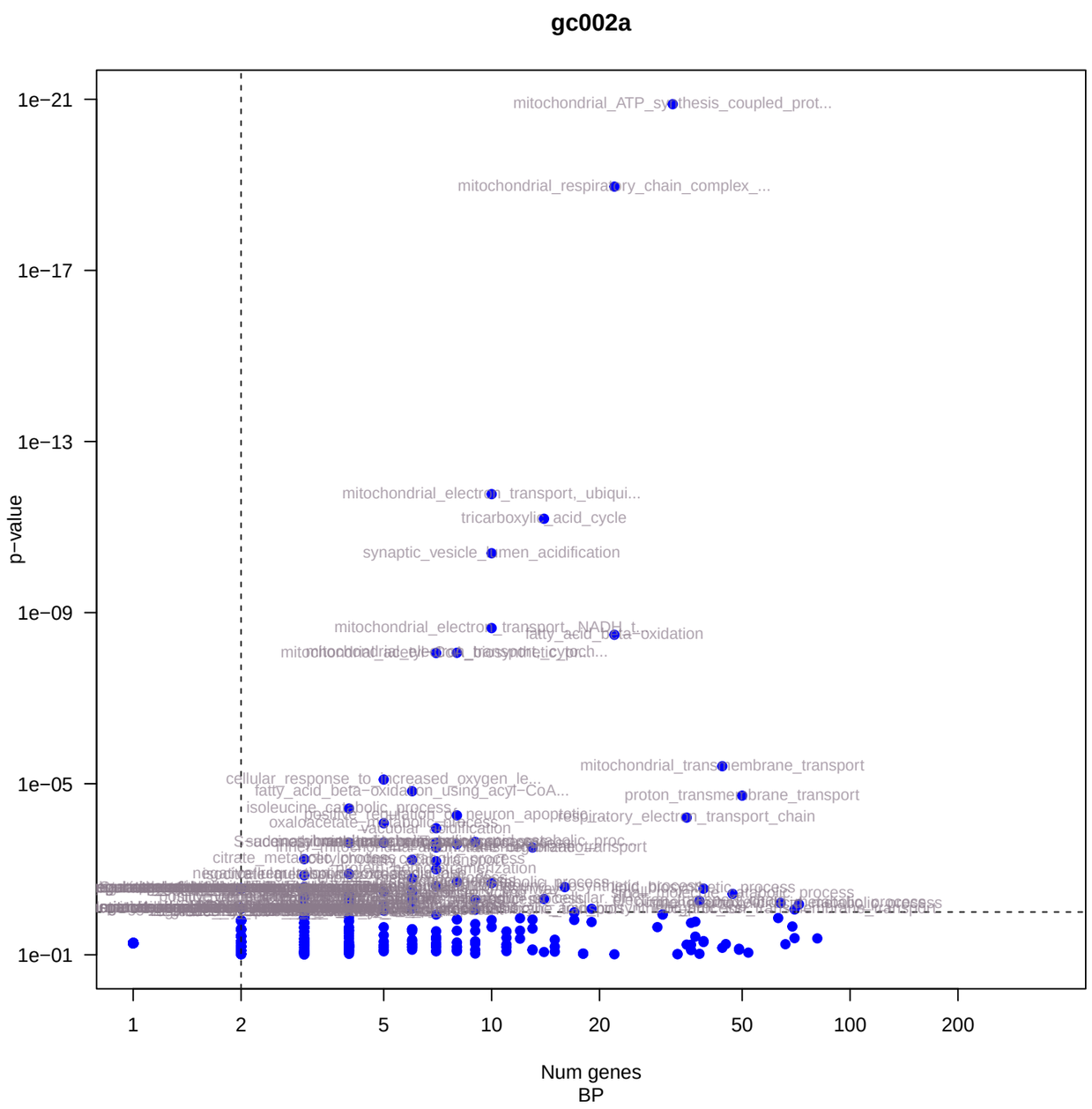
gc002a
p−value
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1e−13
1e−09
1e−05
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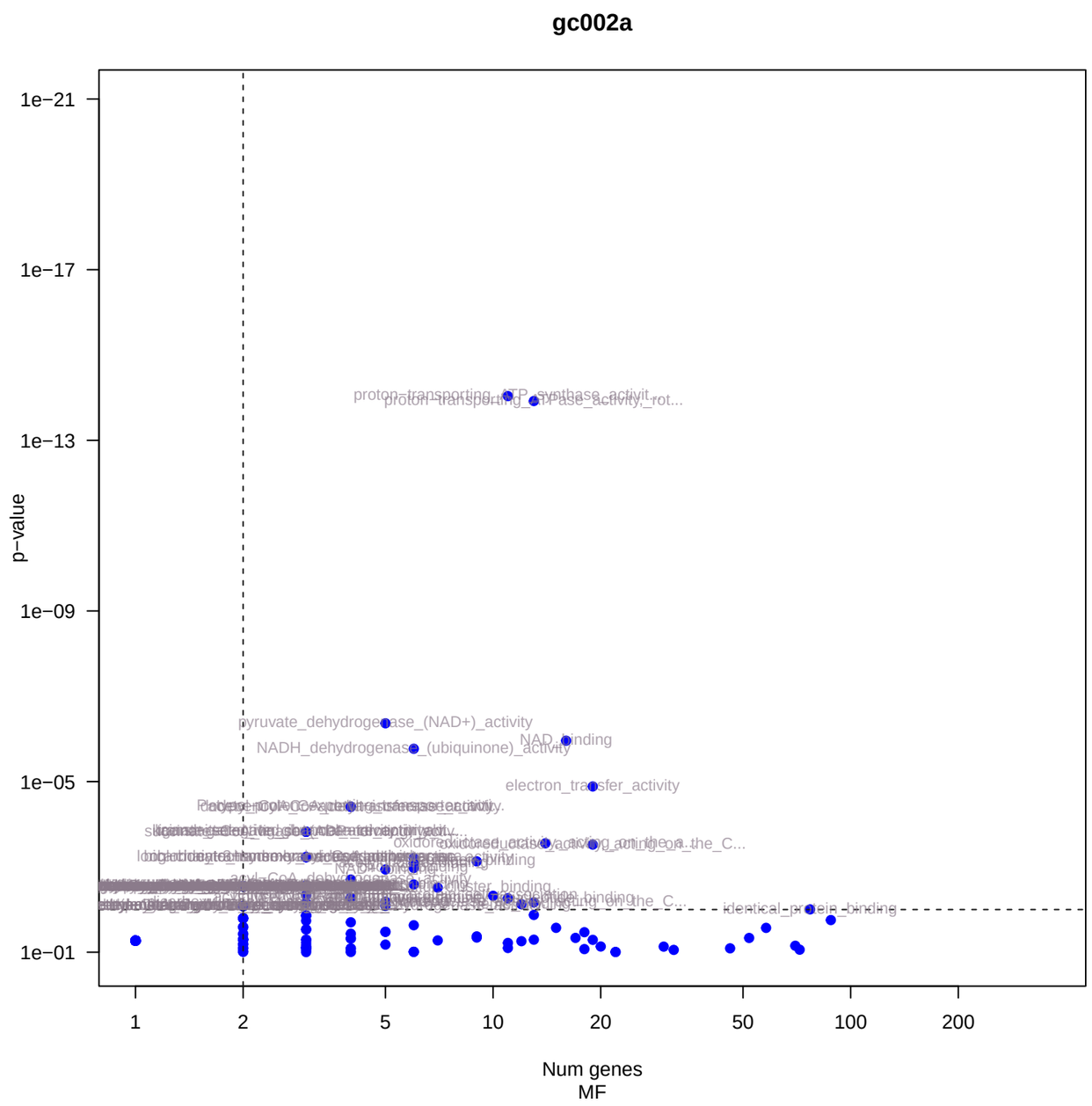
mitochondrial_respiratory_chain_complex_...
mitochondrial_electron_transport,_ubiqui...
tricarboxylic_acid_cycle
synaptic_vesicle_lumen_acidification
mitochondrial_electron_transport,_NADH_t...
fatty_acid_beta−oxidation
mitochondrial_transmembrane_transport
cellular_response_to_increased_oxygen_le...
fatty_acid_beta−oxidation_using_acyl−CoA...
