

Elav\_Nvec\_vc1.1\_XM\_032378928.2

fraction genes in fg and bg

3Beta_HSD	p=1.0E-01	n=1
AC_N	p=1.0E-01	n=2
AcetylCoA_hyd_C	p=1.0E-01	n=1
AcetylCoA_hydro	p=1.0E-01	n=1
AdoHcyase	p=1.0E-01	n=1
AdoHcyase_NAD	p=1.0E-01	n=1
Aldo_ket_red	p=1.0E-01	n=2
AP3B1_C	p=1.0E-01	n=1
Biotin_lipoyl_2	p=1.0E-01	n=1
CBX7_C	p=1.0E-01	n=2
CDC50	p=1.0E-01	n=1
CDK5_activator	p=1.0E-01	n=1
Chromo	p=1.0E-01	n=2
CortBP2	p=1.0E-01	n=1
CUPID	p=1.0E-01	n=1
DinB_2	p=1.0E-01	n=1
Dis3l2_C_term	p=1.0E-01	n=1
DOR	p=1.0E-01	n=1
DUF1011	p=1.0E-01	n=1
DUF3543	p=1.0E-01	n=1
DUF4495	p=1.0E-01	n=1
DUF758	p=1.0E-01	n=1
EMP70	p=1.0E-01	n=1
FERM_M	p=1.0E-01	n=3
FXa_inhibition	p=1.0E-01	n=2
HSL_N	p=1.0E-01	n=1
Kelch_1	p=1.0E-01	n=3
KIND	p=1.0E-01	n=1
KLRAQ	p=1.0E-01	n=1
KOW	p=1.0E-01	n=1

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
fg=0.00	bg=0.00
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fg=0.00	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=209/157 input genes with annotations



fraction