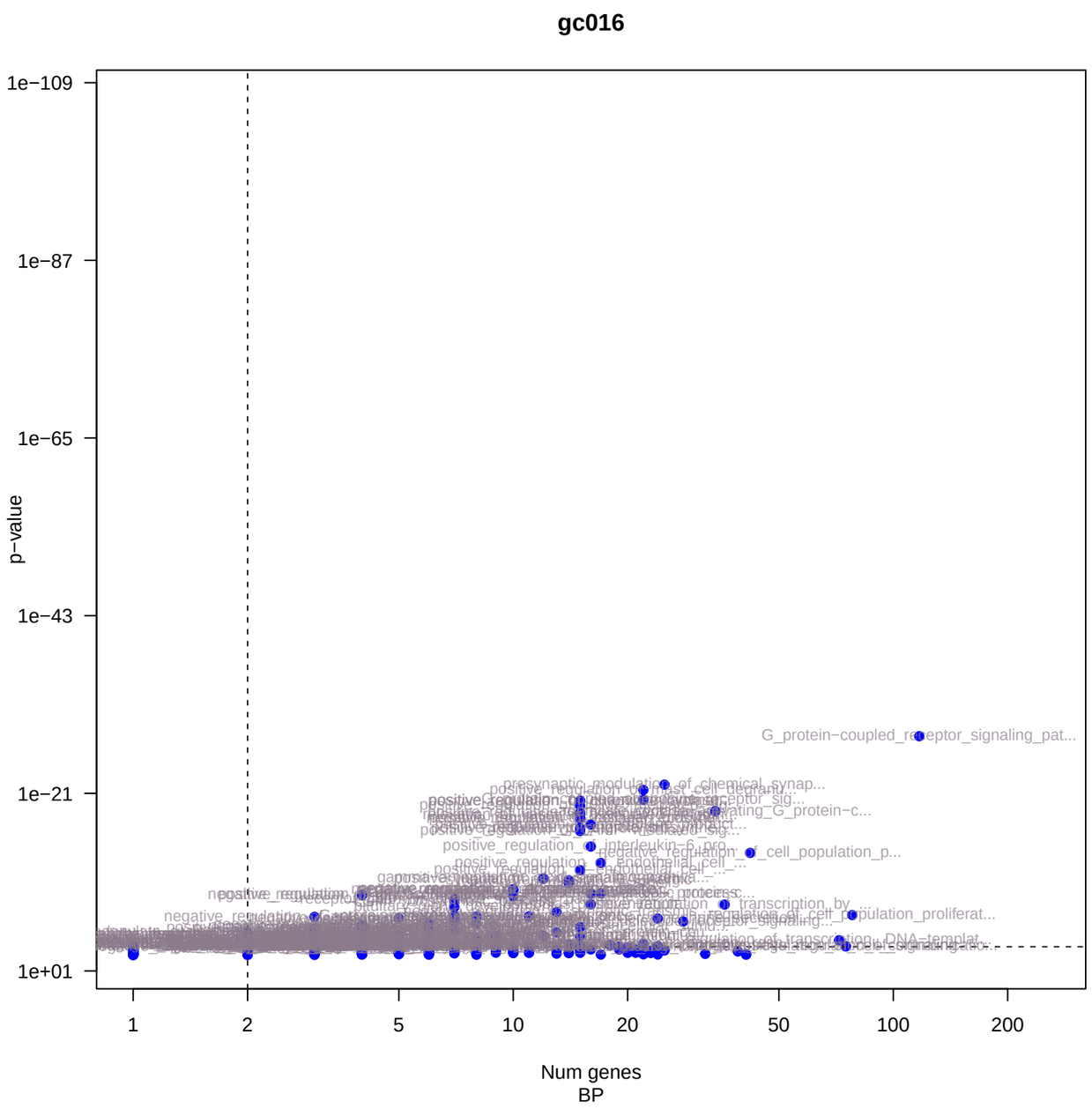
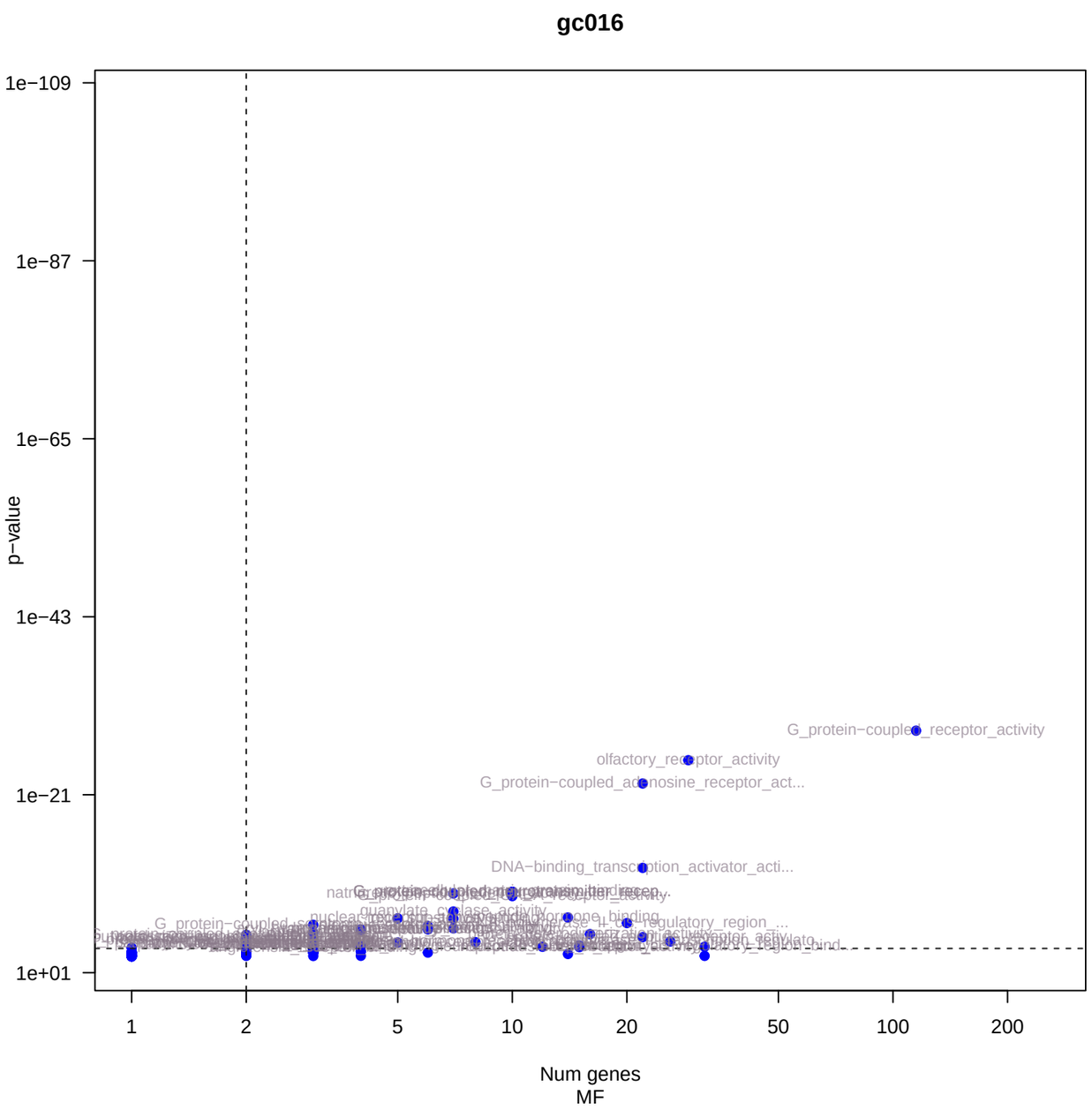
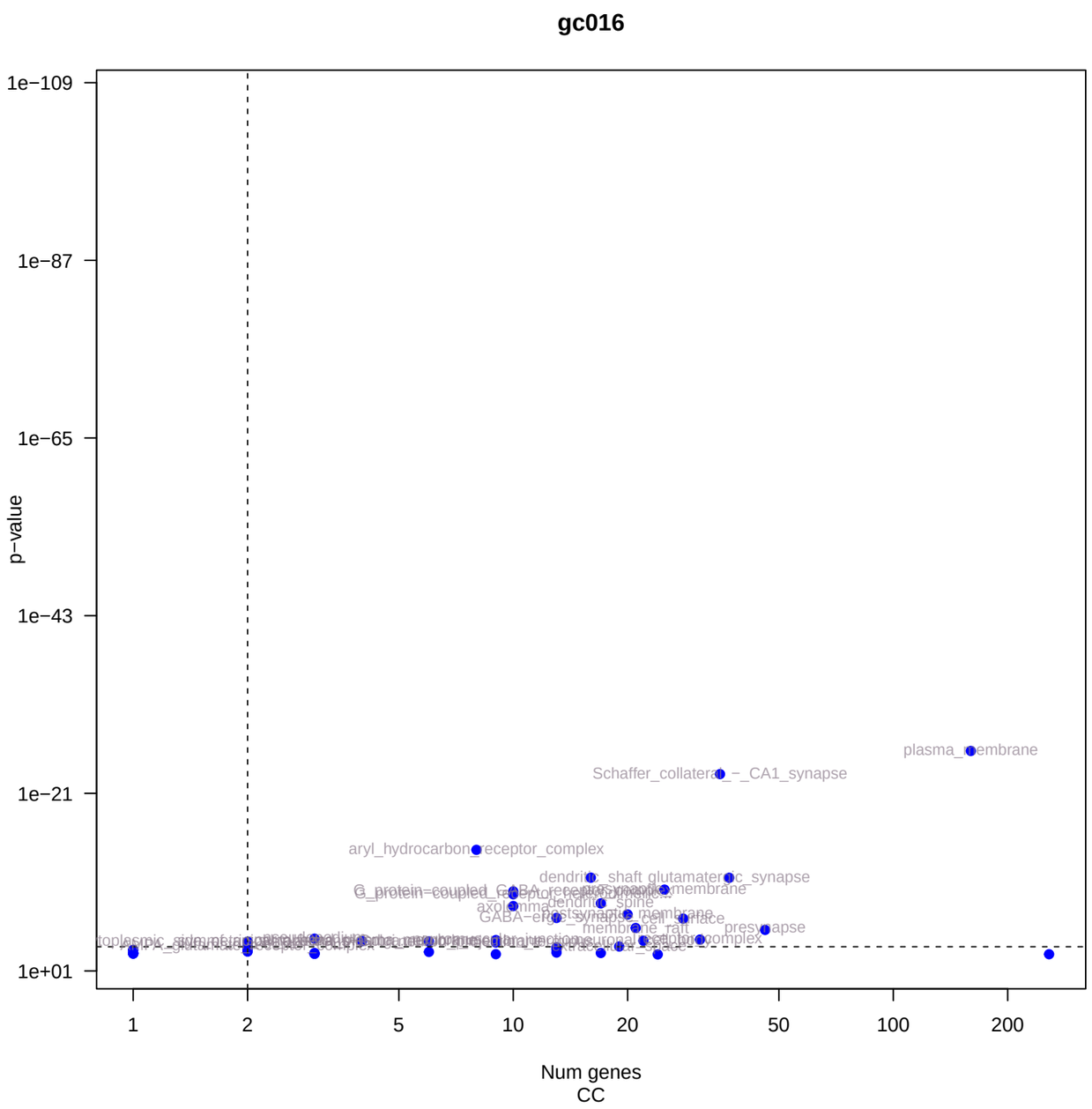
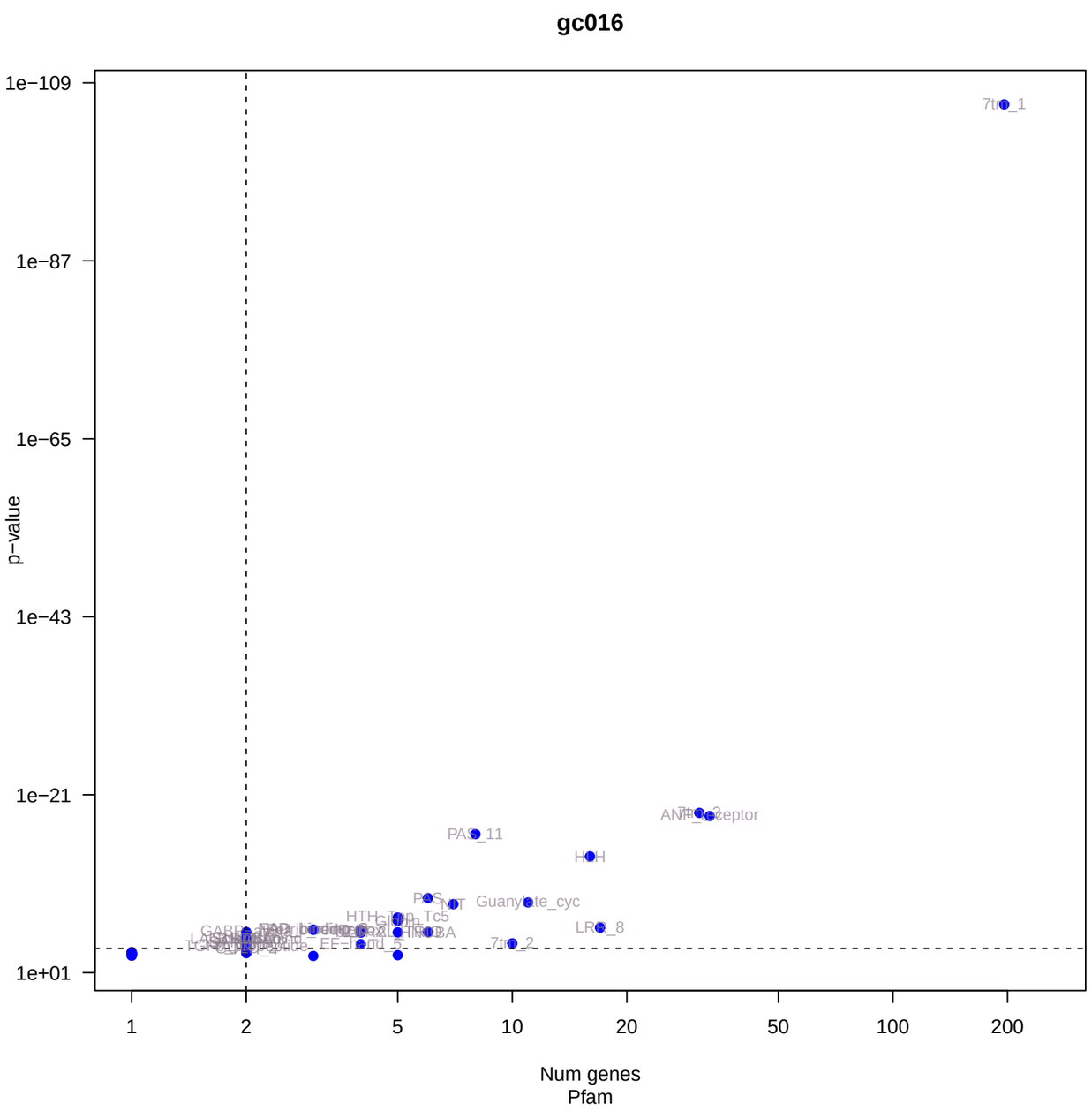

gc016
p−value
1e−109
1e−87
1e−65
1e−43
1e−21
1e+01
G_protein−coupled_receptor_signaling_pat...
presynaptic_modulation_of_chemical_synap...
positive_regulation_of_interleukin−6_pro...
