335 | Tetraspannin | PF00335 | Tetraspannin | 0.01 | 5 | 0.01 |
| PF00340 | IL1 | PF01582 | TIR | 0.01 | 5 | 0.07 |
| PF00412 | LIM | PF00412 | LIM | 0.01 | 6 | 0.89 |
| PF00431 | CUB | PF01391 | Collagen | 0.01 | 5 | 0.16 |
| PF00503 | G-alpha | PF00621 | RhoGEF | 0.01 | 6 | 0.94 |
| PF00505 | HMG_box | PF00514 | Arm | 0.01 | 6 | 0.49 |
| PF00514 | Arm | PF00514 | Arm | 0.01 | 6 | 0.68 |
| PF00515 | TPR_1 | PF00515 | TPR_1 | 0.01 | 5 | 0.24 |
| PF00520 | Ion_trans | PF02060 | ISK_Channel | 0.01 | 5 | 0.03 |
| PF00531 | Death | PF00619 | CARD | 0.01 | 6 | 0.7 |
| PF00560 | LRR_1 | PF01463 | LRRCT | 0.01 | 5 | 0.24 |
| PF00560 | LRR_1 | PF01582 | TIR | 0.01 | 5 | 0.35 |
| PF00594 | Gla | PF07732 | Cu-oxidase_3 | 0.01 | 5 | 0.03 |
| PF00619 | CARD | PF02758 | PAAD_DAPIN | 0.01 | 5 | 0.01 |
| PF00621 | RhoGEF | PF01403 | Sema | 0.01 | 5 | 0.12 |
| PF00808 | CBFD_NFYB_HMF | PF00808 | CBFD_NFYB_HMF | 0.01 | 5 | 0.01 |
| PF00850 | Hist_deacetyl | PF02671 | PAH | 0.01 | 5 | 0.13 |
| PF00928 | Adap_comp_sub | PF02883 | Alpha_adaptinC2 | 0.01 | 5 | 0.07 |
| PF01217 | Clat_adaptor_s | PF02883 | Alpha_adaptinC2 | 0.01 | 5 | 0.01 |
| PF01391 | Collagen | PF01839 | FG-GAP | 0.01 | 6 | 0.91 |
| PF01412 | ArfGap | PB000002 | Pfam-B_2 | 0.01 | 5 | 0.33 |
| PF01821 | ANATO | PF07645 | EGF_CA | 0.01 | 5 | 0.2 |
| PF02931 | Neur_chan_LBD | PF02931 | Neur_chan_LBD | 0.01 | 5 | 0 |
| PF03165 | MH1 | PF03165 | MH1 | 0.01 | 5 | 0.12 |
| PF05008 | V-SNARE | PF05739 | SNARE | 0.01 | 5 | 0.06 |
| PF05739 | SNARE | PB000002 | Pfam-B_2 | 0.01 | 7 | 0.96 |
| PF07645 | EGF_CA | PF07974 | EGF_2 | 0.01 | 7 | 1.38 |
| PF07647 | SAM_2 | PB000002 | Pfam-B_2 | 0.01 | 7 | 0.99 |
| PF07686 | V-set | PF07974 | EGF_2 | 0.01 | 7 | 1.21 |
| PF07714 | Pkinase_Tyr | PF07686 | V-set | 0.01 | 14 | 6.28 |


| PF00005 | ABC_tran | PF00005 | ABC_tran | 0.02 | 4 | 0.02 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00008 | EGF | PF00262 | Calreticulin | 0.02 | 5 | 0.59 |
| PF00013 | KH_1 | PF00018 | SH3_1 | 0.02 | 9 | 3.47 |
| PF00017 | SH2 | PF00168 | C2 | 0.02 | 18 | 10.92 |
| PF00018 | SH3_1 | PF00611 | FCH | 0.02 | 7 | 1.94 |
| PF00023 | Ank | PF00412 | LIM | 0.02 | 7 | 1.88 |
| PF00025 | Arf | PF05351 | GMP_PDE_delta | 0.02 | 4 | 0.01 |
| PF00038 | Filament | PF00130 | C1_1 | 0.02 | 7 | 1.69 |
| PF00040 | fn2 | PF01839 | FG-GAP | 0.02 | 5 | 0.37 |
| PF00047 | ig | PF02189 | ITAM | 0.02 | 5 | 0.53 |
| PF00053 | Laminin_EGF | PF07645 | EGF_CA | 0.02 | 5 | 0.55 |
| PF00071 | Ras | PF00564 | PB1 | 0.02 | 6 | 1.16 |
