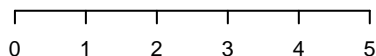


## Fox\_Nvec\_vc1.1\_XM\_001628634.3

## fraction genes in fg and bg

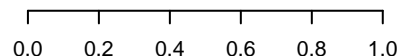
COP1coated_ERV	p=4.4E-02	n=1
DSL	p=4.4E-02	n=1
ERGIC_N	p=4.4E-02	n=1
Hid1	p=4.4E-02	n=1
MNNL	p=4.4E-02	n=1
NPDC1	p=4.4E-02	n=1
PfkB	p=4.4E-02	n=1
SapA	p=4.4E-02	n=1
SapB_1	p=4.4E-02	n=1
SapB_2	p=4.4E-02	n=1
SBF2	p=4.4E-02	n=1
Ski_Sno	p=4.4E-02	n=1
TRP_2	p=4.4E-02	n=1
7tm_1	p=6.1E-02	n=7
OAR	p=6.3E-02	n=1
WES_acyltransf	p=6.3E-02	n=1
WS_DGAT_C	p=6.3E-02	n=1
zf-met	p=6.3E-02	n=1
AA_permease_C	p=1.1E-01	n=1
VWA	p=1.1E-01	n=1
Ion_trans_2	p=1.2E-01	n=2
Cadherin	p=1.3E-01	n=1
AA_permease_2	p=1.3E-01	n=1
Lig_chan	p=1.3E-01	n=1
Lig_chan-Glu_bd	p=1.3E-01	n=1
zf-C3HC4	p=1.3E-01	n=1
F-box-like	p=1.4E-01	n=1
Kinesin	p=1.7E-01	n=1
LRR_6	p=1.7E-01	n=1
Ion_trans	p=1.9E-01	n=2



-log10(p)

n=35/39 input genes with annotations

fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
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fg=0.02	bg=0.00
fg=0.02	bg=0.00
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fg=0.02	bg=0.00
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fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.02	bg=0.00
fg=0.16	bg=0.07
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=0.05	bg=0.01
fg=0.02	bg=0.00
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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=0.05	bg=0.02



fraction