

Ncol_Nvec_vc1.1_XM_032386243.2

fraction genes in fg and bg

SMP_LBD	p=6.5E-03	n=3
AcetylCoA_hyd_C	p=1.1E-01	n=1
AcetylCoA_hydro	p=1.1E-01	n=1
Aminotran_3	p=1.1E-01	n=2
AP3D1	p=1.1E-01	n=1
APOBEC_N	p=1.1E-01	n=1
ATP-grasp_3	p=1.1E-01	n=2
Carm_PH	p=1.1E-01	n=1
CARMIL_C	p=1.1E-01	n=1
CCDC66	p=1.1E-01	n=1
Cofac_haem_bdg	p=1.1E-01	n=1
COX7C	p=1.1E-01	n=1
CP2	p=1.1E-01	n=2
Cpn10	p=1.1E-01	n=1
DDDD	p=1.1E-01	n=1
DNA_pol_alpha_N	p=1.1E-01	n=1
Dopey_N	p=1.1E-01	n=1
DUF3752	p=1.1E-01	n=1
DUF4476	p=1.1E-01	n=1
DUF4495	p=1.1E-01	n=1
DUF758	p=1.1E-01	n=1
Elf1	p=1.1E-01	n=1
Elong_Iki1	p=1.1E-01	n=1
FAO_M	p=1.1E-01	n=2
Fasciclin	p=1.1E-01	n=2
FERM_N	p=1.1E-01	n=3
Filamin	p=1.1E-01	n=3
Formyl_trans_C	p=1.1E-01	n=1
GCV_T	p=1.1E-01	n=2
GCV_T_C	p=1.1E-01	n=2

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



-log10(p)

n=290/257 input genes with annotations



fraction