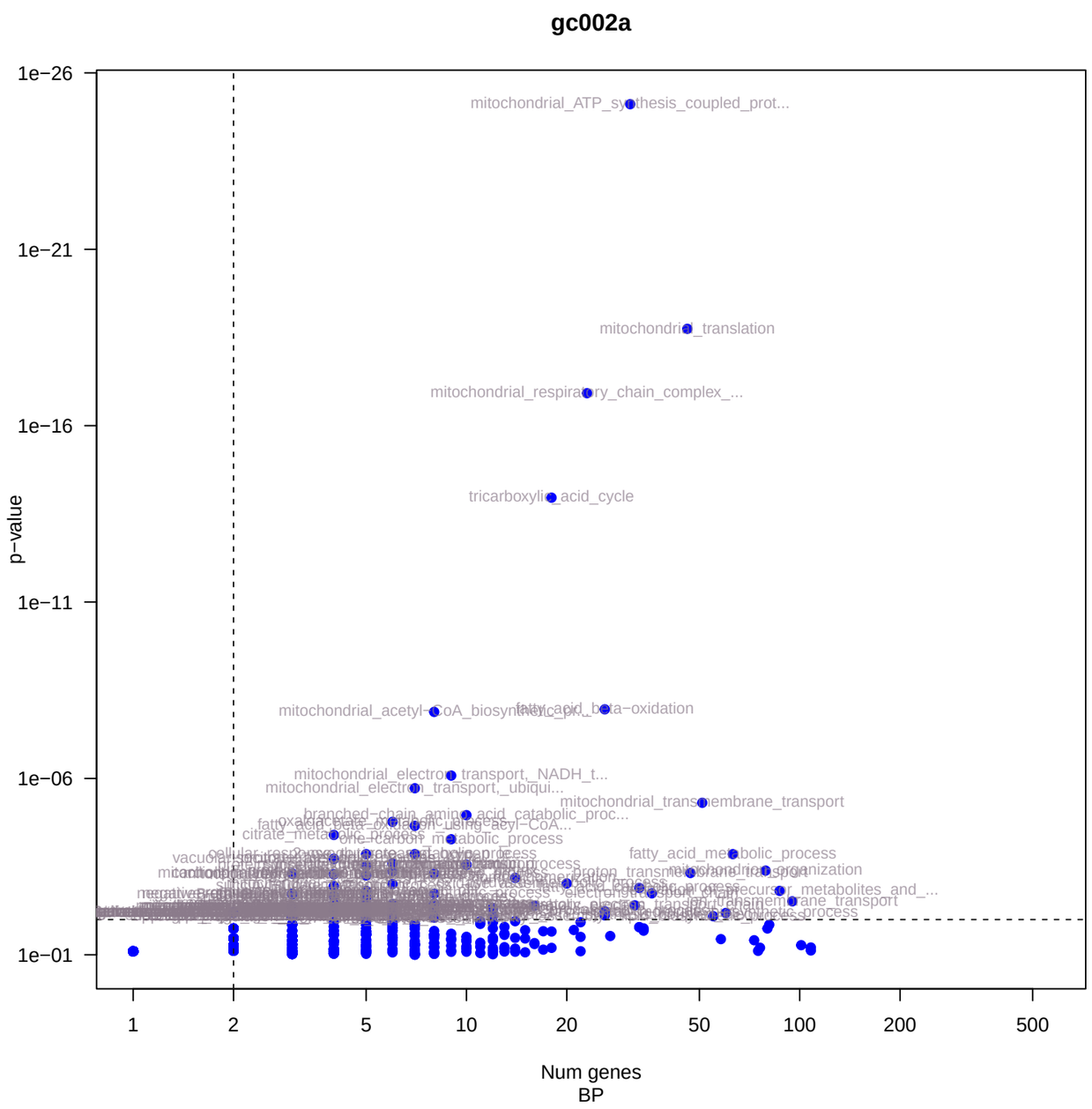
gc002a
p−value
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1e−16
1e−11
1e−06
1e−01
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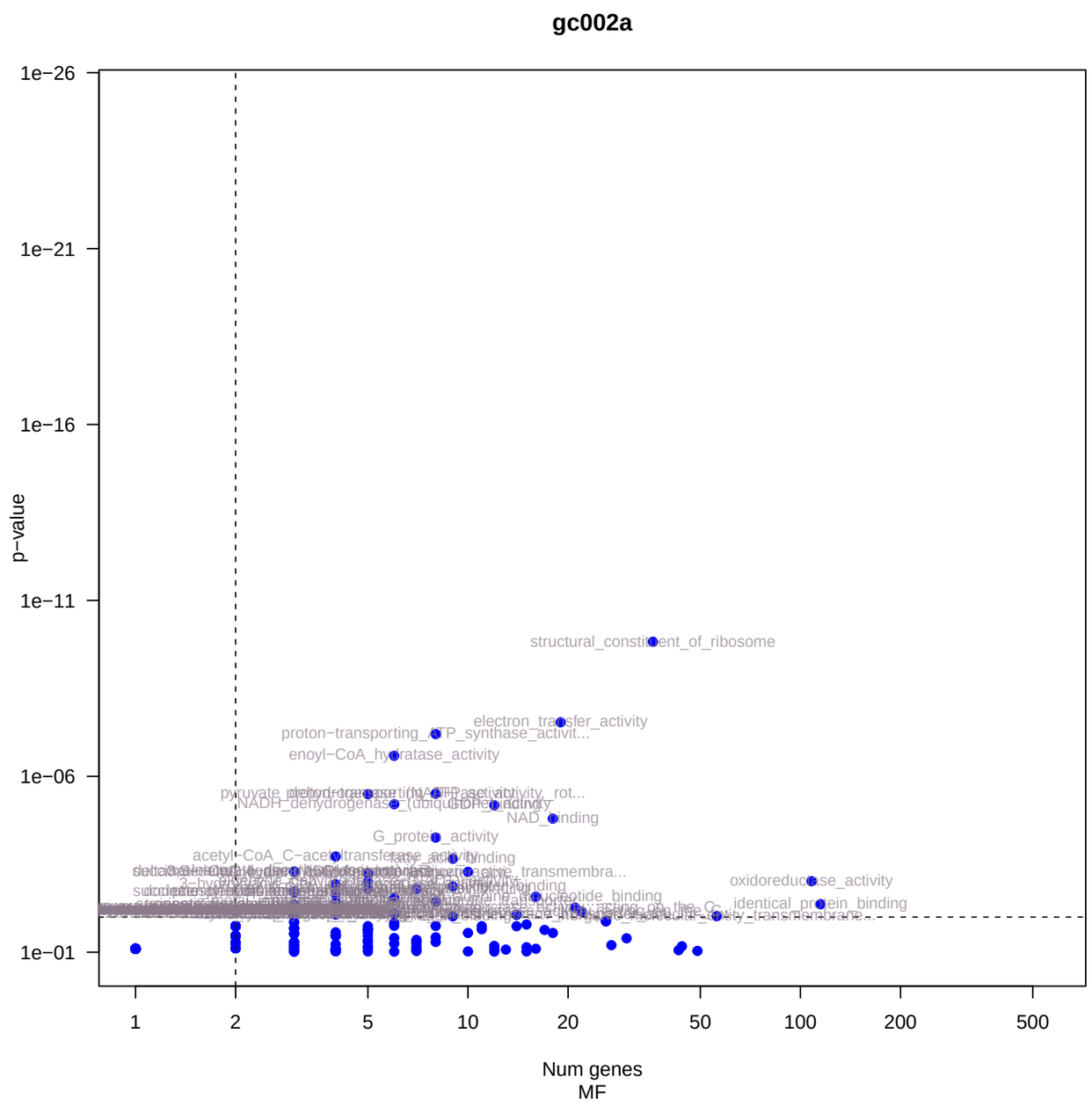
mitochondrial_translation
mitochondrial_respiratory_chain_complex_...
tricarboxylic_acid_cycle
mitochondrial_acetyl−CoA_biosynthetic_pr...
fatty_acid_beta−oxidation
mitochondrial_electron_transport,_NADH_t...
mitochondrial_electron_transport,_ubiqui...
mitochondrial_transmembrane_transport
branched−chain_amino_acid_catabolic_proc...
fatty_acid_metabolic_process
mitochondrion_organization
proton_transmembrane_transport
precursor_metabolites_and_...
