

## Ncol\_Nvec\_vc1.1\_XM\_032383717.2

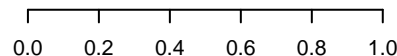
## fraction genes in fg and bg

GTP_EFTU_D2	p=3.1E-02	n=5
GTP_EFTU	p=3.7E-02	n=5
2Fe-2S_thioredx	p=1.2E-01	n=1
AA_kinase	p=1.2E-01	n=1
AAA	p=1.2E-01	n=5
AAA_assoc_2	p=1.2E-01	n=1
AHD	p=1.2E-01	n=1
ALG11_N	p=1.2E-01	n=1
Alg14	p=1.2E-01	n=1
Archease	p=1.2E-01	n=1
ASF1_hist_chap	p=1.2E-01	n=1
Avl9	p=1.2E-01	n=1
AZUL	p=1.2E-01	n=1
B3_4	p=1.2E-01	n=1
B5	p=1.2E-01	n=1
BACK	p=1.2E-01	n=6
BTB	p=1.2E-01	n=6
bZIP_1	p=1.2E-01	n=3
bZIP_2	p=1.2E-01	n=3
C1_2	p=1.2E-01	n=1
CAP-ZIP_m	p=1.2E-01	n=1
CBF_beta	p=1.2E-01	n=1
CBFD_NFYB_HMF	p=1.2E-01	n=2
CFAP298	p=1.2E-01	n=1
CKS	p=1.2E-01	n=1
CMS1	p=1.2E-01	n=1
CSN5_C	p=1.2E-01	n=1
CTU2	p=1.2E-01	n=1
CUPID	p=1.2E-01	n=1
DHHA1	p=1.2E-01	n=1

fg=0.01	bg=0.00
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-log<sub>10</sub>(p)

n=412/394 input genes with annotations



fraction