

| ID_1 | Name_1 | ID_2 | Name_2 | ID_3 | Name_3 | FDR | Count in yeast | Avg count in random | Instance in human? |
|----------|-------------|---------|---------------|---------|---------------|------|-------------------|------------------------|-----------------------|
| PF00018 | SH3_1 | PF00018 | SH3_1 | PF02205 | WH2 | 0 | 22 | 5.54 | Y |
| PF00069 | Pkinase | PF00735 | GTP_CDC | PF00735 | GTP_CDC | 0 | 9 | 0.22 | N |
| PF00125 | Histone | PF00125 | Histone | PF00439 | Bromodomain | 0 | 10 | 0.75 | N |
| PF01423 | LSM | PF01423 | LSM | PF01423 | LSM | 0 | 64 | 26.65 | Y |
| PF05739 | SNARE | PF00957 | Synaptobrevin | PF05008 | V-SNARE | 0 | 17 | 3.73 | N |
| PB000521 | Pfam-B_521 | PF00324 | AA_permease | PF00674 | PF00674 | 0 | 4 | 0.04 | N |
| PB042766 | PB042766 | PF00069 | Pkinase | PF00069 | Pkinase | 0 | 3 | 0.13 | N |
| PF00004 | AAA | PF00004 | AAA | PF00004 | AAA | 0 | 4 | 0.03 | Y |
| PF00018 | SH3_1 | PF00018 | SH3_1 | PF00018 | SH3_1 | 0 | 7 | 1.24 | Y |
| PF00018 | SH3_1 | PF00018 | SH3_1 | PF00069 | Pkinase | 0 | 4 | 0.16 | Y |
| PF00018 | SH3_1 | PF00018 | SH3_1 | PF00071 | Ras | 0 | 3 | 0.05 | Y |
| PF00022 | Actin | PF00018 | SH3_1 | PF00018 | SH3_1 | 0 | 3 | 0.07 | N |
| PF00069 | Pkinase | PF00069 | Pkinase | PF00069 | Pkinase | 0 | 3 | 0.1 | Y |
| PF00069 | Pkinase | PF00069 | Pkinase | PF00134 | Cyclin_N | 0 | 5 | 0.03 | Y |
| PF00069 | Pkinase | PF00169 | PH | PF00564 | PB1 | 0 | 3 | 0 | Y |
| PF00169 | PH | PF00564 | PB1 | PF00786 | PBD | 0 | 3 | 0.31 | N |
| PF00324 | AA_permease | PF01105 | EMP24_GP25L | PF01598 | Sterol_desat | 0 | 3 | 0.26 | N |
| PF00735 | GTP_CDC | PF00735 | GTP_CDC | PF00786 | PBD | 0 | 7 | 1.63 | N |
| PF05739 | SNARE | PF00957 | Synaptobrevin | PF00995 | Sec1 | 0 | 7 | 0.64 | Y |
| PF05739 | SNARE | PF05739 | SNARE | PF00957 | Synaptobrevin | 0 | 7 | 2.54 | Y |
| PF05739 | SNARE | PF05739 | SNARE | PF05008 | V-SNARE | 0 | 7 | 2.07 | Y |
| PF07647 | SAM_2 | PF00018 | SH3_1 | PF00069 | Pkinase | 0 | 3 | 0.02 | Y |
| PF07690 | MFS_1 | PF00137 | ATP-synt_C | PF01105 | EMP24_GP25L | 0 | 8 | 0.08 | N |
| PF00018 | SH3_1 | PF00071 | Ras | PF00071 | Ras | 0.01 | 2 | 0 | Y |
| PF00125 | Histone | PF00125 | Histone | PF03810 | IBN_N | 0.01 | 2 | 0.21 | N |
| PF07690 | MFS_1 | PF00324 | AA_permease | PF00674 | PF00674 | 0.01 | 2 | 0.06 | N |