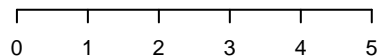


gc003b

fraction genes in fg and bg

| | | |
|-----------------|-----------|------|
| 7tm_1 | p=1.3E-04 | n=10 |
| Exo_endo_phos | p=7.2E-03 | n=2 |
| ANF_receptor | p=9.0E-03 | n=3 |
| Nfu_N | p=9.0E-03 | n=1 |
| NifU | p=9.0E-03 | n=1 |
| RT_RNaseH_2 | p=9.0E-03 | n=1 |
| PAX | p=1.5E-02 | n=1 |
| ADP_ribosyl_GH | p=2.4E-02 | n=1 |
| G-gamma | p=2.4E-02 | n=1 |
| CarboxypepD_reg | p=3.7E-02 | n=1 |
| 7tm_3 | p=3.9E-02 | n=2 |
| Peptidase_M14 | p=5.3E-02 | n=1 |
| Arrestin_C | p=9.0E-02 | n=1 |
| Arrestin_N | p=9.0E-02 | n=1 |
| Galactosyl_T | p=9.4E-02 | n=1 |
| Kringle | p=9.4E-02 | n=1 |
| Abhydrolase_1 | p=9.5E-02 | n=1 |
| Homeodomain | p=9.8E-02 | n=1 |
| fn3 | p=1.2E-01 | n=1 |
| I-set | p=1.2E-01 | n=1 |
| Pkinase_Tyr | p=1.2E-01 | n=1 |
| Ig_3 | p=1.2E-01 | n=1 |
| SRCR | p=1.4E-01 | n=1 |



$-\log_{10}(p)$
n=23/40 input genes with annotations

| | |
|---------|---------|
| fg=0.28 | bg=0.05 |
| fg=0.06 | bg=0.00 |
| fg=0.08 | bg=0.01 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.06 | bg=0.01 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |
| fg=0.03 | bg=0.00 |



fraction