

Ncol\_Nvec\_vc1.1\_XM\_032383907.2

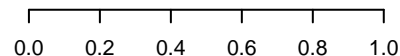
fraction genes in fg and bg

2OG-Fell_Oxy	p=7.8E-02	n=2
DIOX_N	p=7.8E-02	n=2
Glyco_transf_92	p=7.8E-02	n=2
UMPH-1	p=7.8E-02	n=2
5-nucleotidase	p=1.2E-01	n=1
Acetyltransf_13	p=1.2E-01	n=1
Aida_C2	p=1.2E-01	n=1
Aida_N	p=1.2E-01	n=1
ATG16	p=1.2E-01	n=1
Autophagy_act_C	p=1.2E-01	n=1
Biotin_lipoyl_2	p=1.2E-01	n=1
BTB_2	p=1.2E-01	n=4
CAP-ZIP_m	p=1.2E-01	n=1
CCSAP	p=1.2E-01	n=1
CDC50	p=1.2E-01	n=1
CENP-T_C	p=1.2E-01	n=1
Cl-B14_5a	p=1.2E-01	n=1
CLU	p=1.2E-01	n=1
CLU_N	p=1.2E-01	n=1
CTF_NFI	p=1.2E-01	n=1
DHFR_1	p=1.2E-01	n=1
DNA_primase_lrg	p=1.2E-01	n=1
DUF2205	p=1.2E-01	n=1
DUF4485	p=1.2E-01	n=1
DUF4611	p=1.2E-01	n=1
EF-hand_5	p=1.2E-01	n=3
eIF3_p135	p=1.2E-01	n=1
ERO1	p=1.2E-01	n=1
Es2	p=1.2E-01	n=1
FAM72	p=1.2E-01	n=1

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

 $-\log_{10}(p)$ 

n=322/304 input genes with annotations



fraction