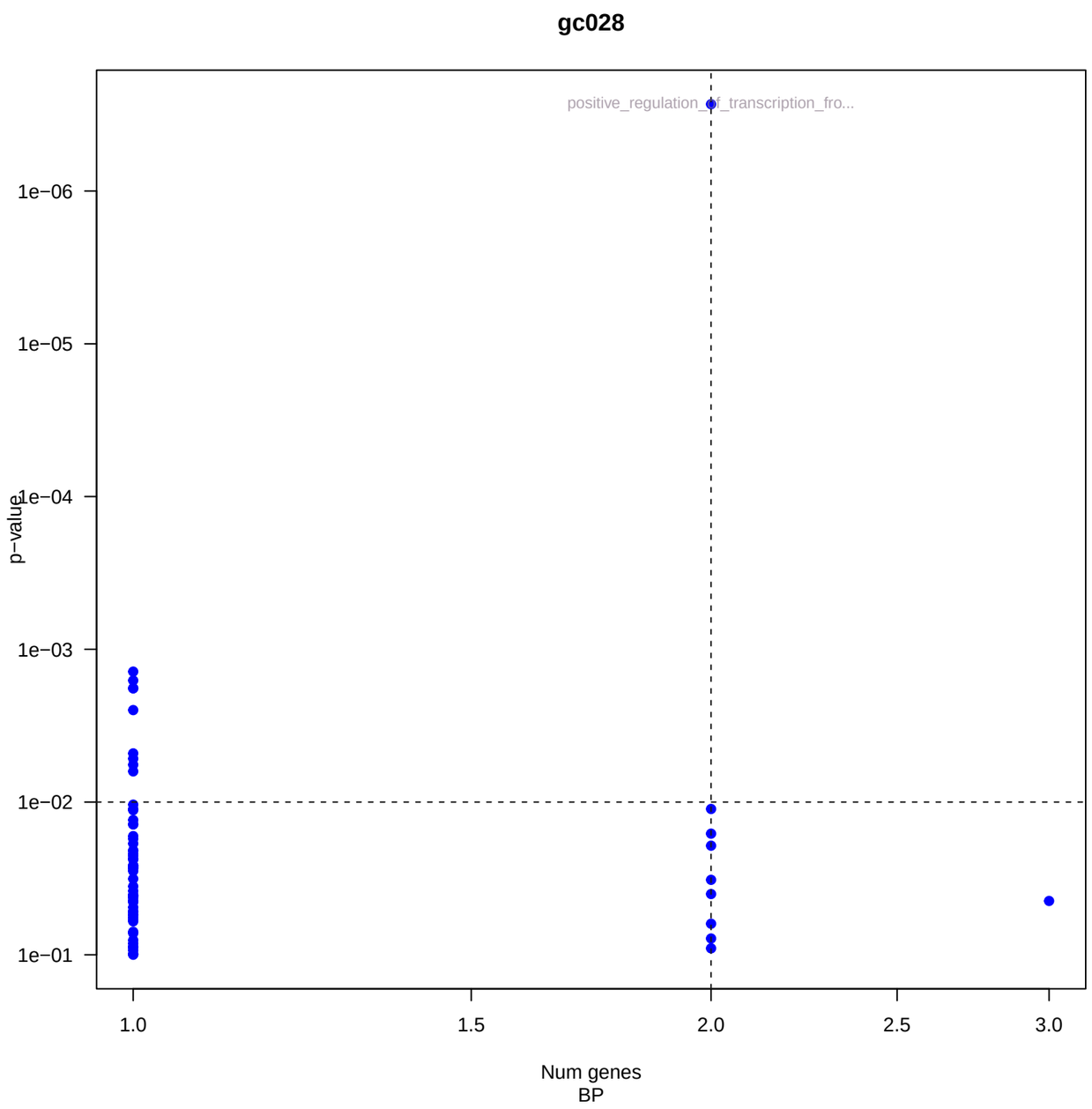
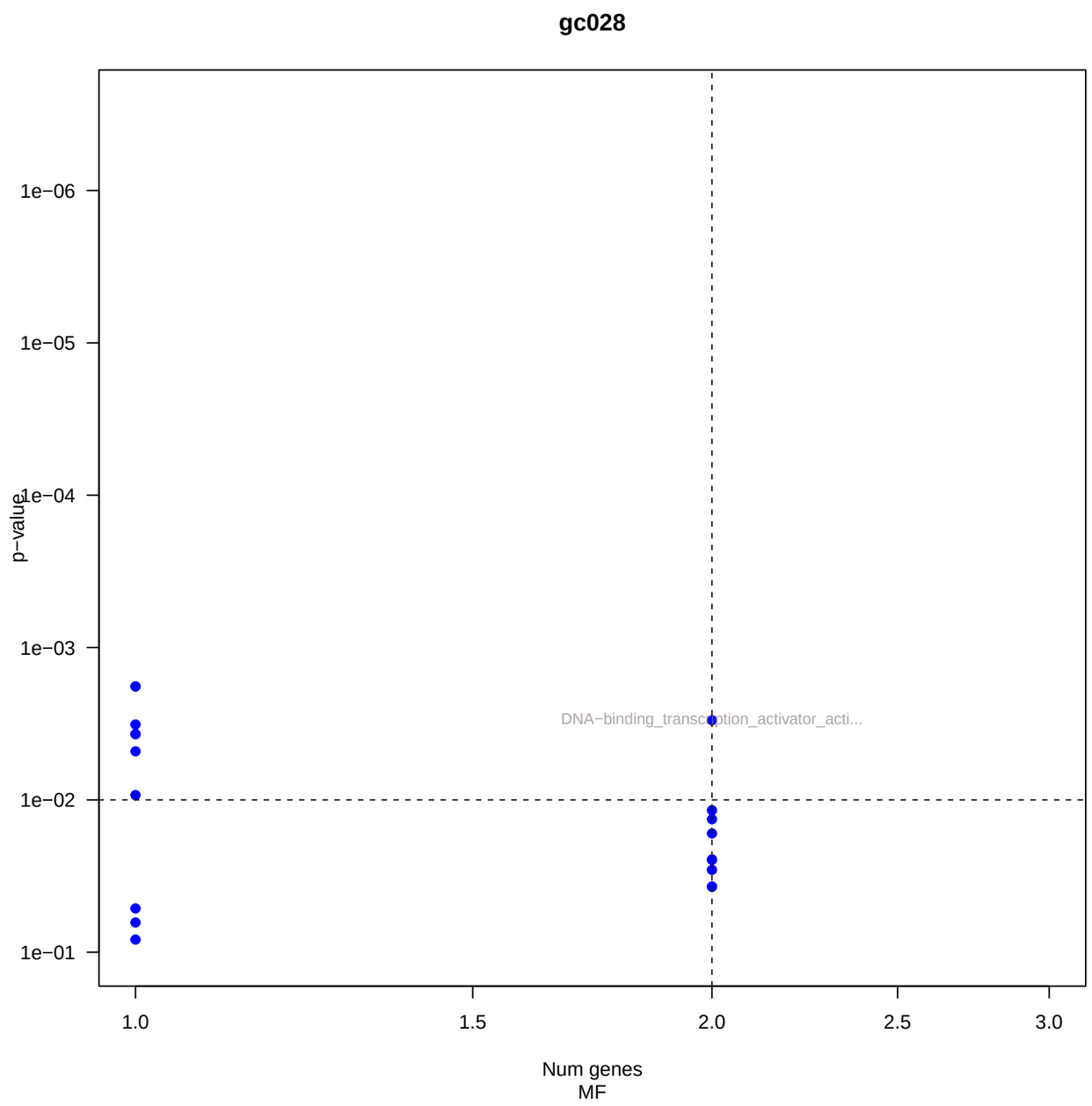