| PF00071 | Ras | PF02185 | HR1 | 0.02 | 5 | 0.53 |
| PF00076 | RRM_1 | PF00271 | Helicase_C | 0.02 | 6 | 1.18 |
| PF00084 | Sushi | PF07645 | EGF_CA | 0.02 | 6 | 1.19 |
| PF00090 | TSP_1 | PF00089 | Trypsin | 0.02 | 5 | 0.53 |
| PF00102 | Y_phosphatase | PF07647 | SAM_2 | 0.02 | 5 | 0.56 |
| PF00130 | C1_1 | PF00130 | C1_1 | 0.02 | 6 | 1.18 |
| PF00244 | 14-3-3 | PF00850 | Hist_deacetyl | 0.02 | 5 | 0.55 |
| PF00249 | Myb_DNA-binding | PF00439 | Bromodomain | 0.02 | 5 | 0.6 |
| PF00307 | CH | PF07974 | EGF_2 | 0.02 | 6 | 1.14 |
| PF00571 | CBS | PF04739 | AMPKBI | 0.02 | 4 | 0.01 |
| PF00626 | Gelsolin | PF00626 | Gelsolin | 0.02 | 4 | 0.01 |
| PF00635 | Motile_Sperm | PF00957 | Synaptobrevin | 0.02 | 4 | 0 |
| PF00640 | PID | PF07679 | I-set | 0.02 | 6 | 1.15 |
| PF01652 | IF4E | PF05456 | eIF_4EBP | 0.02 | 4 | 0 |
| PF02136 | NTF2 | PB006713 | Pfam-B_6713 | 0.02 | 4 | 0 |
| PF00004 | AAA | PF00004 | AAA | 0.03 | 4 | 0.05 |
| PF00008 | EGF | PF00014 | Kunitz_BPTI | 0.03 | 5 | 0.83 |
| PF00008 | EGF | PF00093 | VWC | 0.03 | 5 | 0.7 |
| PF00010 | HLH | PF00412 | LIM | 0.03 | 7 | 2.15 |
| PF00012 | HSP70 | PF00226 | DnaJ | 0.03 | 4 | 0.06 |
| PF00012 | HSP70 | PF00240 | ubiquitin | 0.03 | 4 | 0.2 |
| PF00012 | HSP70 | PF02179 | BAG | 0.03 | 4 | 0.1 |
| PF00014 | Kunitz_BPTI | PF00040 | fn2 | 0.03 | 4 | 0.14 |
| PF00017 | SH2 | PF07679 | I-set | 0.03 | 14 | 7.98 |
| PF00019 | TGF_beta | PF00093 | VWC | 0.03 | 4 | 0.13 |
| PF00019 | TGF_beta | PF07974 | EGF_2 | 0.03 | 5 | 0.68 |
| PF00025 | Arf | PF02883 | Alpha_adaptinC2 | 0.03 | 4 | 0.09 |
| PF00038 | Filament | PF01477 | PLAT | 0.03 | 4 | 0.12 |
| PF00038 | Filament | PF02185 | HR1 | 0.03 | 4 | 0.2 |
| PF00041 | fn3 | PF00084 | Sushi | 0.03 | 7 | 2.26 |
| PF00041 | fn3 | PF00621 | RhoGEF | 0.03 | 8 | 2.83 |
| PF00041 | fn3 | PF01291 | LIF_OSM | 0.03 | 4 | 0.1 |
| PF00046 | Homeobox | PF00505 | HMG_box | 0.03 | 5 | 0.69 |
| PF00046 | Homeobox | PF02037 | SAP | 0.03 | 5 | 0.68 |
| PF00047 | ig | PF00048 | IL8 | 0.03 | 6 | 1.53 |
| PF00053 | Laminin_EGF | PF00053 | Laminin_EGF | 0.03 | 4 | 0.07 |
| PF00057 | Ldl_recept_a | PF00207 | A2M | 0.03 | 4 | 0.12 |
| PF00058 | Ldl_recept_b | PF00595 | PDZ | 0.03 | 7 | 2.21 |
| PF00059 | Lectin_C | PF00129 | MHC_I | 0.03 | 4 | 0.21 |
| PF00059 | Lectin_C | PF00431 | CUB | 0.03 | 4 | 0.09 |
| PF00059 | Lectin_C | PF07645 | EGF_CA | 0.03 | 6 | 1.28 |


| PF00060 | Lig_chan | PF00373 | Band_41 | 0.03 | 4 | 0.23 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00084 | Sushi | PF01391 | Collagen | 0.03 | 5 | 0.82 |
