

## Fox\_Nvec\_vc1.1\_XM\_001639692.3

## fraction genes in fg and bg

WHIM1	p=1.0E-02	n=3
Neurensin	p=5.5E-02	n=2
DDT	p=8.1E-02	n=2
WSD	p=8.1E-02	n=2
Antistatin	p=1.1E-01	n=1
Arm_vescicular	p=1.1E-01	n=1
B12D	p=1.1E-01	n=1
Band_7	p=1.1E-01	n=2
Bromodomain	p=1.1E-01	n=3
CaM_bdg_C0	p=1.1E-01	n=1
CAP	p=1.1E-01	n=2
Cation_ATPase	p=1.1E-01	n=2
CEBP_ZZ	p=1.1E-01	n=1
DBB	p=1.1E-01	n=1
Drc1-Sld2	p=1.1E-01	n=1
DUF11	p=1.1E-01	n=1
DUF3583	p=1.1E-01	n=1
Dynactin	p=1.1E-01	n=1
ELM2	p=1.1E-01	n=2
ETS_PEA3_N	p=1.1E-01	n=1
FAM117	p=1.1E-01	n=1
Fes1	p=1.1E-01	n=1
Fzo_mitofusin	p=1.1E-01	n=1
HMMR_C	p=1.1E-01	n=1
Il2rg	p=1.1E-01	n=1
Ion_trans	p=1.1E-01	n=5
KCT2	p=1.1E-01	n=1
Morc6_S5	p=1.1E-01	n=1
NosD	p=1.1E-01	n=1
PHD	p=1.1E-01	n=3

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

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

n=185/155 input genes with annotations



fraction