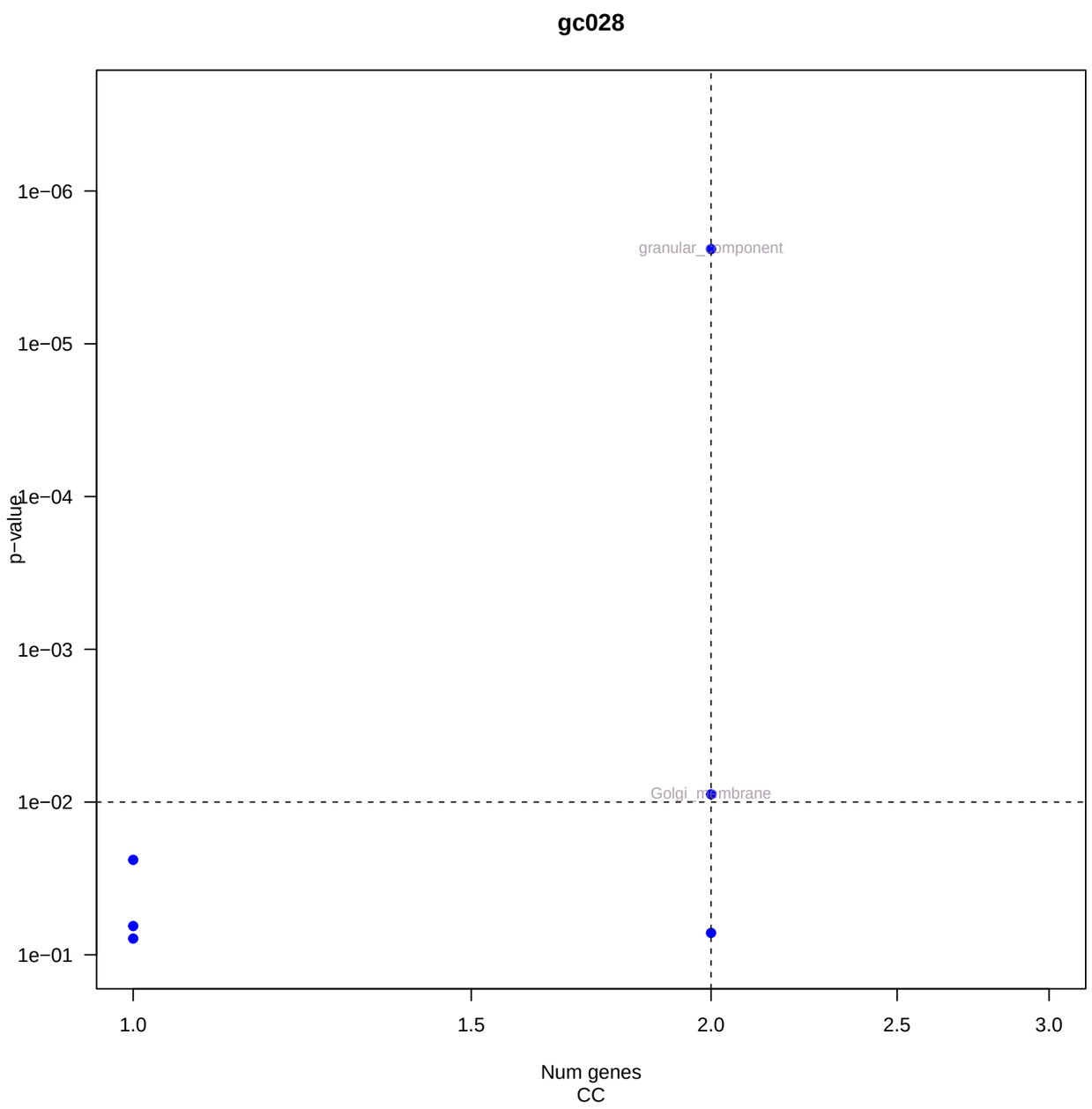

gc028
positive_regulation_of_transcription_fro...
p−value
1e−06
1e−05
1e−04
1e−03
1e−02
1e−01
1.0
1.5
2.0
2.5
3.0
Num genes
BP


gc028
DNA−binding_transcription_activator_acti...
