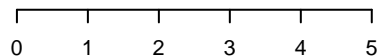


Ncol\_Nvec\_vc1.1\_XM\_001634326.3

fraction genes in fg and bg

CSRNP_N	p=2.2E-02	n=2
AdoHcyase	p=5.4E-02	n=1
AdoHcyase_NAD	p=5.4E-02	n=1
BAR_3_WASP_bdg	p=5.4E-02	n=1
CCDC50_N	p=5.4E-02	n=1
Cystatin	p=5.4E-02	n=1
DSPc	p=5.4E-02	n=2
EIF4E-T	p=5.4E-02	n=1
Fcf2	p=5.4E-02	n=1
Fic	p=5.4E-02	n=1
Indigoidine_A	p=5.4E-02	n=1
Inhibitor_I29	p=5.4E-02	n=2
JTB	p=5.4E-02	n=1
LBP_BPI_CETP	p=5.4E-02	n=1
MTA_R1	p=5.4E-02	n=1
PAN_1	p=5.4E-02	n=3
ParBc	p=5.4E-02	n=1
Peptidase_C1	p=5.4E-02	n=2
Peptidase_S49	p=5.4E-02	n=1
Peroxin-3	p=5.4E-02	n=1
PIF1	p=5.4E-02	n=1
PUA	p=5.4E-02	n=1
Ras	p=5.4E-02	n=4
Rhodanese	p=5.4E-02	n=2
Skp1	p=5.4E-02	n=1
Somatomedin_B	p=5.4E-02	n=2
Tom5	p=5.4E-02	n=1
TPR_14	p=5.4E-02	n=1
UNC45-central	p=5.4E-02	n=1
7tm_2	p=6.7E-02	n=2

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



-log<sub>10</sub>(p)  
n=111/112 input genes with annotations



fraction