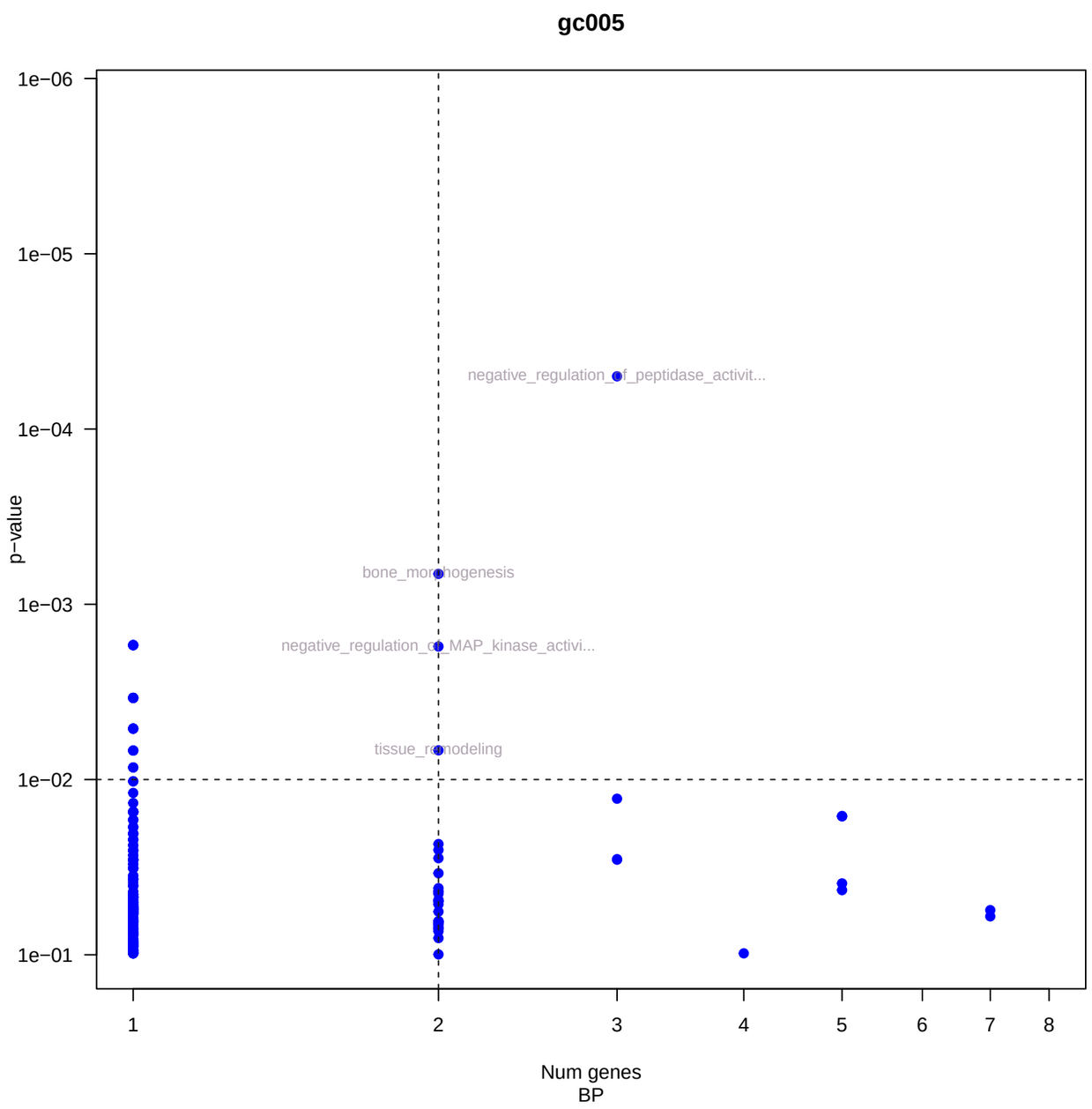
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1e−04
1e−03
1e−02
1e−01
