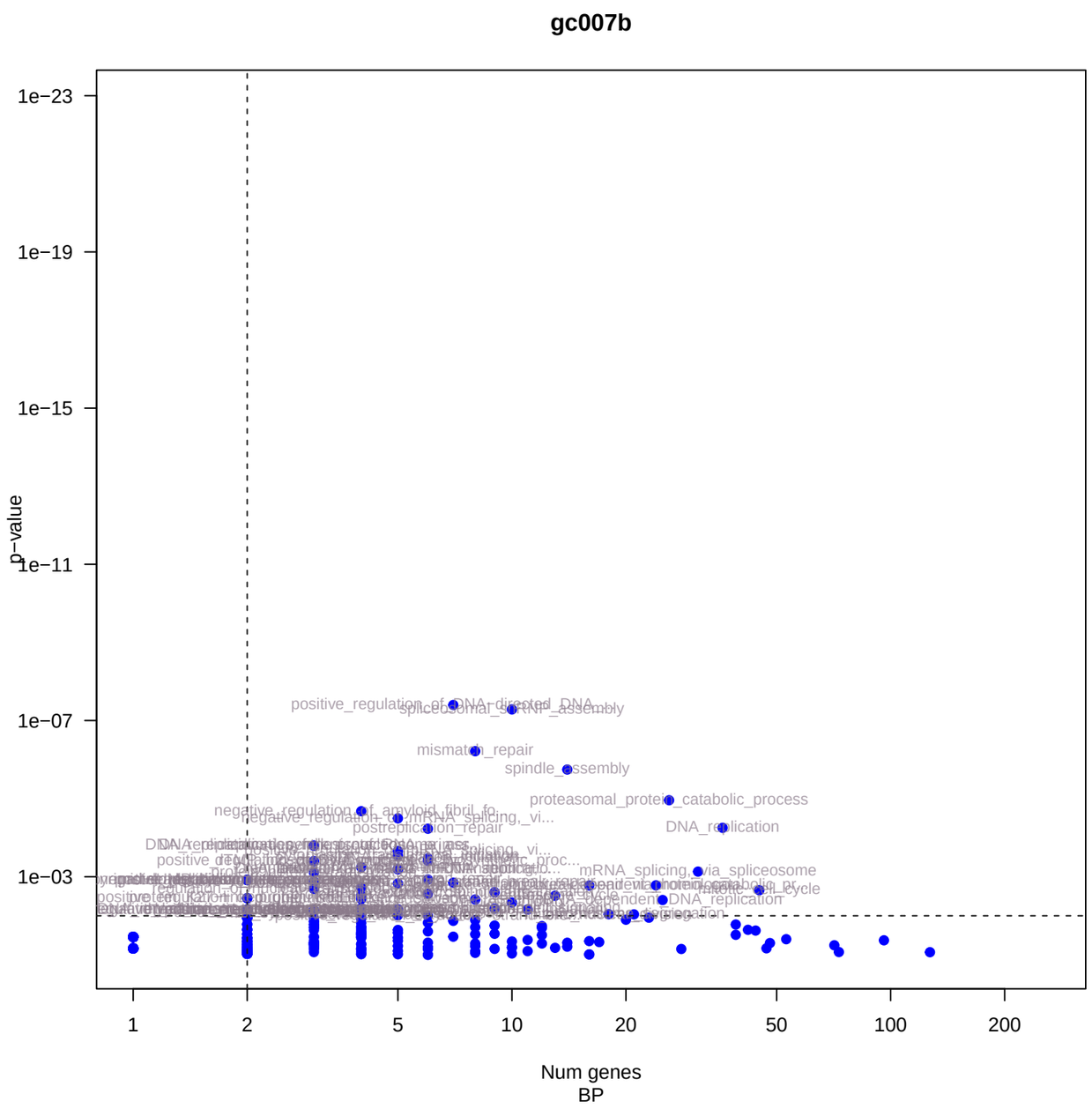
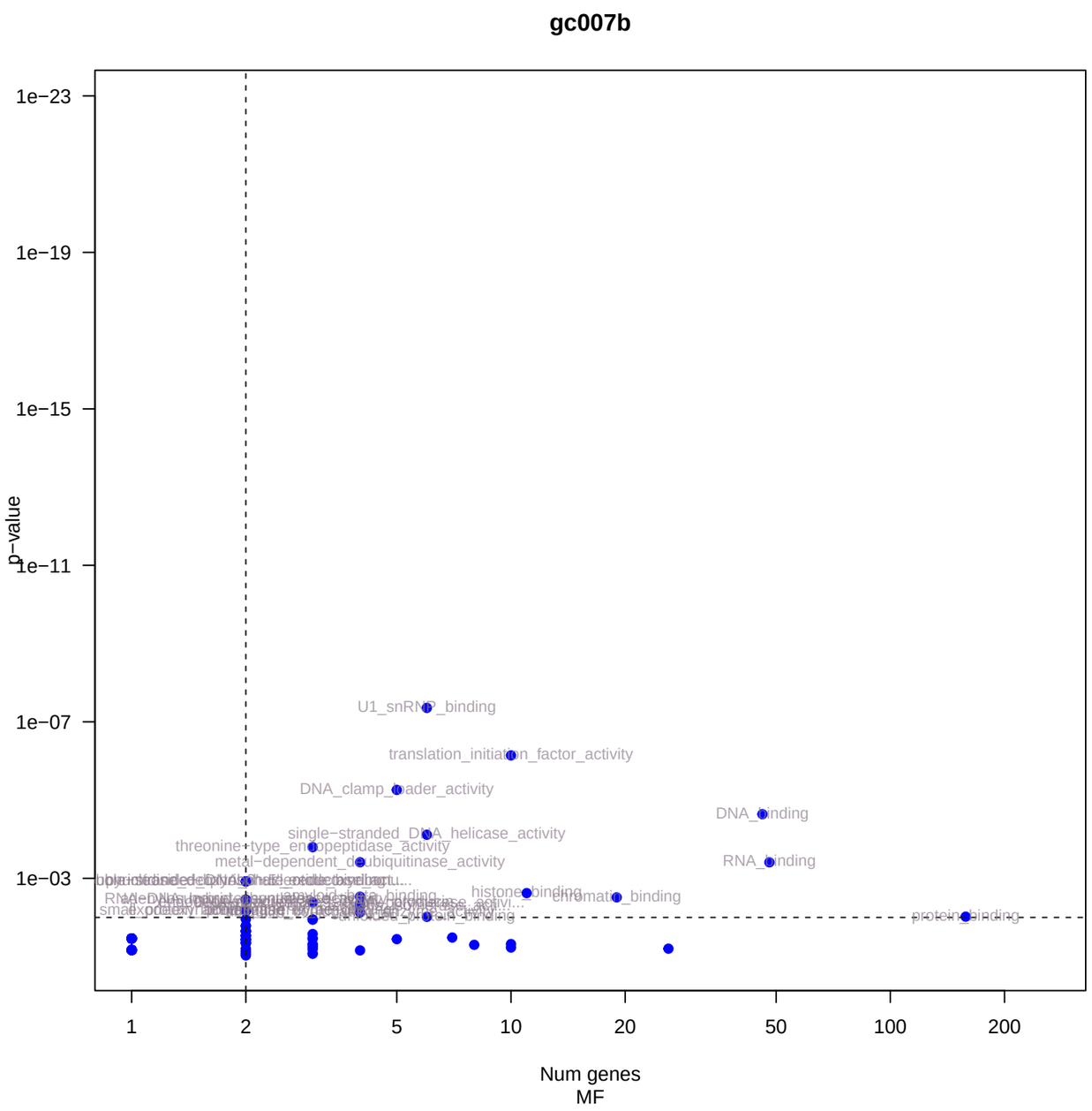
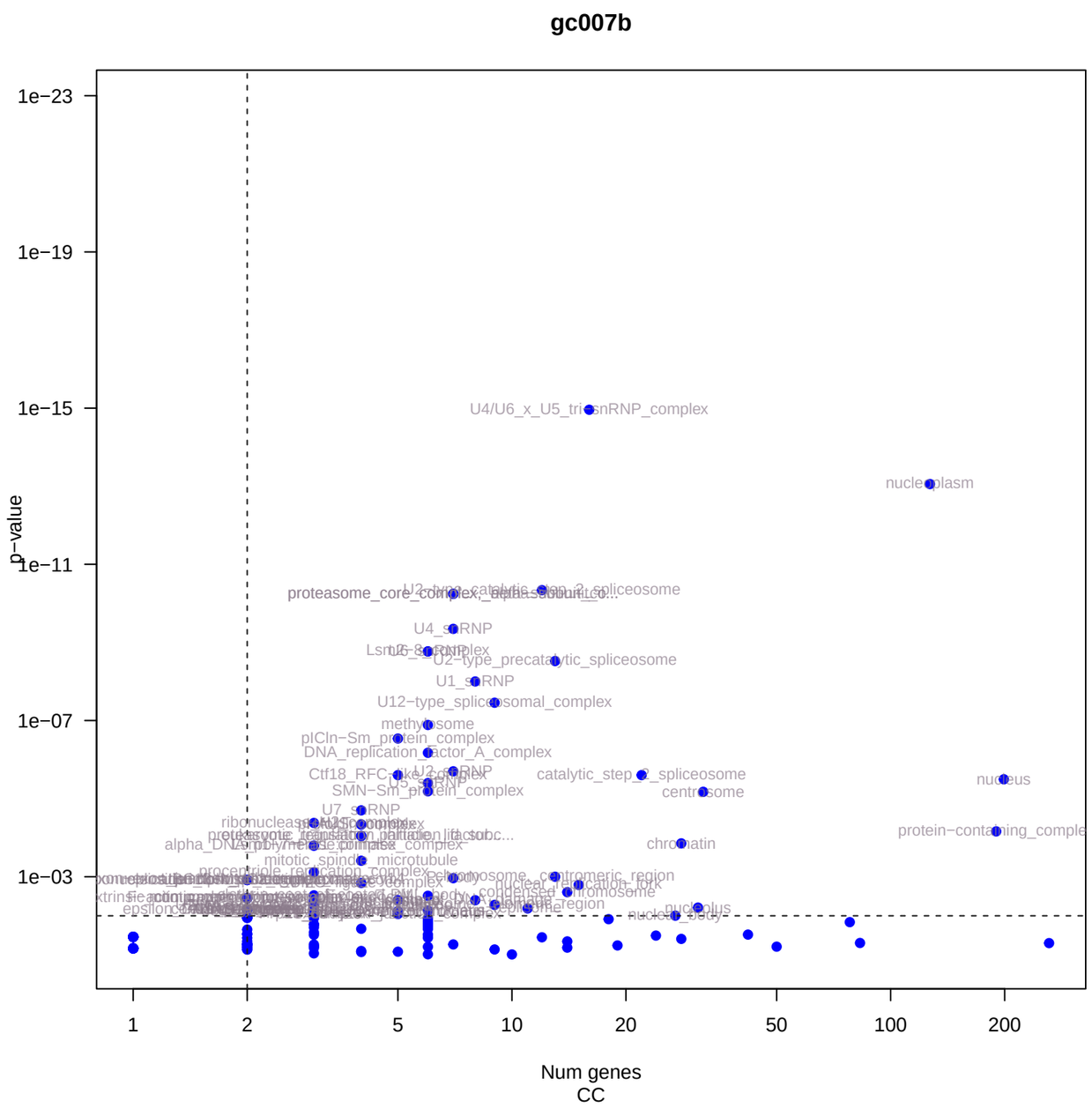
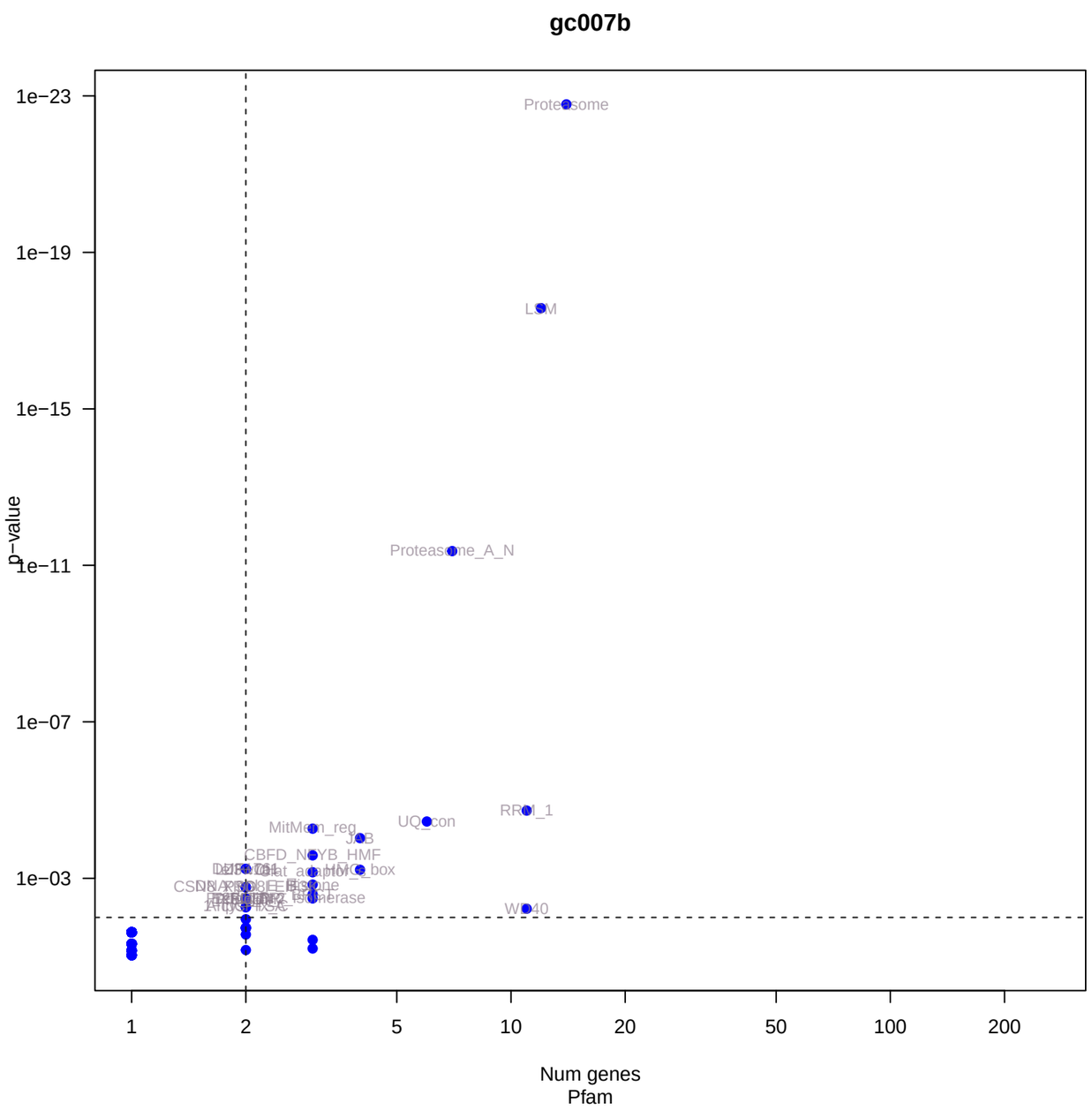

gc007b
p−value
1e−23
1e−19
1e−15
1e−11
1e−07
1e−03
1
2
5
10
20
50
100
200
Num genes
BP
positive_regulation_of_DNA_directed_DNA
spliceosomal_snRNP_assembly
mismatch_repair
spindle_assembly
proteasomal_protein_catabolic_process
