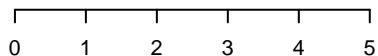


Fox\_Nvec\_vc1.1\_XM\_001625998.3

fraction genes in fg and bg

A_deaminase	p=1.1E-01	n=1
AAA_28	p=1.1E-01	n=1
ACC_central	p=1.1E-01	n=1
ADP_PFK_GK	p=1.1E-01	n=1
Anoct_dimer	p=1.1E-01	n=2
APP_amyloid	p=1.1E-01	n=1
APP_Cu_bd	p=1.1E-01	n=1
APP_E2	p=1.1E-01	n=1
APP_N	p=1.1E-01	n=1
Auts2	p=1.1E-01	n=1
BET	p=1.1E-01	n=1
BRAP2	p=1.1E-01	n=1
BRD4_CDT	p=1.1E-01	n=1
CCDC85	p=1.1E-01	n=1
CD225	p=1.1E-01	n=2
CDC45	p=1.1E-01	n=1
CEBP_ZZ	p=1.1E-01	n=1
Clathrin_lg_ch	p=1.1E-01	n=1
Coiled-coil_56	p=1.1E-01	n=1
CortBP2	p=1.1E-01	n=1
COX6C	p=1.1E-01	n=2
DDE_Tnp_1_7	p=1.1E-01	n=2
DNA_primase_lrg	p=1.1E-01	n=1
DSPc	p=1.1E-01	n=3
DUF155	p=1.1E-01	n=1
DUF1751	p=1.1E-01	n=1
DUF4419	p=1.1E-01	n=1
DUF647	p=1.1E-01	n=1
ETF_alpha	p=1.1E-01	n=1
Exo5	p=1.1E-01	n=1

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



$-\log_{10}(p)$   
n=326/282 input genes with annotations



fraction