p−value
1e−06
1e−05
1e−04
1e−03
1e−02
1e−01
1.0
1.5
2.0
2.5
3.0
Num genes
MF


gc028
granular_component
Golgi_membrane
p−value
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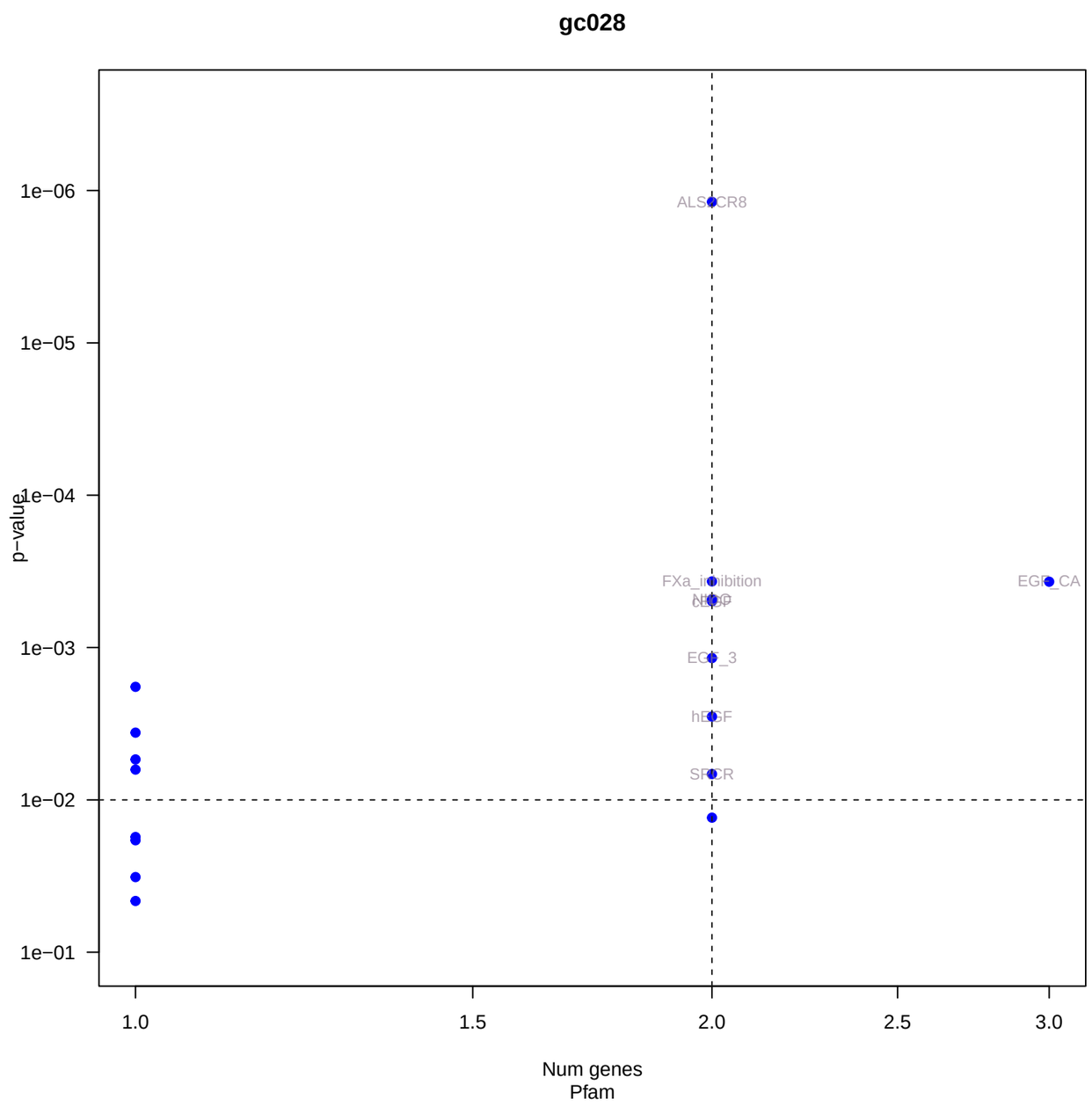
1e−05
1e−04
1e−03
1e−02
1e−01
1.0
1.5
2.0
2.5
3.0
Num genes
CC


gc028
ALSCR8
FXa_inhibition
EGF_CA
EGF_3
hEGF
SPCR
p−value
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1e−05
1e−04
1e−03
1e−02
1e−01
1.0
1.5
2.0
2.5
3.0
Num genes
Pfam