negative_regulation_of_amyloid_fibril_fo
postreplication_repair
DNA_replication
mRNA_splicing,_via_spliceosome
DNA_replication


gc007b
p−value
1e−23
1e−19
1e−15
1e−11
1e−07
1e−03
1
2
5
10
20
50
100
200
Num genes
MF
U1_snRNP_binding
translation_initiation_factor_activity
DNA_clamp_loader_activity
DNA_binding
single−stranded_DNA_helicase_activity
threonine−type_endopeptidase_activity
metal−dependent_deubiquitinase_activity
RNA_binding
histone_binding
chromatin_binding
protein_binding


gc007b
p−value
1e−23
1e−19
1e−15
1e−11
1e−07
1e−03
1
2
5
10
20
50
100
200
Num genes
CC
U4/U6_x_U5_tri_snRNP_complex
nucleoplasm
proteasome_core_complex,_alpha_subunit_co...
U4_snRNP
U2−type_precatalytic_spliceosome
U1_snRNP
U12−type_spliceosomal_complex
methyosome
pICln−Sm_protein_complex
DNA_replication_factor_A_complex
catalytic_step_2_spliceosome
nucleus
centrosome
SMN−Sm_protein_complex
U7_snRNP
protein−containing_comple
chromatin
mitotic_spindle_microtubule
centromeric_region
condensed_chromosome
nucleolus


gc007b
p−value
1e−23
1e−19
1e−15
1e−11
1e−07
1e−03
1
2
5
10
20
50
100
200
Num genes
Pfam
Proteasome
LOM
Proteasome_A_N
UQ_con
RRM_1
MitMem_reg
JAB
CBFD_NFYB_HMF
W40