

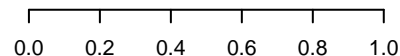
Acetyltransf_13	p=9.5E-02	n=1
Actin	p=9.5E-02	n=2
Adaptin_binding	p=9.5E-02	n=1
AlbA_2	p=9.5E-02	n=1
AMP-binding	p=9.5E-02	n=2
ANAPC1	p=9.5E-02	n=1
APC1_C	p=9.5E-02	n=1
APG12	p=9.5E-02	n=1
Asp_protease	p=9.5E-02	n=1
BCCT	p=9.5E-02	n=1
BCL_N	p=9.5E-02	n=1
CBF_beta	p=9.5E-02	n=1
CDC50	p=9.5E-02	n=1
cNMP_binding	p=9.5E-02	n=3
Coatmer_g_Cpla	p=9.5E-02	n=1
Complex1_51K	p=9.5E-02	n=1
COP-gamma_platf	p=9.5E-02	n=1
Cript	p=9.5E-02	n=1
Cse1	p=9.5E-02	n=1
Dis3l2_C_term	p=9.5E-02	n=1
DUF1908	p=9.5E-02	n=1
DUF4200	p=9.5E-02	n=1
DUF4457	p=9.5E-02	n=1
DUF92	p=9.5E-02	n=1
EF-hand_like	p=9.5E-02	n=2
eIF-1a	p=9.5E-02	n=1
Endosulfine	p=9.5E-02	n=1
Ephrin	p=9.5E-02	n=2
FANCI_HD1	p=9.5E-02	n=1
FANCI_HD2	p=9.5E-02	n=1

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-log10(p)

n=231/171 input genes with annotations



fraction