

Ncol\_Nvec\_vc1.1\_XM\_001635015.3

fraction genes in fg and bg

Aida_C2	p=6.7E-02	n=1
Aida_N	p=6.7E-02	n=1
ATP_synt_F	p=6.7E-02	n=1
BAR_3_WASP_bdg	p=6.7E-02	n=1
cNMP_binding	p=6.7E-02	n=3
Complex1_49kDa	p=6.7E-02	n=1
Cu_amine_oxid	p=6.7E-02	n=1
Cu_amine_oxidN2	p=6.7E-02	n=1
DER1	p=6.7E-02	n=1
DS	p=6.7E-02	n=1
DUF2205	p=6.7E-02	n=1
DUF572	p=6.7E-02	n=1
EF-hand_7	p=6.7E-02	n=4
Gelsolin	p=6.7E-02	n=2
GFA	p=6.7E-02	n=1
HAUS4	p=6.7E-02	n=1
Ion_trans	p=6.7E-02	n=5
Peroxin-3	p=6.7E-02	n=1
Phospholip_A2_3	p=6.7E-02	n=1
Serinc	p=6.7E-02	n=2
Snurportin1	p=6.7E-02	n=1
SPRY	p=6.7E-02	n=2
Tfb2	p=6.7E-02	n=1
Tfb2_C	p=6.7E-02	n=1
Tim29	p=6.7E-02	n=1
TRAP_beta	p=6.7E-02	n=1
VHP	p=6.7E-02	n=2
YhhN	p=6.7E-02	n=1
bZIP_1	p=7.9E-02	n=2
SET	p=7.9E-02	n=2

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
fg=0.01	bg=0.00
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fg=0.01	bg=0.00
fg=0.03	bg=0.01
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.03	bg=0.01
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=0.01	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



-log<sub>10</sub>(p)  
n=118/120 input genes with annotations



fraction