| PF00089 | Trypsin | PF02210 | Laminin_G_2 | 0.03 | 5 | 0.63 |
| PF00089 | Trypsin | PF07645 | EGF_CA | 0.03 | 7 | 2.04 |
| PF00089 | Trypsin | PF07732 | Cu-oxidase_3 | 0.03 | 4 | 0.14 |
| PF00096 | zf-C2H2 | PF00610 | DEP | 0.03 | 5 | 0.84 |
| PF00102 | Y_phosphatase | PF02985 | HEAT | 0.03 | 5 | 0.79 |
| PF00104 | Hormone_recep | PF00240 | ubiquitin | 0.03 | 6 | 1.42 |
| PF00104 | Hormone_recep | PF00643 | zf-B_box | 0.03 | 6 | 1.39 |
| PF00147 | Fibrinogen_C | PF01839 | FG-GAP | 0.03 | 4 | 0.2 |
| PF00168 | C2 | PF05739 | SNARE | 0.03 | 6 | 1.32 |
| PF00169 | PH | PF00620 | RhoGAP | 0.03 | 7 | 2.2 |
| PF00178 | Ets | PF07716 | bZIP_2 | 0.03 | 4 | 0.15 |
| PF00219 | IGFBP | PF07974 | EGF_2 | 0.03 | 4 | 0.15 |
| PF00243 | NGF | PF07679 | I-set | 0.03 | 4 | 0.07 |
| PF00320 | GATA | PB003900 | Pfam-B_3900 | 0.03 | 4 | 0.03 |
| PF00335 | Tetraspannin | PF07974 | EGF_2 | 0.03 | 4 | 0.2 |
| PF00439 | Bromodomain | PF05030 | SSXT | 0.03 | 4 | 0.06 |
| PF00452 | Bcl-2 | PF06553 | BNIP3 | 0.03 |  | 0.03 |
| PF00503 | G-alpha | PF00595 | PDZ | 0.03 | 9 | 3.62 |
| PF00520 | Ion_trans | PF00595 | PDZ | 0.03 | 8 | 3.04 |
| PF00531 | Death | PF00560 | LRR_1 | 0.03 | 6 | 1.26 |
| PF00536 | SAM_1 | PF00536 | SAM_1 | 0.03 | 4 | 0.03 |
| PF00536 | SAM_1 | PF07647 | SAM_2 | 0.03 | 4 | 0.15 |
| PF00595 | PDZ | PF03165 | MH1 | 0.03 | 7 | 2.06 |
| PF00595 | PDZ | PF07653 | SH3_2 | 0.03 | 8 | 2.83 |
| PF00605 | IRF | PF01833 | TIG | 0.03 | 4 | 0.07 |
| PF00619 | CARD | PF00653 | BIR | 0.03 | 4 | 0.16 |
| PF00638 | Ran_BP1 | PF03810 | IBN_N | 0.03 | 4 | 0.09 |
| PF00995 | Sec1 | PF05739 | SNARE | 0.03 | 4 | 0.05 |
| PF01391 | Collagen | PF07546 | EMI | 0.03 |  | 0.08 |
| PF01582 | TIR | PF01582 | TIR | 0.03 | 4 | 0.07 |
| PF01602 | Adaptin_N | PF01602 | Adaptin_N | 0.03 |  | 0.05 |
| PF01833 | TIG | PF01833 | TIG | 0.03 | 4 | 0.23 |
| PF01857 | RB_B | PB000002 | Pfam-B_2 | 0.03 | 5 | 0.61 |
| PF02319 | E2F_TDP | PF01857 | RB_B | 0.03 | 4 | 0.09 |
| PF02437 | Ski_Sno | PF03165 | MH1 | 0.03 | 4 | 0.1 |
| PF02985 | HEAT | PB000002 | Pfam-B_2 | 0.03 | 6 | 1.35 |
| PF03810 | IBN_N | PB020723 | Pfam-B_20723 | 0.03 | 4 | 0.05 |
| PF03921 | ICAM_N | PF07974 | EGF_2 | 0.03 | 4 | 0.13 |
| PF05507 | MAGP | PF07645 | EGF_CA | 0.03 | 4 | 0.08 |
| PF07716 | bZIP_2 | PF07716 | bZIP_2 | 0.03 | 4 | 0.11 |
| PF00010 | HLH | PF00319 | SRF-TF | 0.04 | 4 | 0.32 |
| PF00013 | KH_1 | PF00017 | SH2 | 0.04 | 8 | 3.24 |
| PF00017 | SH2 | PB001239 | Pfam-B_1239 | 0.04 |  | 0.31 |
