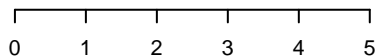


## Fox\_Nvec\_vc1.1\_XM\_032385496.2

## fraction genes in fg and bg

7tm_3	p=1.4E-01	n=2
DSL	p=1.4E-01	n=1
LRR_4	p=1.4E-01	n=1
MNNL	p=1.4E-01	n=1
PDEase_I_N	p=1.4E-01	n=1
Plug_translocon	p=1.4E-01	n=1
SecY	p=1.4E-01	n=1
GPS	p=1.6E-01	n=2
LRR_9	p=1.6E-01	n=1
Neurensin	p=1.6E-01	n=1
PHR	p=1.6E-01	n=1
Sulfotransfer_1	p=1.6E-01	n=1
AA_permease_C	p=1.7E-01	n=1
ANF_receptor	p=1.7E-01	n=2
DUF885	p=1.7E-01	n=1
GAIN	p=1.7E-01	n=1
Lactamase_B	p=1.7E-01	n=1
P_proprotein	p=1.7E-01	n=1
Peptidase_S8	p=1.7E-01	n=1
REJ	p=1.7E-01	n=1
RILa	p=1.7E-01	n=1
S8_pro-domain	p=1.7E-01	n=1
AA_permease_2	p=1.9E-01	n=1
Astacin	p=1.9E-01	n=1
BTB_2	p=1.9E-01	n=2
DEP	p=1.9E-01	n=1
MFS_1_like	p=1.9E-01	n=1
PDEase_I	p=1.9E-01	n=1
PKD_channel	p=1.9E-01	n=1
WSC	p=1.9E-01	n=1

-log<sub>10</sub>(p)

n=48/41 input genes with annotations

fg=0.04	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.04	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.04	bg=0.01
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.04	bg=0.01
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00



fraction