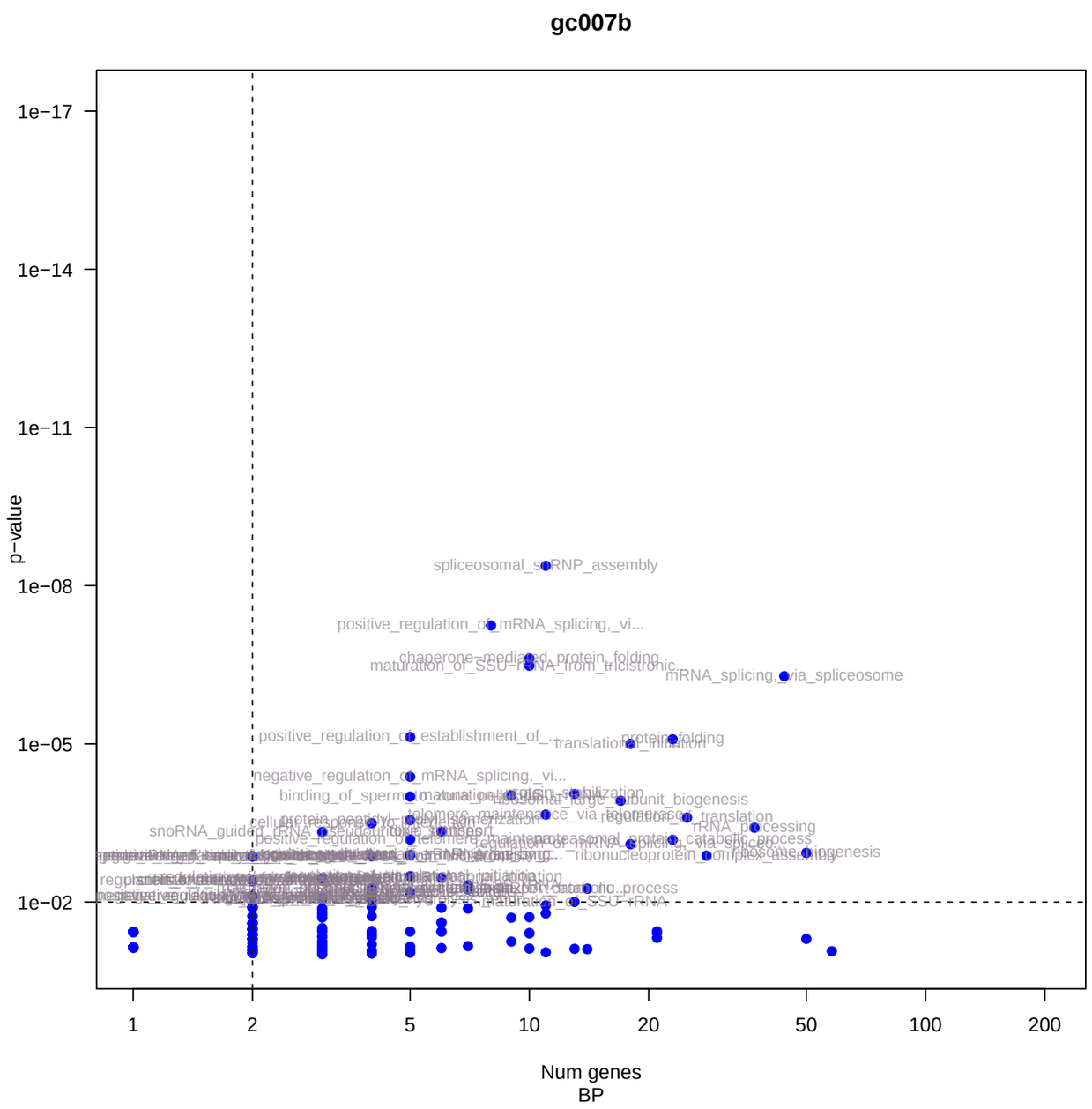
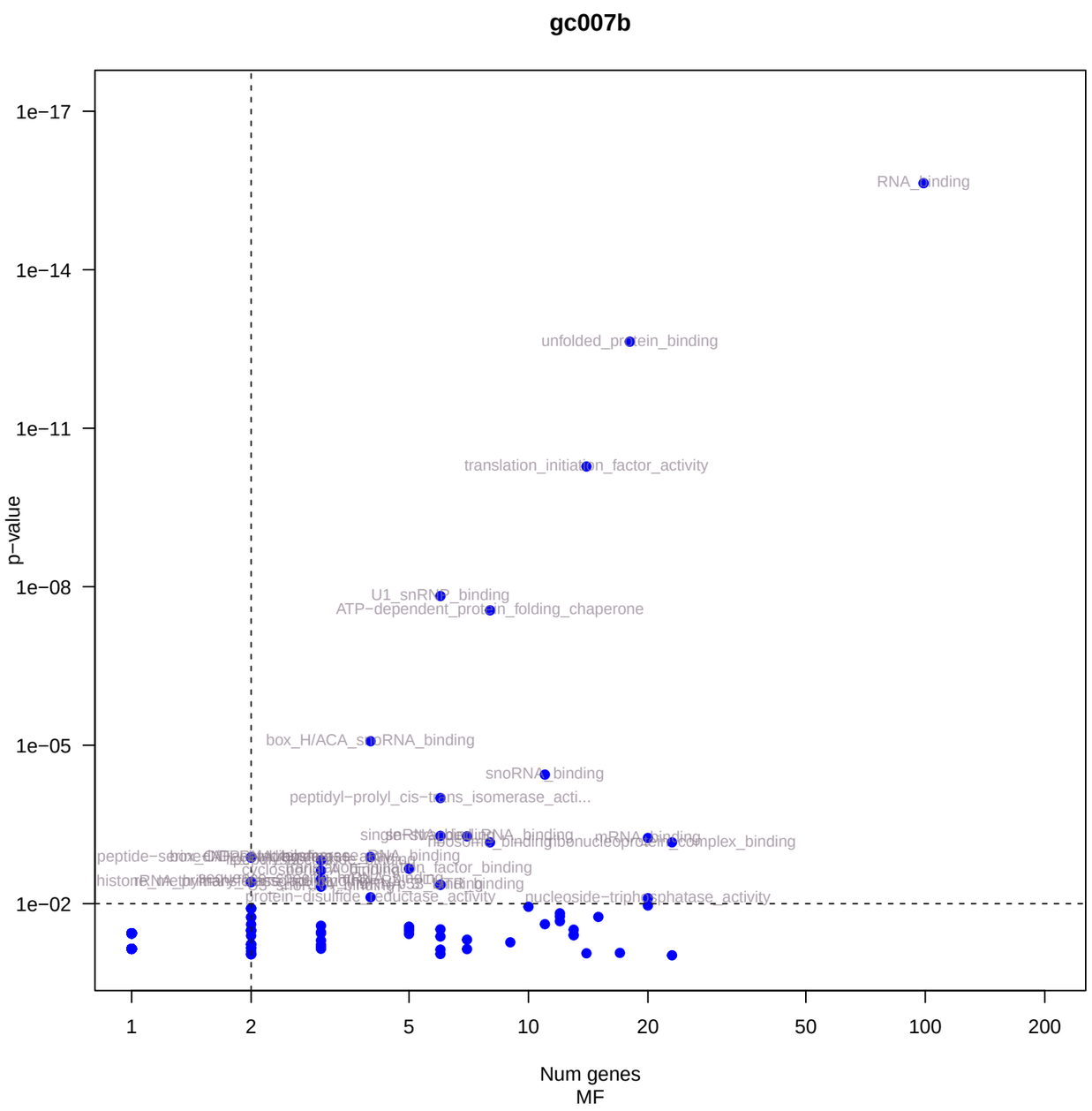
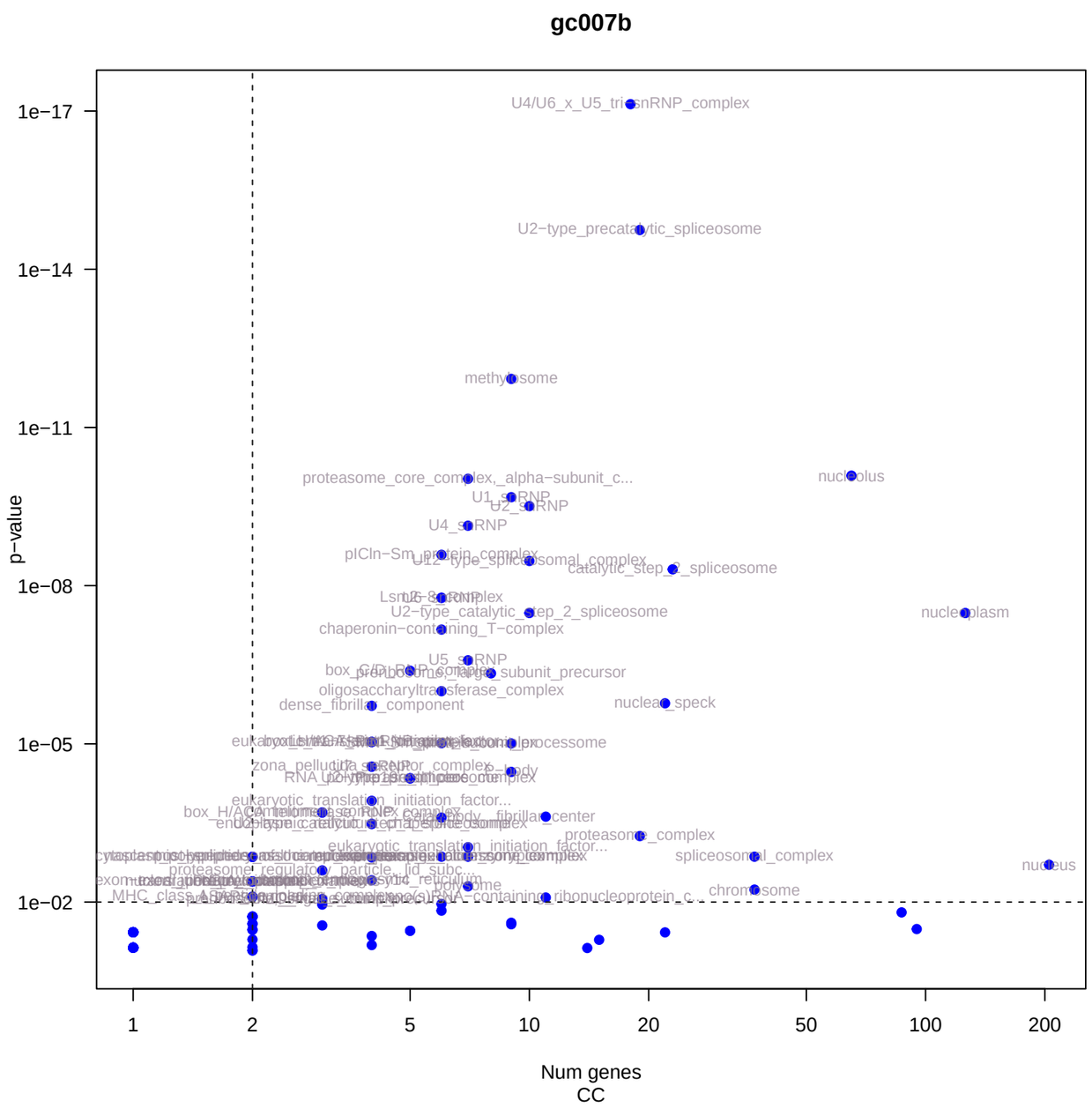
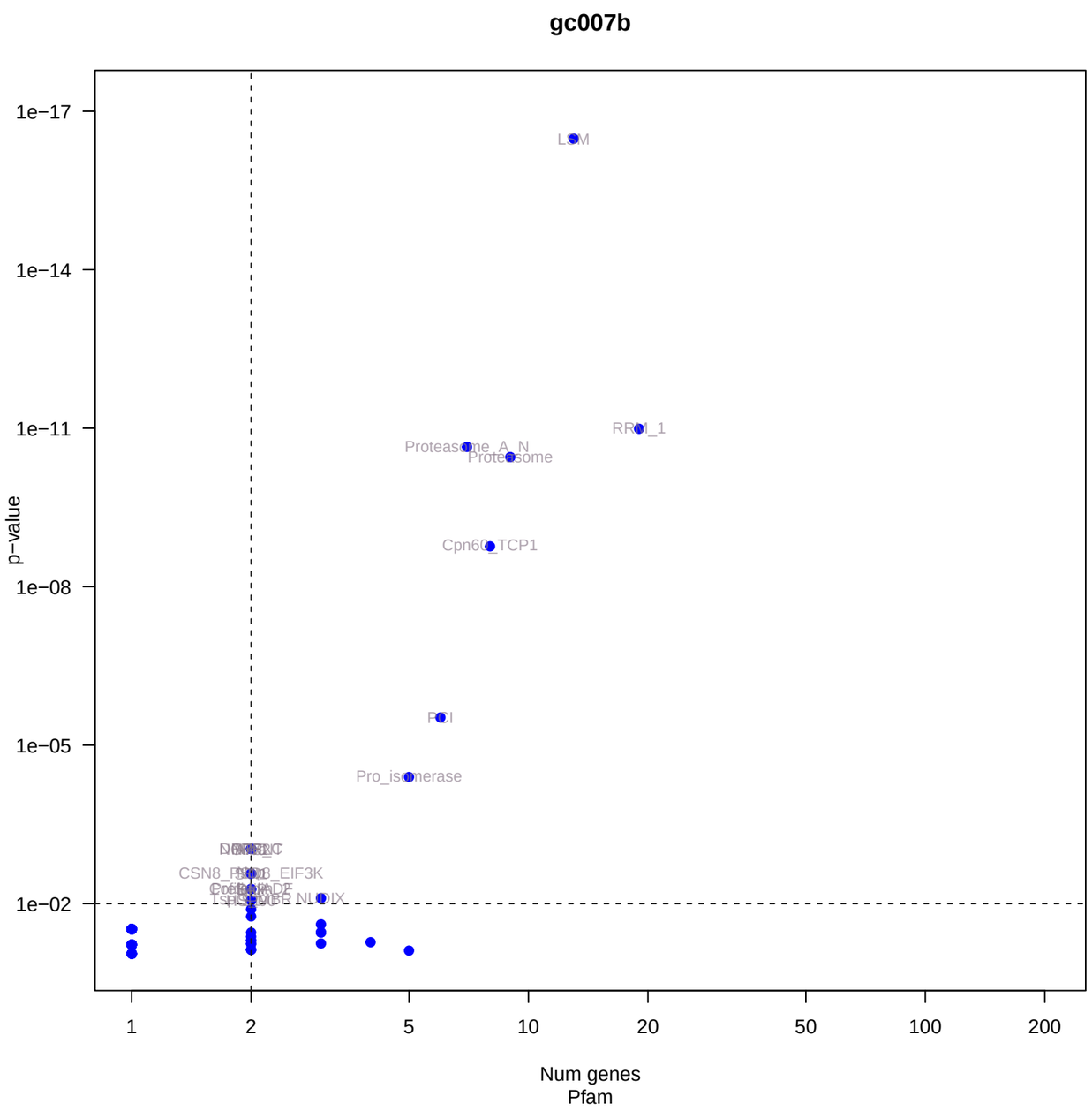

gc007b
p−value
1e−17
1e−14
1e−11
1e−08
1e−05
1e−02
spliceosomal_snRNP_assembly
positive_regulation_of_mRNA_splicing,_vi...
mRNA_splicing,_via_spliceosome
protein_folding
translation
negative_regulation_of_mRNA_splicing,_vi...
