|  |  |  |  | Num | Avg num in |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| ID_1 | Name_1 | ID_2 | Name_2 | FDR | in yeast | random networks | in human?


| PF00324 | AA_permease | PF03198 | PF03198 | 0 | 7 | 0.77 | N |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF00400 | WD40 | PF00400 | WD40 | 0 | 8 | 2.05 | Y |
| PF00674 | PF00674 | PF00674 | PF00674 | 0 | 7 | 1.26 | N |
| PF00735 | GTP_CDC | PF00735 | GTP_CDC | 0 | 9 | 0.15 | Y |
| PF00735 | GTP_CDC | PF00786 | PBD | 0 | 8 | 0.47 | Y |
| PF00786 | PBD | PB063688 | PB063688 | 0 | 5 | 0.25 | N |
| PF00957 | Synaptobrevin | PF05008 | V-SNARE | 0 | 7 | 0.26 | N |
| PF01174 | PF01174 | PF01680 | PF01680 | 0 | 7 | 0.02 | N |
| PF01398 | Mov34 | PF01399 | PCI | 0 | 5 | 0.09 | Y |
| PF01423 | LSM | PF01423 | LSM | 0 | 41 | 2.75 |  |
| PF02463 | SMC_N | PF02463 | SMC_N | 0 | 6 | 0.07 | N |
| PF02985 | HEAT | PF04096 | Nucleoporin2 | 0 | 6 | 0.23 |  |
| PF05739 | SNARE | PF00957 | Synaptobrevin | 0 | 18 | 0.78 |  |
| PF05739 | SNARE | PF00995 | Sec1 | 0 | 7 | 0.22 |  |
| PF05739 | SNARE | PF05008 | V-SNARE | 0 | 11 | 0.4 |  |
| PF05739 | SNARE | PF05739 | SNARE | 0 | 6 | 0.56 |  |
| PF06371 | Drf_GBD | PF00018 | SH3_1 | 0 | 8 | 0.91 | N |
| PF07647 | SAM_2 | PF00069 | Pkinase | 0 | 9 | 1.84 |  |
| PF07690 | MFS_1 | PF00137 | ATP-synt_C | 0 | 9 | 0.52 | N |
| PF07690 | MFS_1 | PF00674 | PF00674 | 0 | 20 | 2.41 | N |
| PF07690 | MFS_1 | PF01105 | EMP24_GP25L | 0 | 8 | 1.32 | N |
| PB046617 | PB046617 | PF00957 | Synaptobrevin | 0.01 | 4 | 0.14 | N |
| PB073262 | PB073262 | PF00169 | PH | 0.01 | 4 | 0.15 | N |
| PF00012 | HSP70 | PF00515 | TPR_1 | 0.01 | 5 | 0.62 |  |
| PF00018 | SH3_1 | PF00611 | FCH | 0.01 | 7 | 1.61 |  |
| PF00025 | Arf | PF00790 | VHS | 0.01 | 4 | 0.07 |  |
| PF00069 | Pkinase | PF00400 | WD40 | 0.01 | 17 | 10.91 |  |
| PF00071 | Ras | PF00612 | IQ | 0.01 | 5 | 0.46 |  |
| PF00071 | Ras | PF00786 | PBD | 0.01 | 6 | 1.25 |  |
| PF00076 | RRM_1 | PF02854 | MIF4G | 0.01 | 5 | 0.34 |  |
| PF00091 | Tubulin | PF01302 | CAP_GLY | 0.01 | 4 | 0.02 |  |
| PF00096 | zf-C2H2 | PF00515 | TPR_1 | 0.01 | 5 | 0.52 | N |
| PF00102 | Y_phosphatase | PF00069 | Pkinase | 0.01 | 5 | 0.32 |  |
| PF00136 | DNA_pol_B | PF04042 | DNA_pol_E_B | 0.01 | 4 | 0 |  |
| PF00153 | Mito_carr | PF00674 | PF00674 | 0 | 6 | 0.76 | N |
