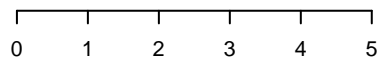
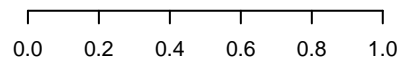


2Fe-2S_thioredx	p=1.0E-01	n=1
Alg14	p=1.0E-01	n=1
Alpha-2-MRAP_C	p=1.0E-01	n=1
Alpha-2-MRAP_N	p=1.0E-01	n=1
Amino_oxidase	p=1.0E-01	n=2
ANAPC5	p=1.0E-01	n=1
Arm_vesicular	p=1.0E-01	n=1
ATP-grasp_2	p=1.0E-01	n=1
ATP-synt_F	p=1.0E-01	n=1
BTB_2	p=1.0E-01	n=4
CBF_beta	p=1.0E-01	n=1
CDO_I	p=1.0E-01	n=1
CG-1	p=1.0E-01	n=1
DUF1242	p=1.0E-01	n=2
DUF4505	p=1.0E-01	n=1
eIF-1a	p=1.0E-01	n=1
ELM2	p=1.0E-01	n=2
F5_F8_type_C	p=1.0E-01	n=4
Glycohydro_20b2	p=1.0E-01	n=1
GOLD_2	p=1.0E-01	n=1
Headcase	p=1.0E-01	n=1
HECA	p=1.0E-01	n=1
Hexapep	p=1.0E-01	n=1
HSL_N	p=1.0E-01	n=1
Ion_trans	p=1.0E-01	n=7
KCT2	p=1.0E-01	n=1
Leg1	p=1.0E-01	n=1
Ligase_CoA	p=1.0E-01	n=1
Lyase_aromatic	p=1.0E-01	n=1
Mab-21	p=1.0E-01	n=2

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 $-\log_{10}(p)$ 

n=214/210 input genes with annotations



fraction