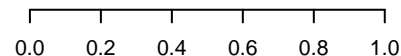
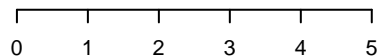


gc002a

fraction genes in fg and bg

ECH_1	p=1.2E-30	n=39
Acyl-CoA_dh_N	p=1.4E-24	n=37
Acyl-CoA_dh_M	p=1.5E-23	n=43
Acyl-CoA_dh_1	p=1.2E-21	n=37
adh_short	p=9.5E-19	n=63
ATP-synt_ab	p=3.3E-15	n=15
ATP-synt_ab_N	p=3.3E-15	n=15
ATP-synt_C	p=3.1E-13	n=14
Thiolase_C	p=4.4E-12	n=16
Thiolase_N	p=4.4E-12	n=16
Mito_carr	p=1.0E-11	n=36
Ligase_CoA	p=1.4E-11	n=13
Glyoxalase	p=7.7E-11	n=11
2-oxoacid_dh	p=1.5E-10	n=12
adh_short_C2	p=4.3E-10	n=18
Iso_dh	p=4.3E-10	n=13
Abhydrolase_1	p=2.4E-09	n=32
Complex1_LYR	p=5.8E-09	n=13
COesterase	p=8.9E-09	n=26
E3_binding	p=1.0E-08	n=9
3HCDH	p=1.4E-08	n=8
Acyltransf_C	p=1.4E-08	n=10
L51_S25_Cl-B8	p=1.4E-08	n=10
Redoxin	p=1.4E-08	n=8
Aldedh	p=8.2E-08	n=18
Lig_chan-Glu_bd	p=1.2E-07	n=16
Lig_chan	p=1.8E-07	n=16
zf-Tim10_DDP	p=3.1E-07	n=11
ADH_N	p=4.3E-07	n=21
Biotin_lipoyl	p=7.8E-07	n=12

fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.01	bg=0.00
fg=0.02	bg=0.00
fg=0.00	bg=0.00
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fg=0.00	bg=0.00
fg=0.01	bg=0.00
fg=0.00	bg=0.00



-log₁₀(p)
n=1110/3420 input genes with annotations

fraction