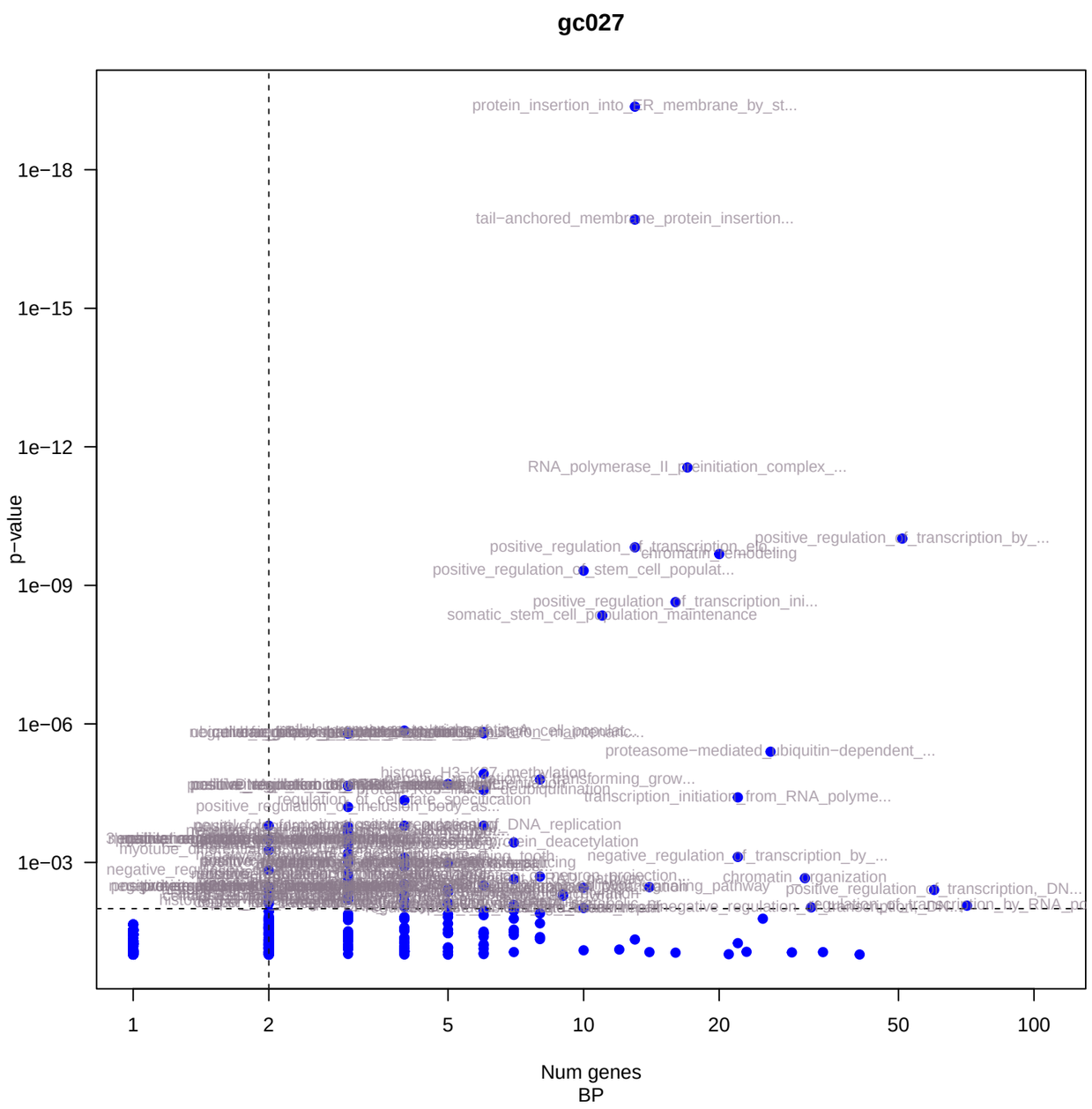
gc027
p−value
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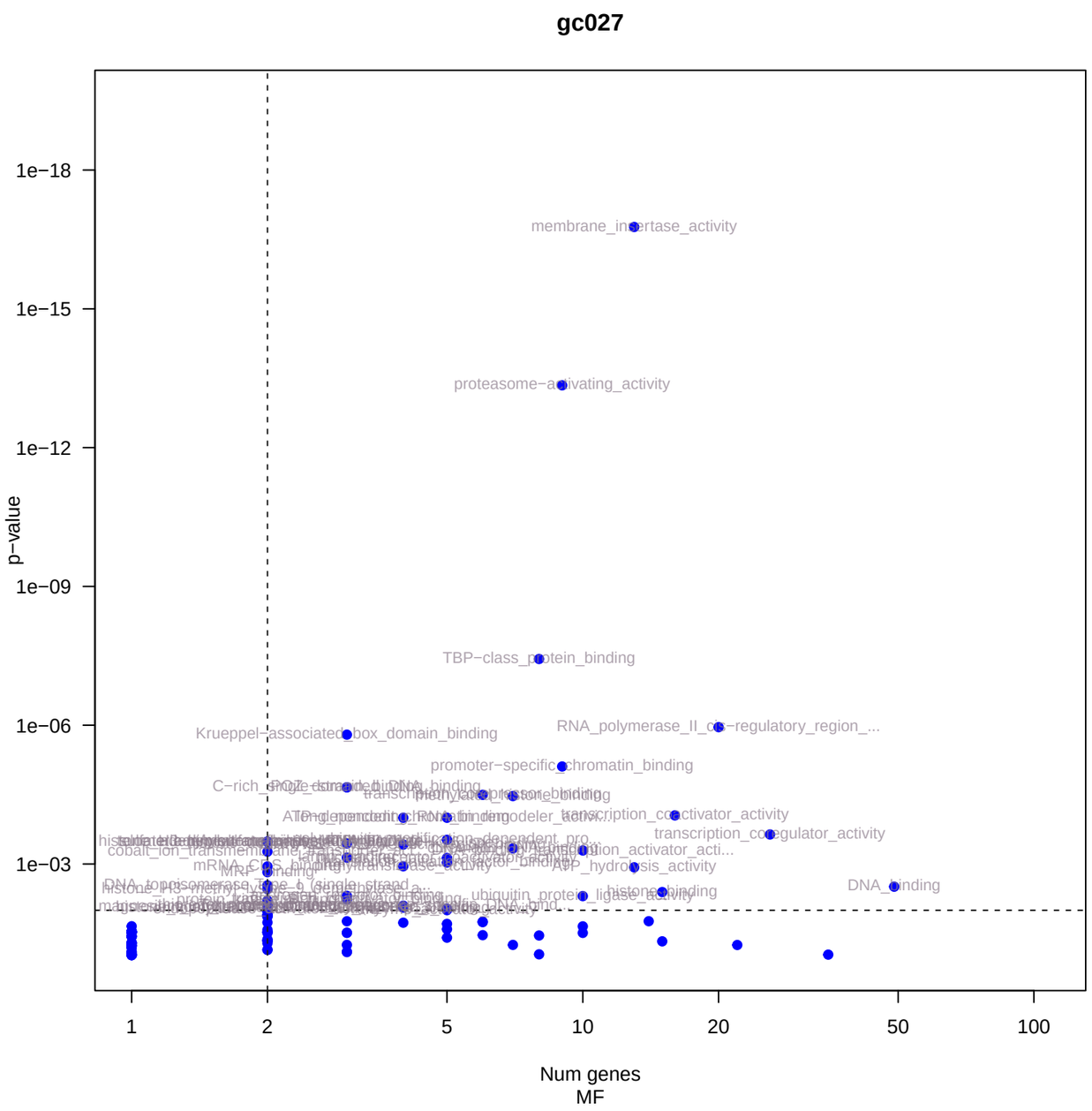
protein_insertion_into_ER_membrane_by_st...
tail−anchored_membrane_protein_insertion...
RNA_polymerase_II_preinitiation_complex_...
positive_regulation_of_transcription_by_...
positive_regulation_of_transcription_ele...
chromatin_remodeling
positive_regulation_of_stem_cell_populat...
positive_regulation_of_transcription_ini...
somatic_stem_cell_population_maintenance
proteasome−mediated_ubiquitin−dependent_...
histone_H3−K27_methylation
transcription_initiation_from_RNA_polyme...
positive_regulation_of_DNA_replication
negative_regulation_of_transcription_by_...
chromatin_organization
positive_regulation_of_transcription,_DN...