snoRNA_guided_rRNA_pseudouridine_synthesis
ribonucleoprotein_complex_assembly
1
2
5
10
20
50
100
200
Num genes
BP


gc007b
p−value
1e−17
1e−14
1e−11
1e−08
1e−05
1e−02
RNA_binding
unfolded_protein_binding
translation_initiation_factor_activity
U1_snRNP_binding
ATP−dependent_protein_folding_chaperone
box_H/ACA_snoRNA_binding
snoRNA_binding
peptidyl−prolyl_cis−trans_isomerase_acti...
mRNA_binding
ribonucleoprotein_complex_binding
protein_disulfide_oxidoreductase_activity
nucleoside−triphosphatase_activity
1
2
5
10
20
50
100
200
Num genes
MF


gc007b
p−value
1e−17
1e−14
1e−11
1e−08
1e−05
1e−02
U4/U6_x_U5_tri_snRNP_complex
U2−type_precatalytic_spliceosome
methyosome
proteasome_core_complex,_alpha−subunit_c...
nucleolus
U1_snRNP
U2_snRNP
U4_snRNP
U2−type_catalytic_step_2_spliceosome
nucleoplasm
chaperonin−containing_T−complex
U5_snRNP
oligosaccharyltransferase_complex
dense_fibrillar_component
nucleus_speck
eukaryotic_translation_initiation_factor...
proteasome_complex
spliceosomal_complex
nucleus
chromosome
1
2
5
10
20
50
100
200
Num genes
CC


gc007b
p−value
1e−17
1e−14
1e−11
1e−08
1e−05
1e−02
LIM
RRM_1
Proteasome_A_N
Proteasome
Cpn60_TCP1
PCI
Pro_isomerase
1
2
5
10
20
50
100
200
Num genes
Pfam