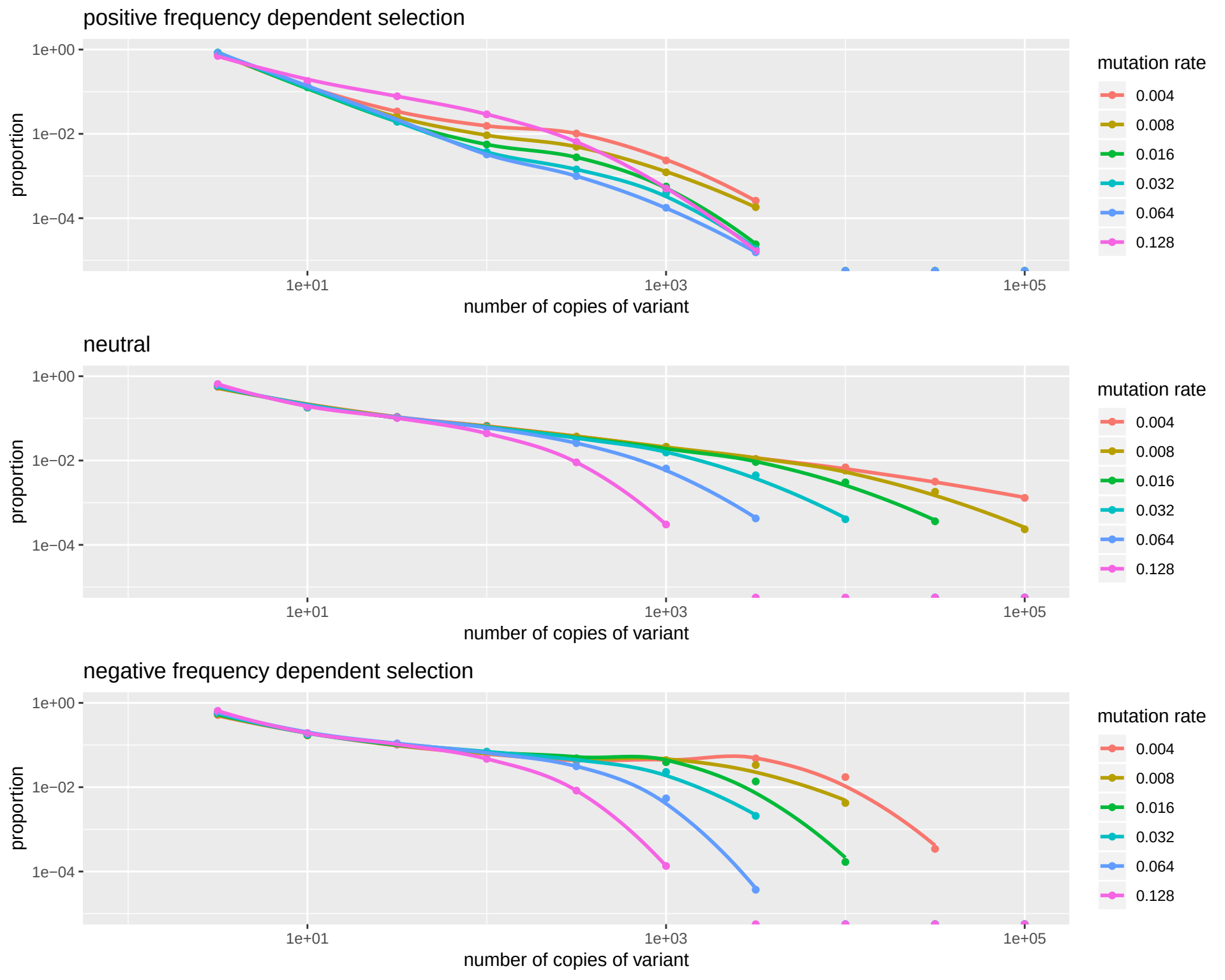
positive frequency dependent selection
proportion
1e+00
1e−02
1e−04
1e+01
1e+03
1e+05
number of copies of variant
mutation rate
0.004
0.008
0.016
0.032
0.064
0.128
neutral
proportion
1e+00
1e−02
1e−04
1e+01
1e+03
1e+05
number of copies of variant
mutation rate
0.004
0.008
0.016
0.032
0.064
0.128
negative frequency dependent selection
proportion
1e+00
1e−02
1e−04
1e+01
1e+03
1e+05
number of copies of variant
mutation rate
0.004
0.008
0.016
0.032
0.064
0.128