negative_regulation_of_cell_population_p...
positive_regulation_of_endothelial_cell_...
transcription_by_...
cell_population_proliferat...
1
2
5
10
20
50
100
200
Num genes
BP


gc016
p−value
1e−109
1e−87
1e−65
1e−43
1e−21
1e+01
G_protein−coupled_receptor_activity
olfactory_receptor_activity
G_protein−coupled_adenosine_receptor_act...
DNA−binding_transcription_activator_acti...
quanylate_cyclase_activity
G_protein−coupled_peptide_receptor_activity
1
2
5
10
20
50
100
200
Num genes
MF


gc016
p−value
1e−109
1e−87
1e−65
1e−43
1e−21
1e+01
plasma_membrane
Schaffer_collateral_−_CA1_synapse
aryl_hydrocarbon_receptor_complex
dendritic_shaft_glutamatergic_synapse
G_protein−coupled_GABA_receptor_complex
axolemma
dendritic_spine
GABA−ergic_synapse
postsynaptic_membrane
membrane_raft
presynapse
1
2
5
10
20
50
100
200
Num genes
CC


gc016
p−value
1e−109
1e−87
1e−65
1e−43
1e−21
1e+01
7tm_1
7tm_3
PAS_11
HOH
Guanylate_cyc
HTH_Tc5
LRR_8
1
2
5
10
20
50
100
200
Num genes
Pfam