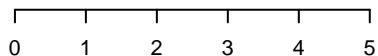


Ncol_Nvec_vc1.1_XM_001639701.3

fraction genes in fg and bg

4HBT_2	p=9.2E-02	n=1
6PF2K	p=9.2E-02	n=1
A_deaminase	p=9.2E-02	n=1
Agenet	p=9.2E-02	n=1
Alg14	p=9.2E-02	n=1
AMP-binding_C	p=9.2E-02	n=2
APP_amyloid	p=9.2E-02	n=1
APP_Cu_bd	p=9.2E-02	n=1
APP_E2	p=9.2E-02	n=1
APP_N	p=9.2E-02	n=1
ARPC4	p=9.2E-02	n=1
ARTD15_N	p=9.2E-02	n=1
Asp_Arg_Hydrox	p=9.2E-02	n=1
Biotin_lipoyl	p=9.2E-02	n=2
BLOC1_2	p=9.2E-02	n=1
BRO1	p=9.2E-02	n=2
CBF_beta	p=9.2E-02	n=1
Cmc1	p=9.2E-02	n=1
Coiled-coil_56	p=9.2E-02	n=1
Complex1_51K	p=9.2E-02	n=1
COQ7	p=9.2E-02	n=1
COX5B	p=9.2E-02	n=1
Cu_amine_oxid	p=9.2E-02	n=1
Cu_amine_oxidN2	p=9.2E-02	n=1
CybS	p=9.2E-02	n=1
Cytochrom_C1	p=9.2E-02	n=1
DDDD	p=9.2E-02	n=1
Drf_FH3	p=9.2E-02	n=2
Drf_GBD	p=9.2E-02	n=2
DS	p=9.2E-02	n=1

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



-log₁₀(p)
n=276/271 input genes with annotations



fraction