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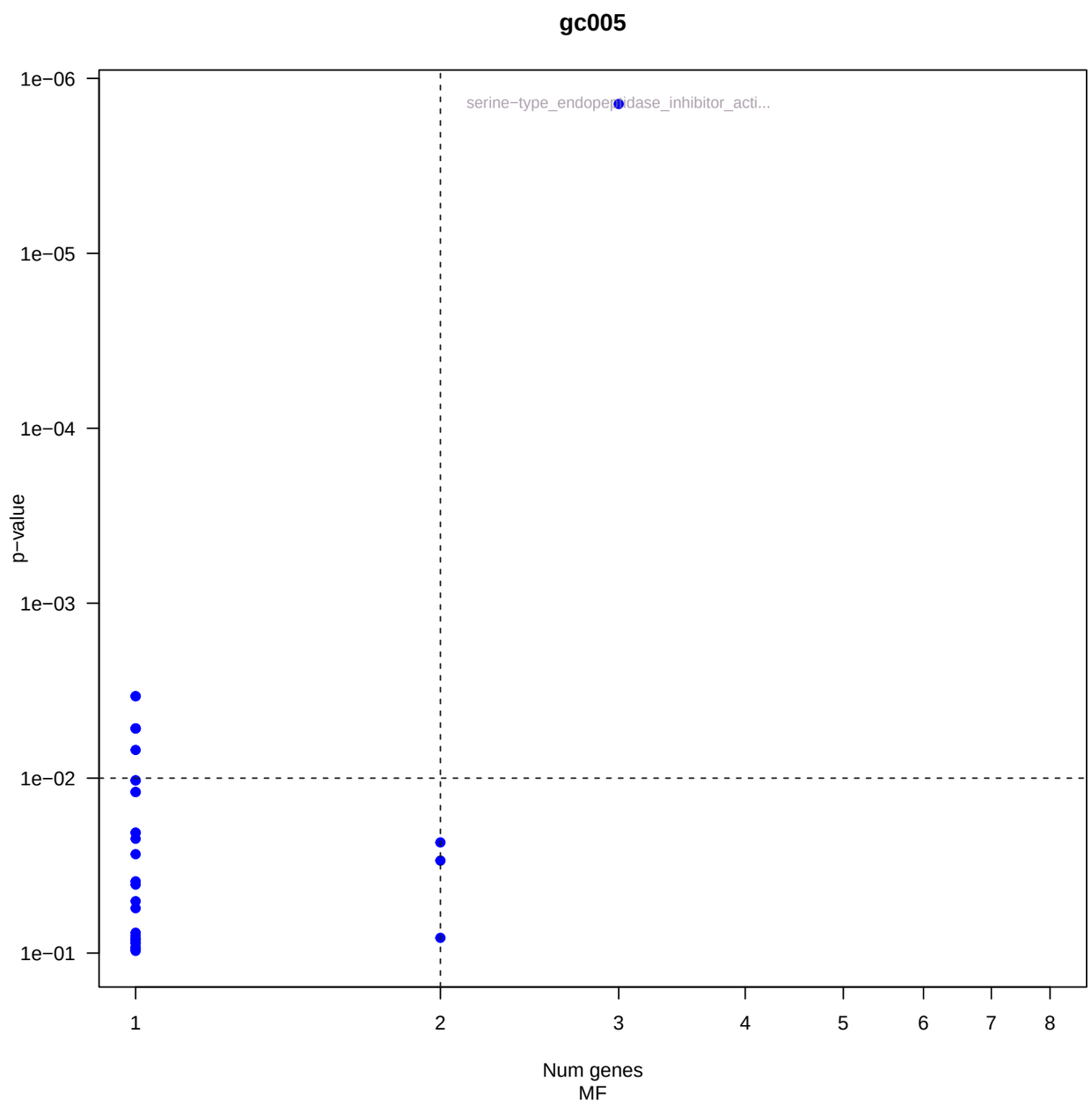
bone_morphogenesis
negative_regulation_of_MAP_kinase_activi...