1
2
5
10
20
50
100
200
500
Num genes
BP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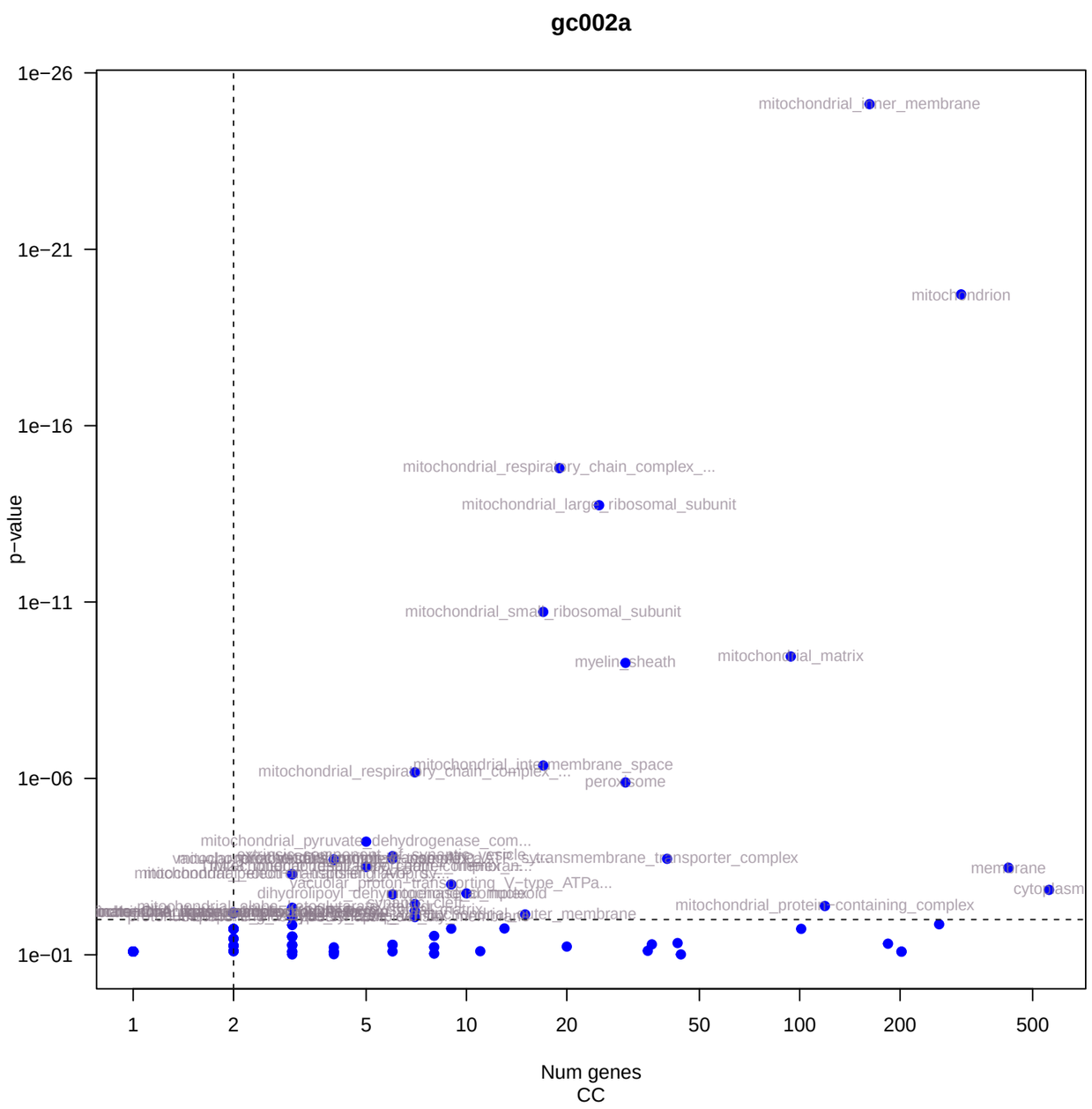
electron_transfer_activity
proton−transporting_ATP_synthase_activit...
