

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

# fraction genes in fg and bg

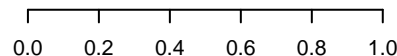
WHIM1	p=4.3E-03	n=4
DCX	p=1.7E-02	n=3
DDT	p=1.7E-02	n=3
Ribosomal_S15	p=1.7E-02	n=3
WSD	p=1.7E-02	n=3
Neurensin	p=9.5E-02	n=2
AICARFT_IMPCHas	p=1.6E-01	n=1
Antistasin	p=1.6E-01	n=1
Archease	p=1.6E-01	n=1
Arm_vescicular	p=1.6E-01	n=1
B12D	p=1.6E-01	n=1
Band_7	p=1.6E-01	n=3
Bromodomain	p=1.6E-01	n=4
BTB_2	p=1.6E-01	n=6
CaM_bdg_C0	p=1.6E-01	n=1
CBFB_NFYA	p=1.6E-01	n=1
CDK5_activator	p=1.6E-01	n=1
CEBP_ZZ	p=1.6E-01	n=1
CENP-T_C	p=1.6E-01	n=1
CG-1	p=1.6E-01	n=1
DBB	p=1.6E-01	n=1
DNA_pol3_delta2	p=1.6E-01	n=1
DOT1	p=1.6E-01	n=1
DPCD	p=1.6E-01	n=1
Drc1-Sld2	p=1.6E-01	n=1
DUF11	p=1.6E-01	n=1
DUF1977	p=1.6E-01	n=1
DUF3583	p=1.6E-01	n=1
Dynactin	p=1.6E-01	n=1
EF-hand_7	p=1.6E-01	n=7

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



-log10(p)

n=367/354 input genes with annotations



fraction