tissue_remodeling
1
2
3
4
5
6
7
8
Num genes
BP


gc005
p−value
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1e−04
1e−03
1e−02
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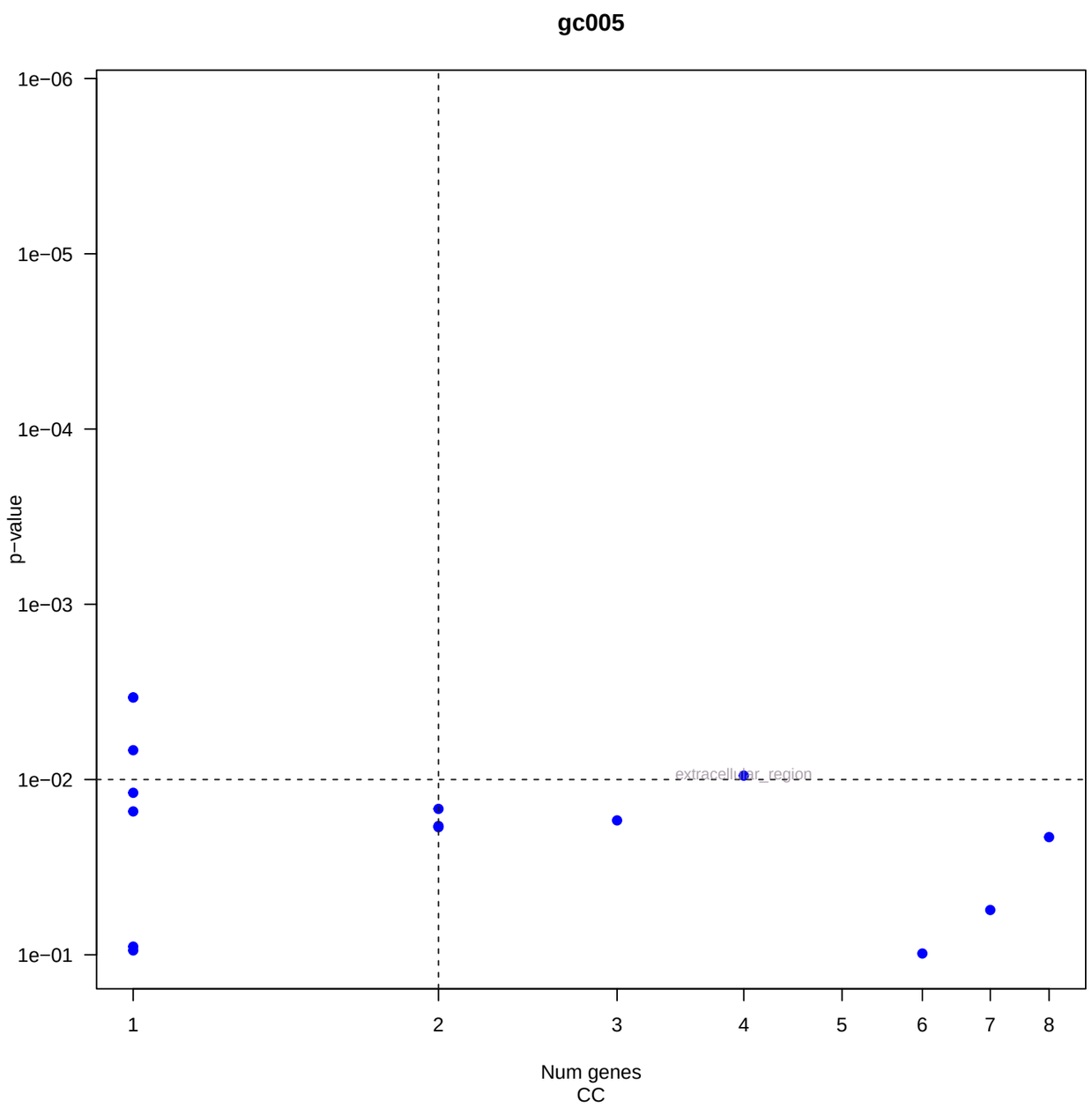
1
2
3
4
5
6
7
8
Num genes
MF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1e−01