enoyl−CoA_hydratase_activity
NAD_binding
G_protein_activity
oxidoreductase_activity
identical_protein_binding
1
2
5
10
20
50
100
200
500
Num genes
MF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1e−16
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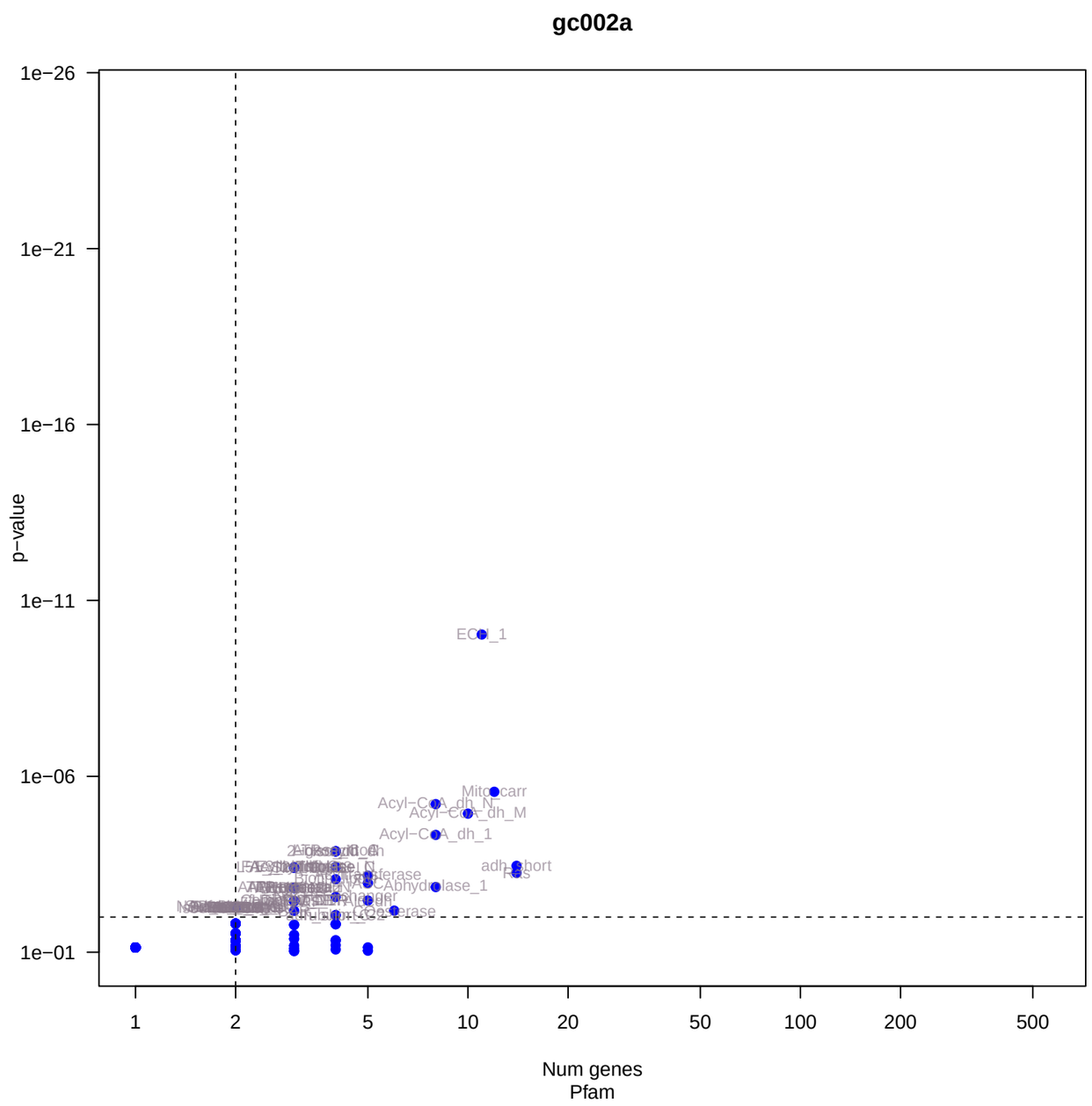
mitochondrion
mitochondrial_respiratory_chain_complex_...
mitochondrial_large_ribosomal_subunit
mitochondrial_small_ribosomal_subunit
myelin_sheath
mitochondrial_matrix
mitochondrial_intermembrane_space
peroxisome
mitochondrial_pyruvate_dehydrogenase_com...
membrane
cytoplasm
mitochondrial_protein_containing_complex
1
2
5
10
20
50
100
200
500
Num genes
CC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1e−21
1e−16
1e−11
1e−06
1e−01
ECH_1
Mitocarr
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Acyl−CoA_dh_M
Acyl−CoA_dh_1
adh_short
1
2
5
10
20
50
100
200
500
Num genes
Pfam