

Ncol_Nvec_vc1.1_XM_001637272.3

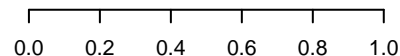
fraction genes in fg and bg

5-nucleotidase	p=7.6E-02	n=1
6PF2K	p=7.6E-02	n=1
AA_kinase	p=7.6E-02	n=1
AdoHcyase	p=7.6E-02	n=1
AdoHcyase_NAD	p=7.6E-02	n=1
ARL2_Bind_BART	p=7.6E-02	n=1
BLOC1_2	p=7.6E-02	n=1
bZIP_1	p=7.6E-02	n=3
CBF_beta	p=7.6E-02	n=1
CDO_I	p=7.6E-02	n=1
CLIP1_ZNF	p=7.6E-02	n=1
DOR	p=7.6E-02	n=1
DUF2045	p=7.6E-02	n=1
DUF2052	p=7.6E-02	n=1
Dynein_attach_N	p=7.6E-02	n=1
Filamin	p=7.6E-02	n=3
GGACT	p=7.6E-02	n=1
Gln-synt_C	p=7.6E-02	n=1
Gln-synt_N	p=7.6E-02	n=1
Granulin	p=7.6E-02	n=1
His_Phos_1	p=7.6E-02	n=1
IMD	p=7.6E-02	n=1
JAMP	p=7.6E-02	n=1
Leu_zip	p=7.6E-02	n=1
MitMem_reg	p=7.6E-02	n=1
Neur_chan_LBD	p=7.6E-02	n=4
Neur_chan_memb	p=7.6E-02	n=4
Nt_Gln_amidase	p=7.6E-02	n=1
Oxysterol_BP	p=7.6E-02	n=2
PNP_phzG_C	p=7.6E-02	n=1

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

 $-\log_{10}(p)$

n=171/153 input genes with annotations



fraction