| PF00149 | Metallophos | PF03095 | PTPA | 0.01 | 4 | 0.06 |  |
| PF00156 | Pribosyltran | PF00156 | Pribosyltran | 0.01 | 4 | 0.02 |  |
| PF00169 | PH | PF00735 | GTP_CDC | 0.01 | 5 | 0.48 | Y |
| PF00179 | UQ_con | PF02037 | SAP | 0.01 | 4 | 0.06 |  |
| PF00271 | Helicase_C | PF00271 | Helicase_C | 0.01 | 5 | 0.62 | N |
| PF00271 | Helicase_C | PF01423 | LSM | 0.01 | 8 | 2.66 |  |
| PF00400 | WD40 | PF00515 | TPR_1 | 0.01 | 7 | 1.83 | Y |
| PF00439 | Bromodomain | PF04433 | SWIRM | 0.01 | 4 | 0.02 | Y |
| PF00454 | PI3_PI4_kinase | PF00498 | FHA | 0.01 | 4 | 0.06 | N |
| PF00454 | PI3_PI4_kinase | PF01336 | tRNA_anti | 0.01 | 4 | 0.07 | Y |
| PF00493 | MCM | PF00493 | MCM | 0.01 | 4 | 0.03 | Y |
| PF00564 | PB1 | PF00786 | PBD | 0.01 | 4 | 0.11 | N |
| PF00928 | Adap_comp_sub | PF01602 | Adaptin_N | 0.01 | 4 | 0.02 |  |


| PF00957 | Synaptobrevin | PF00995 | Sec1 | 0.01 | 4 | 0.11 | Y |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| PF01105 | EMP24_GP25L | PF01066 | CDP-OH_P_transf | 0.01 | 5 | 0.4 | N |
| PF03162 | PF03162 | PF03162 | PF03162 | 0.01 | 4 | 0 | N |
| PF06398 | PF06398 | PF06398 | PF06398 | 0.01 | 4 | 0 | N |
| PB000002 | Pfam-B_2 | PF00036 | efhand | 0.02 | 5 | 0.96 | Y |
| PB000031 | Pfam-B_31 | PF00271 | Helicase_C | 0.02 | 4 | 0.25 | Y |
| PB000521 | Pfam-B_521 | PF04811 | Sec23_trunk | 0.02 | 4 | 0.28 | N |
| PB028889 | PB028889 | PF00069 | Pkinase | 0.02 | 4 | 0.32 | N |
| PF00005 | ABC_tran | PF01423 | LSM | 0.02 | 6 | 1.65 | N |
| PF00018 | SH3_1 | PF00581 | Rhodanese | 0.02 | 5 | 0.79 | Y |
| PF00022 | Actin | PF00018 | SH3_1 | 0.02 | 6 | 1.6 | Y |
| PF00023 | Ank | PF00069 | Pkinase | 0.02 | 7 | 2.26 | Y |
| PF00069 | Pkinase | PF00481 | PP2C | 0.02 | 5 | 0.87 | Y |
| PF00069 | Pkinase | PF04153 | NOT2_3_5 | 0.02 | 5 | 0.91 | N |
| PF00071 | Ras | PF00564 | PB1 | 0.02 | 4 | 0.37 | Y |
| PF00076 | RRM_1 | PF02985 | HEAT | 0.02 | 6 | 1.63 | Y |
| PF00083 | Sugar_tr | PF00515 | TPR_1 | 0.02 | 5 | 1 | N |
| PF00091 | Tubulin | PF02985 | HEAT | 0.02 | 4 | 0.17 | Y |
| PF00611 | FCH | PF02205 | WH2 | 0.02 | 4 | 0.2 | Y |
| PF00787 | PX | PF04893 | Yip1 | 0.02 | 4 | 0.25 | N |
| PF07651 | ANTH | PF00036 | efhand | 0.02 | 4 | 0.18 | Y |
| PF08171 | PF08171 | PF00400 | WD40 | 0.02 | 4 | 0.2 | N |