proton_transmembrane_transport
isoleucine_catabolic_process
respiratory_electron_transport_chain
oxaloacetate_metabolic_process
citrate_metabolic_process
1
2
5
10
20
50
100
200
Num genes
BP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1e−13
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1e−05
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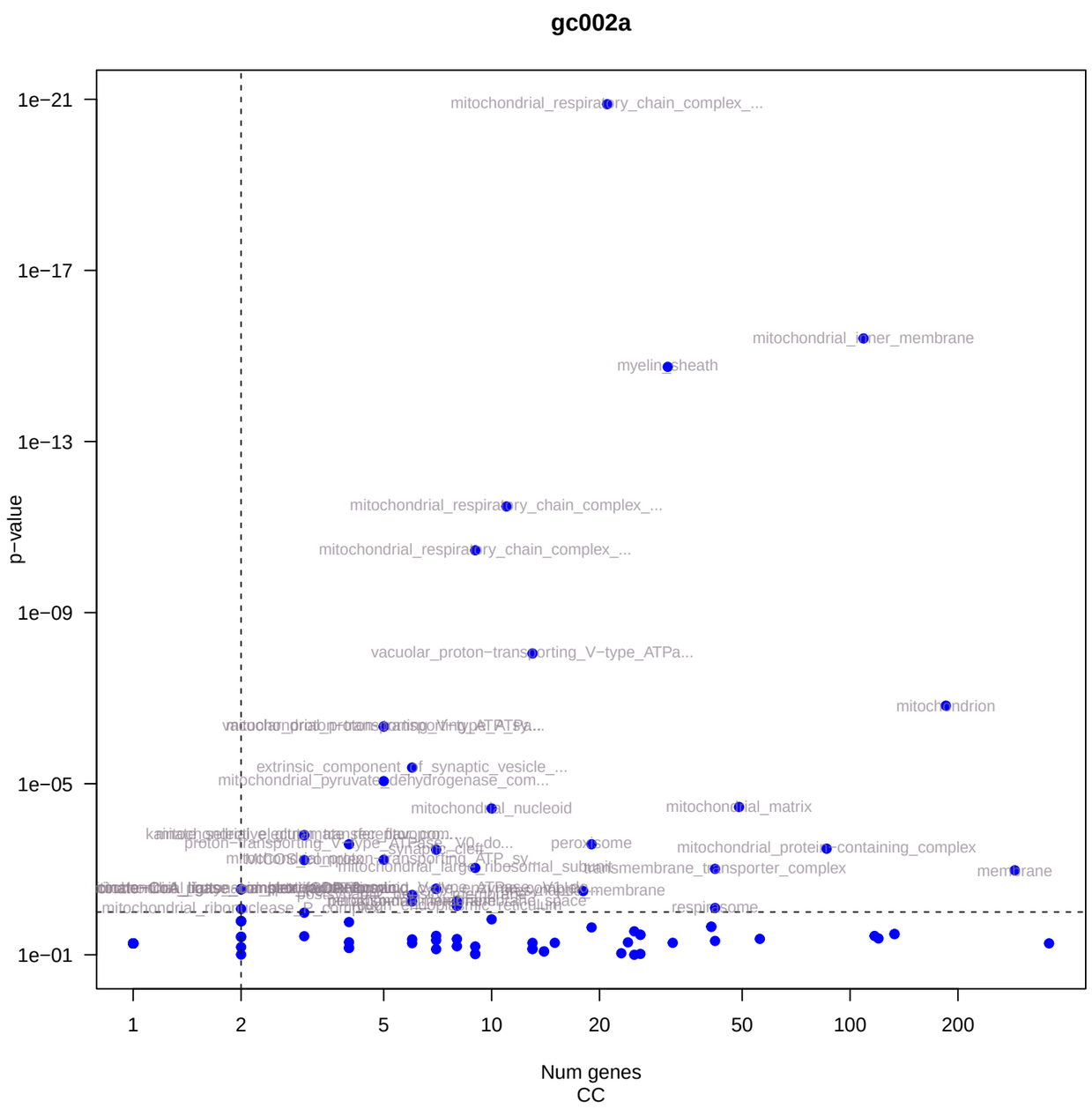
pyruvate_dehydrogenase_(NAD+)_activity
NAD_binding
NADH_dehydrogenase_(ubiquinone)_activity
electron_transfer_activity
identical_protein_binding
1
2
5
10
20
50
100
200
Num genes
MF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1e−13
1e−09
1e−05
