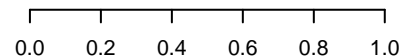


AcetylCoA_hyd_C	p=8.0E-02	n=1
AcetylCoA_hydro	p=8.0E-02	n=1
AdoHcyase	p=8.0E-02	n=1
AdoHcyase_NAD	p=8.0E-02	n=1
AP3D1	p=8.0E-02	n=1
ATP-grasp_3	p=8.0E-02	n=2
ATP-synt_F	p=8.0E-02	n=1
BCCT	p=8.0E-02	n=1
BTK	p=8.0E-02	n=1
CCDC66	p=8.0E-02	n=1
COX7C	p=8.0E-02	n=1
CP2	p=8.0E-02	n=2
Dopey_N	p=8.0E-02	n=1
DUF4476	p=8.0E-02	n=1
DUF908	p=8.0E-02	n=1
DUF913	p=8.0E-02	n=1
EMP70	p=8.0E-02	n=1
GGACT	p=8.0E-02	n=1
INTS5_C	p=8.0E-02	n=1
INTS5_N	p=8.0E-02	n=1
Mito_carr	p=8.0E-02	n=3
MTA_R1	p=8.0E-02	n=1
MTABC_N	p=8.0E-02	n=1
MutS_IV	p=8.0E-02	n=1
Na_Pi_cotrans	p=8.0E-02	n=1
NAGidase	p=8.0E-02	n=1
OSR1_C	p=8.0E-02	n=1
P66_CC	p=8.0E-02	n=1
PEX11	p=8.0E-02	n=1
PNP_phzG_C	p=8.0E-02	n=1

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



$-\log_{10}(p)$
n=179/145 input genes with annotations



fraction