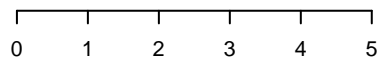
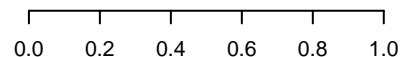


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APG9	p=9.7E-02	n=1
ARS2	p=9.7E-02	n=1
ATP-grasp_2	p=9.7E-02	n=1
BAT2_N	p=9.7E-02	n=1
BORCS8	p=9.7E-02	n=1
COesterase	p=9.7E-02	n=2
COQ7	p=9.7E-02	n=1
DENND11	p=9.7E-02	n=1
DNA_repr_REX1B	p=9.7E-02	n=1
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EIF4E-T	p=9.7E-02	n=1
Fascin	p=9.7E-02	n=1
Fer2_3	p=9.7E-02	n=1
Fer4_17	p=9.7E-02	n=1
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Forkhead	p=9.7E-02	n=3
FOXO-TAD	p=9.7E-02	n=1
Galactosyl_T	p=9.7E-02	n=3
GKAP	p=9.7E-02	n=1
Glyco_transf_8	p=9.7E-02	n=2
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$-\log_{10}(p)$
n=220/223 input genes with annotations



fraction