| PF00069 | Pkinase | PF00224 | PK | 0.03 | 4 | 0.42 | N |
| PF00149 | Metallophos | PF00439 | Bromodomain | 0.03 | 4 | 0.47 | N |
| PF00170 | bZIP_1 | PF02985 | HEAT | 0.03 | 4 | 0.43 | N |
| PF01105 | EMP24_GP25L | PF01598 | Sterol_desat | 0.03 | 4 | 0.39 | N |
| PF07647 | SAM_2 | PF00018 | SH3_1 | 0.03 | 5 | 1.06 | Y |
| PF07690 | MFS_1 | PF01598 | Sterol_desat | 0.03 | 4 | 0.45 | N |
| PF00009 | GTP_EFTU | PF01423 | LSM | 0.04 | 5 | 1.28 | N |
| PF00069 | Pkinase | PF00564 | PB1 | 0.04 | 5 | 1.28 | Y |
| PF06371 | Drf_GBD | PF00071 | Ras | 0.04 | 4 | 0.58 | Y |
| PB008540 | PB008540 | PF02133 | PF02133 | 0.05 | 3 | 0.04 | N |
| PF00012 | HSP70 | PF00012 | HSP70 | 0.05 | 3 | 0.06 | Y |
| PF00036 | efhand | PF00149 | Metallophos | 0.05 | 4 | 0.69 | Y |
| PF00085 | Thioredoxin | PF00578 | AhpC-TSA | 0.05 | 3 | 0.02 | N |
| PF00132 | Hexapep | PF01008 | IF-2B | 0.05 | 3 | 0.01 | Y |
| PF00149 | Metallophos | PF03370 | CBM_21 | 0.05 | 3 | 0.07 | Y |
| PF00240 | ubiquitin | PF00899 | ThiF | 0.05 | 3 | 0.04 | Y |
| PF00428 | Ribosomal_60s | PF00428 | Ribosomal_60s | 0.05 | 3 | 0 | Y |
| PF00432 | Prenyltrans | PF01239 | PPTA | 0.05 | 3 | 0 | Y |
| PF00488 | MutS_V | PF00488 | MutS_V | 0.05 | 3 | 0.03 | Y |
| PF00488 | MutS_V | PF01119 | DNA_mis_repair | 0.05 | 3 | 0.02 | Y |
| PF00562 | RNA_pol_Rpb2_6 | PF01096 | TFIIS_C | 0.05 | 3 | 0.01 | N |
| PF00562 | RNA_pol_Rpb2_6 | PF01193 | RNA_pol_L | 0.05 | 3 | 0.03 | N |
| PF00638 | Ran_BP1 | PF03810 | IBN_N | 0.05 | 3 | 0.06 | Y |
| PF01066 | CDP-OH_P_transf | PF00674 | PF00674 | 0.05 | 4 | 0.65 | N |
| PF01138 | RNase_PH | PF01138 | RNase_PH | 0.05 | 3 | 0 | Y |
| PF01193 | RNA_pol_L | PF01193 | RNA_pol_L | 0.05 | 3 | 0 | N |


| PF01399 | PCI | PF01399 | PCI | 0.05 | 3 | 0.03 | Y |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| PF01853 | MOZ_SAS | PF03366 | YEATS | 0.05 | 3 | 0 | N |
| PF01920 | KE2 | PF02996 | Prefoldin | 0.05 | 3 | 0 | N |
| PF02466 | Tim17 | PF02953 | zf-Tim10_DDP | 0.05 | 3 | 0 | N |
| PF02815 | MIR | PF02815 | MIR | 0.05 | 3 | 0.01 | Y |
| PF03357 | ESCRT-III | PF03357 | ESCRT-III | 0.05 | 3 | 0 | N |
| PF03911 | Sec61_beta | PF04756 | OST3_OST6 | 0.05 | 3 | 0.03 | N |
| PF07651 | ANTH | PF08226 | PF08226 | 0.05 | 3 | 0.04 | N |
| PF08240 | ADH_N | PF08240 | ADH_N | 0.05 | 3 | 0.02 | N |