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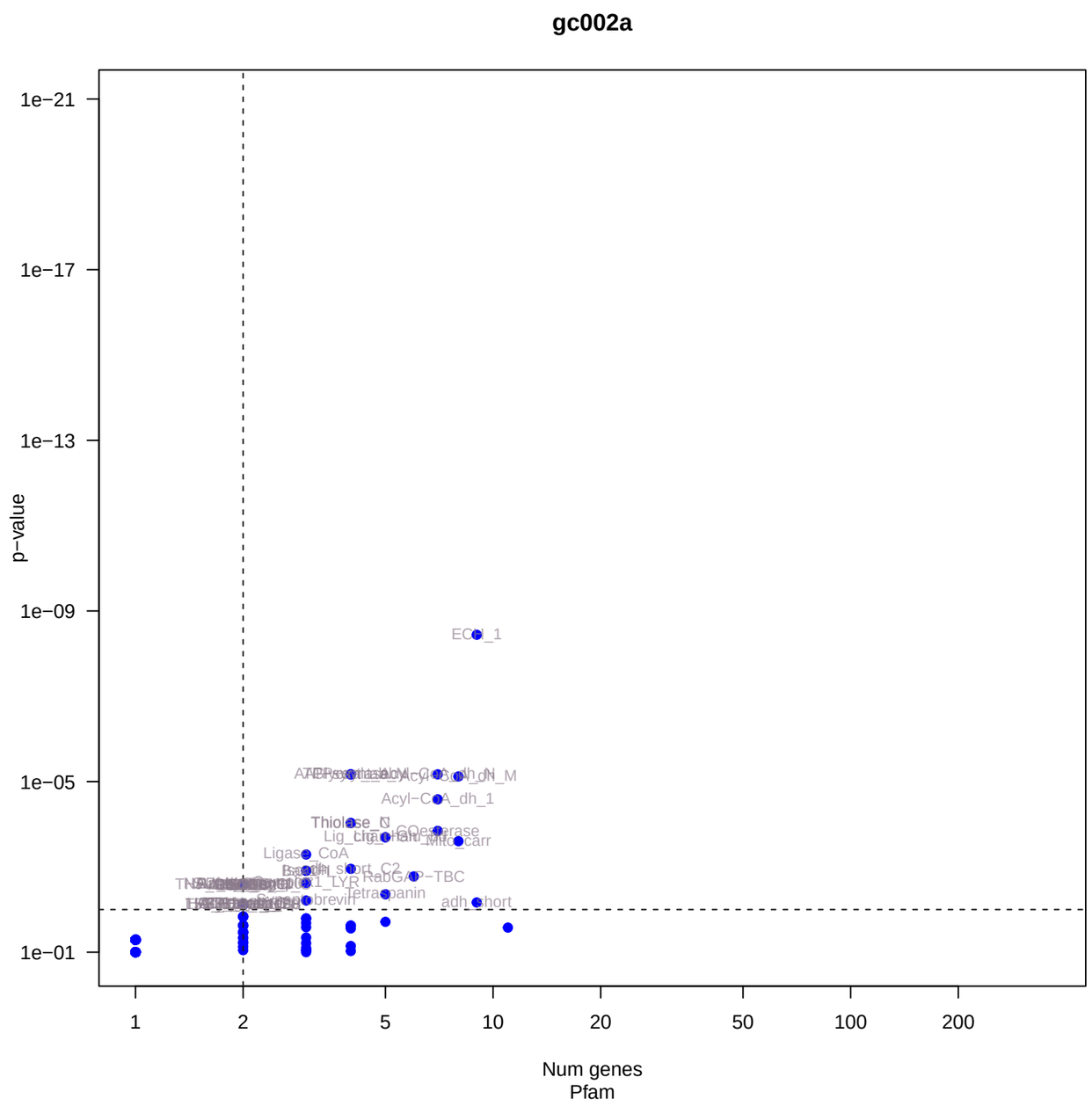
mitochondrial_inner_membrane
myelin_sheath
mitochondrial_respiratory_chain_complex_...
mitochondrial_respiratory_chain_complex_...
vacuolar_proton−transporting_V−type_ATPa...
mitochondrion
extrinsic_component_of_synaptic_vesicle_...
mitochondrial_pyruvate_dehydrogenase_com...
mitochondrial_nucleoid
mitochondrial_matrix
peroxisome
mitochondrial_protein_containing_complex
membrane
respirasome
1
2
5
10
20
50
100
200
Num genes
CC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1e−17
1e−13
1e−09
1e−05
1e−01
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Acyl−CoA_dh_1
Ligase_CoA
Tetraspanin
adh_short
1
2
5
10
20
50
100
200
Num genes
Pfam