1
2
5
10
20
50
100
Num genes
BP


gc027
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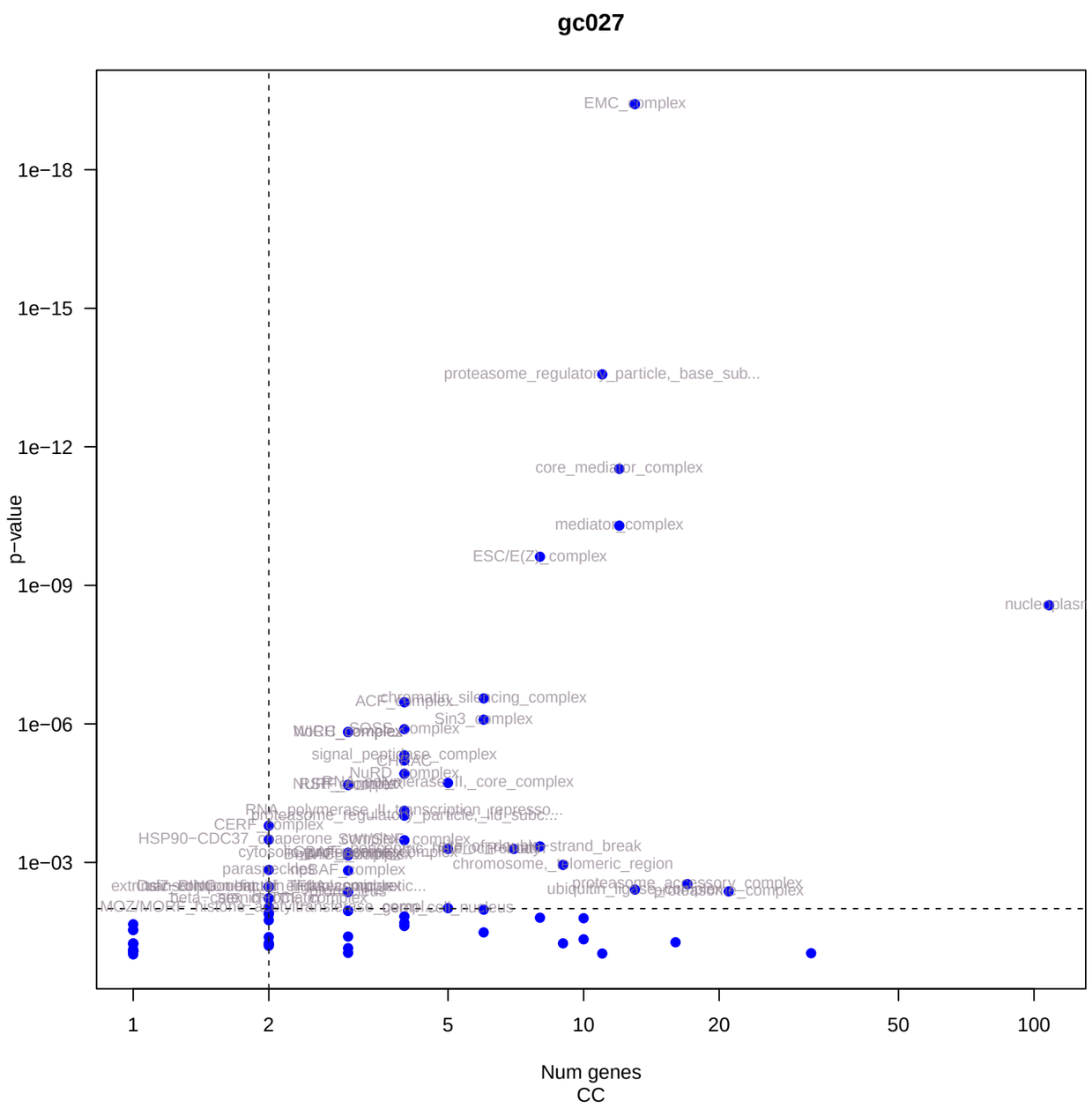
1e−12
1e−09
1e−06
1e−03
membrane_insertase_activity
proteasome−activating_activity
TBP−class_protein_binding
Krueppel−associated_box_domain_binding
RNA_polymerase_II_cis−regulatory_region_...