| PF00018 | SH3_1 | PF07653 | SH3_2 | 0.04 | 10 | 5.09 |
| PF00019 | TGF_beta | PF01462 | LRRNT | 0.04 | 4 | 0.35 |
| PF00040 | fn2 | PF00048 | IL8 | 0.04 | 4 | 0.3 |
| PF00040 | fn2 | PF07645 | EGF_CA | 0.04 | 5 | 0.89 |
| PF00046 | Homeobox | PF00439 | Bromodomain | 0.04 | 6 | 1.7 |
| PF00046 | Homeobox | PF03920 | TLE_N | 0.04 | 4 | 0.37 |
| PF00047 | ig | PF00340 | IL1 | 0.04 | 4 | 0.29 |
| PF00047 | ig | PF00531 | Death | 0.04 | 7 | 2.43 |


| PF00047 | ig | PF01839 | FG-GAP | 0.04 | 6 | 1.73 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| PF00059 | Lectin_C | PF00089 | Trypsin | 0.04 | 5 | 0.94 |
| PF00059 | Lectin_C | PF00147 | Fibrinogen_C | 0.04 | 4 | 0.3 |
| PF00069 | Pkinase | PF01335 | DED | 0.04 | 8 | 3.3 |
| PF00076 | RRM_1 | PF02037 | SAP | 0.04 | 5 | 0.89 |
| PF00089 | Trypsin | PF00207 | A2M | 0.04 | 4 | 0.35 |
| PF00104 | Hormone_recep | PF00096 | zf-C2H2 | 0.04 | 9 | 4.23 |
| PF00147 | Fibrinogen_C | PF07645 | EGF_CA | 0.04 | 4 | 0.38 |
| PF00249 | Myb_DNA-binding | PF07716 | bZIP_2 | 0.04 | 4 | 0.37 |
| PF00320 | GATA | PF00412 | LIM | 0.04 | 4 | 0.24 |
| PF00386 | C1q | PF01391 | Collagen | 0.04 | 4 | 0.26 |
| PF00400 | WD40 | PF00400 | WD40 | 0.04 | 5 | 0.95 |
| PF00400 | WD40 | PF00515 | TPR_1 | 0.04 | 5 | 0.98 |
| PF00400 | WD40 | PF00856 | SET | 0.04 | 4 | 0.28 |
| PF00439 | Bromodomain | PF00641 | zf-RanBP | 0.04 | 4 | 0.37 |
| PF00514 | Arm | PF03810 | IBN_N | 0.04 | 4 | 0.33 |
| PF00564 | PB1 | PF00595 | PDZ | 0.04 | 6 | 1.63 |
| PF00595 | PDZ | PF00858 | ASC | 0.04 | 4 | 0.28 |
| PF00595 | PDZ | PF02210 | Laminin_G_2 | 0.04 | 6 | 1.79 |
| PF00615 | RGS | PF01462 | LRRNT | 0.04 | 4 | 0.35 |
| PF00788 | RA | PF07647 | SAM_2 | 0.04 | 4 | 0.27 |
| PF00928 | Adap_comp_sub | PF07686 | V-set | 0.04 | 4 | 0.36 |
| PF00012 | HSP70 | PF00397 | WW | 0.05 | 4 | 0.41 |
| PF00014 | Kunitz_BPTI | PF00089 | Trypsin | 0.05 | 4 | 0.39 |
| PF00041 | fn3 | PF01839 | FG-GAP | 0.05 | 6 | 1.86 |
| PF00071 | Ras | PF00514 | Arm | 0.05 | 7 | 2.62 |
| PF00097 | zf-C3HC4 | PF00531 | Death | 0.05 | 7 | 2.59 |
| PF00097 | zf-C3HC4 | PF07714 | Pkinase_Tyr | 0.05 | 17 | 11.62 |
| PF00100 | Zona_pellucida | PF00069 | Pkinase | 0.05 | 5 | 1.12 |
| PF00307 | CH | PF00595 | PDZ | 0.05 | 9 | 4.31 |
| PF00335 | Tetraspannin | PF07686 | V-set | 0.05 | 4 | 0.39 |
| PF00454 | PI3_PI4_kinase | PF07714 | Pkinase_Tyr | 0.05 | 6 | 1.81 |
| PF00531 | Death | PF01454 | MAGE | 0.05 | 4 | 0.43 |
| PF00627 | UBA | PF00643 | zf-B_box | 0.05 | 4 | 0.4 |

