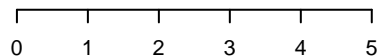


Ncol_Nvec_vc1.1_XM_032373099.2

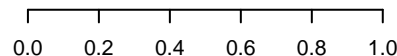
fraction genes in fg and bg

lon_trans	p=1.9E-03	n=9
SNF	p=3.7E-02	n=3
BTB_2	p=6.6E-02	n=5
fn3_4	p=7.1E-02	n=2
Arrestin_C	p=8.0E-02	n=2
Arrestin_N	p=8.0E-02	n=2
PRELI	p=8.0E-02	n=2
6PGD	p=9.0E-02	n=1
Acetyltransf_16	p=9.0E-02	n=1
Aida_C2	p=9.0E-02	n=1
Aida_N	p=9.0E-02	n=1
BAR_3_WASP_bdg	p=9.0E-02	n=1
BNIP3	p=9.0E-02	n=1
Ceramidase	p=9.0E-02	n=1
COX7C	p=9.0E-02	n=1
CTF_NFI	p=9.0E-02	n=1
CUB	p=9.0E-02	n=3
DUF3719	p=9.0E-02	n=1
eIF2_C	p=9.0E-02	n=1
EMI	p=9.0E-02	n=1
Exo_endo_phos_2	p=9.0E-02	n=1
FAIM1	p=9.0E-02	n=1
GF_recep_IV	p=9.0E-02	n=1
GKAP	p=9.0E-02	n=1
ICA69	p=9.0E-02	n=1
IMD	p=9.0E-02	n=1
Lebercilin	p=9.0E-02	n=1
Na_Pi_cotrans	p=9.0E-02	n=1
NAD_binding_2	p=9.0E-02	n=1
Neur_chan_LBD	p=9.0E-02	n=4

fg=0.04	bg=0.01
fg=0.01	bg=0.00
fg=0.02	bg=0.00
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fg=0.00	bg=0.00
fg=0.02	bg=0.00

-log₁₀(p)

n=169/175 input genes with annotations



fraction