promoter−specific_chromatin_binding
transcription_coactivator_activity
transcription_coregulator_activity
ATP_hydrolysis_activity
DNA_binding
1
2
5
10
20
50
100
Num genes
MF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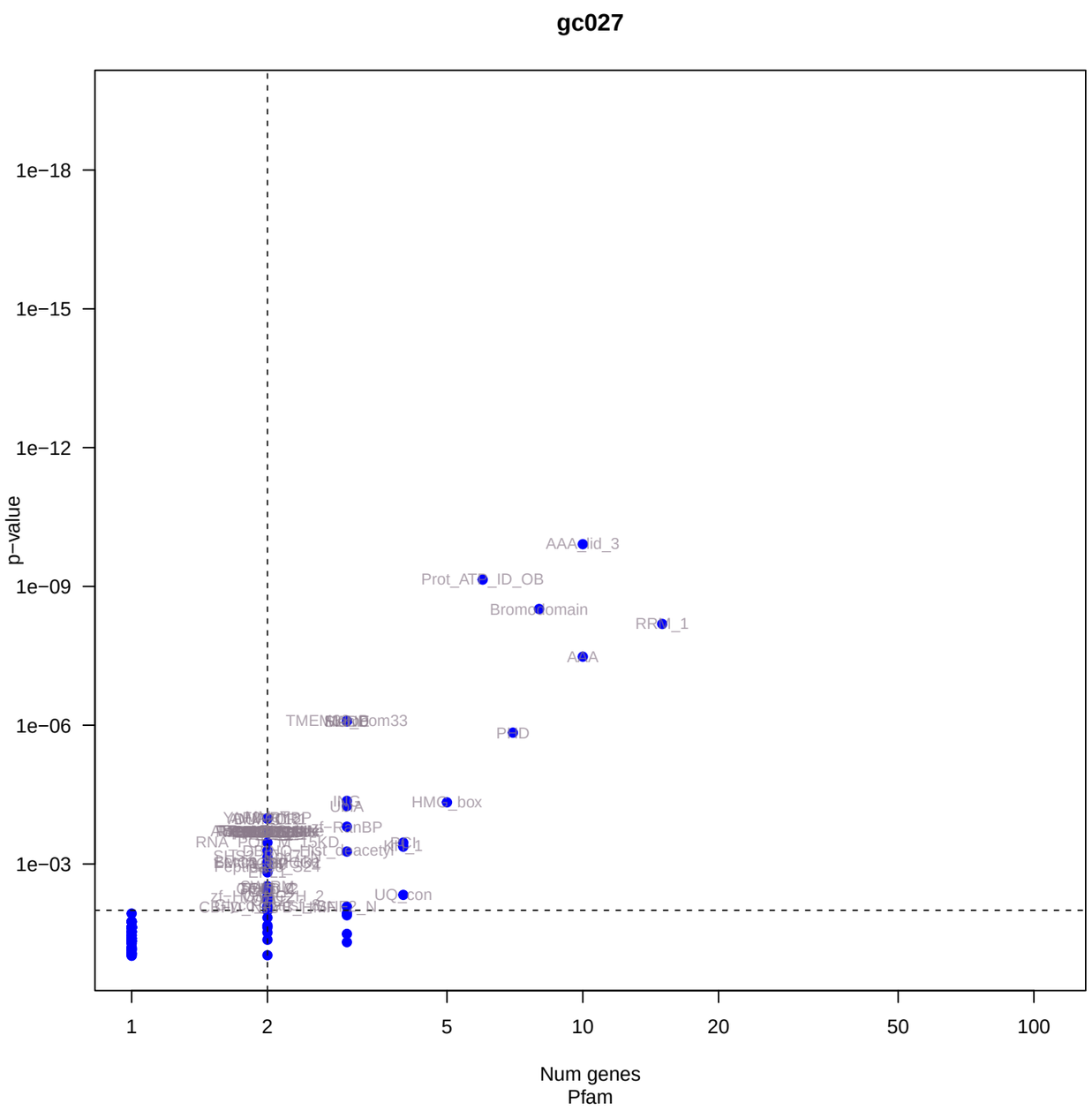
1e−15
1e−12
1e−09
1e−06
1e−03
EMC_complex
proteasome_regulatory_particle,_base_sub...
core_mediator_complex
mediator_complex
ESC/E(Z)_complex
chromatin_silencing_complex
Sin3_complex
signal_peptidase_complex
NuRD_complex
RNA_polymerase_II,_core_complex
CERF_complex
HSP90−CDC37_chaperone_complex
double−strand_break
chromosome,_telomeric_region
proteasome_accessory_complex
1
2
5
10
20
50
100
Num genes
CC


gc027
p−value
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1e−15
1e−12
1e−09
1e−06
1e−03
AAA_lid_3
Prot_ATP_ID_OB
Bromodomain
RRM_1
HMG_box
1
2
5
10
20
50
100
Num genes
Pfam