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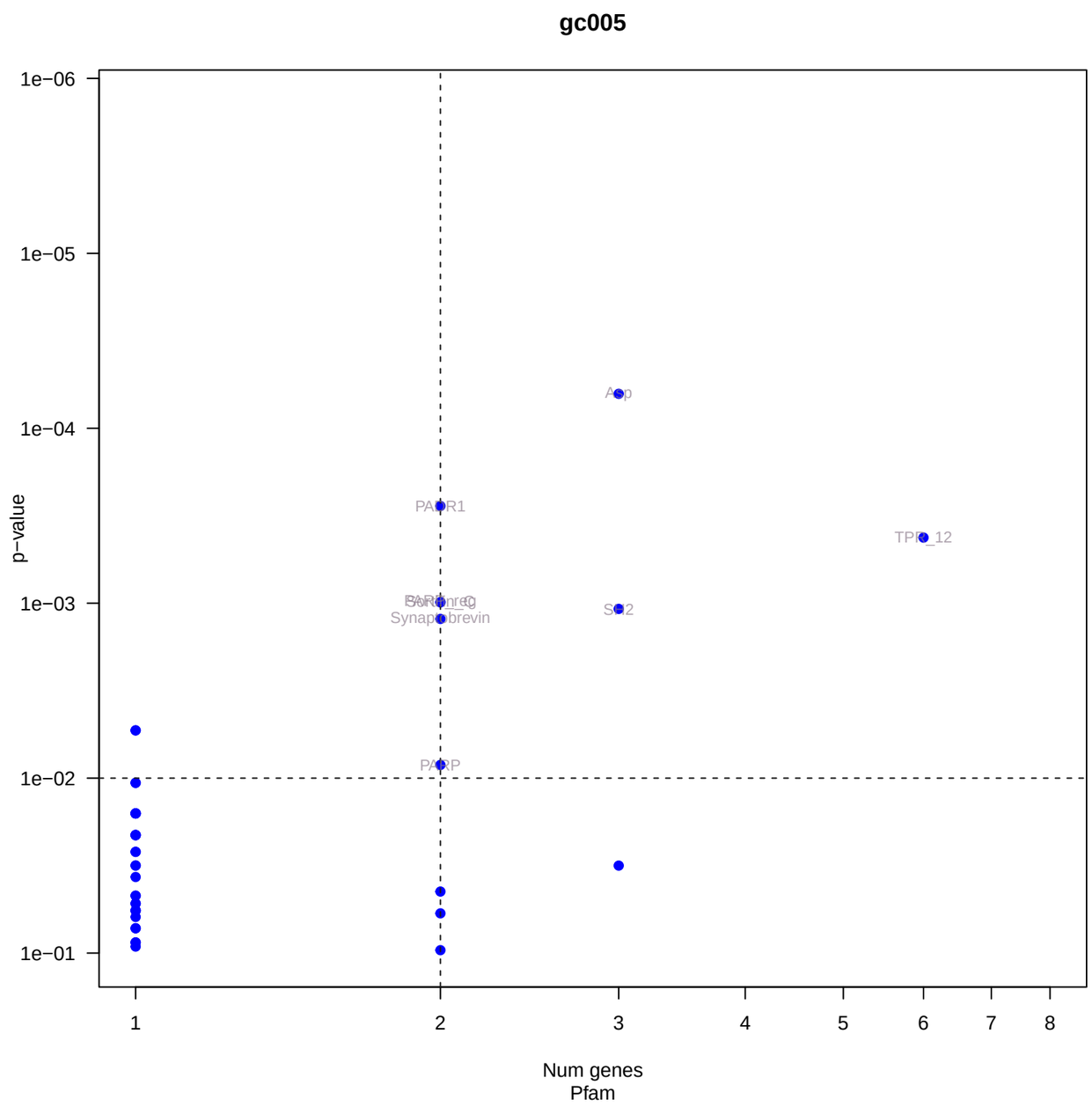
2
3
4
5
6
7
8
Num genes
CC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p−value
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1e−02
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App
PARP1
TPR_12
Synaptobrevin
SNF2
PARP
1
2
3
4
5
6
7
8
Num genes
Pfam