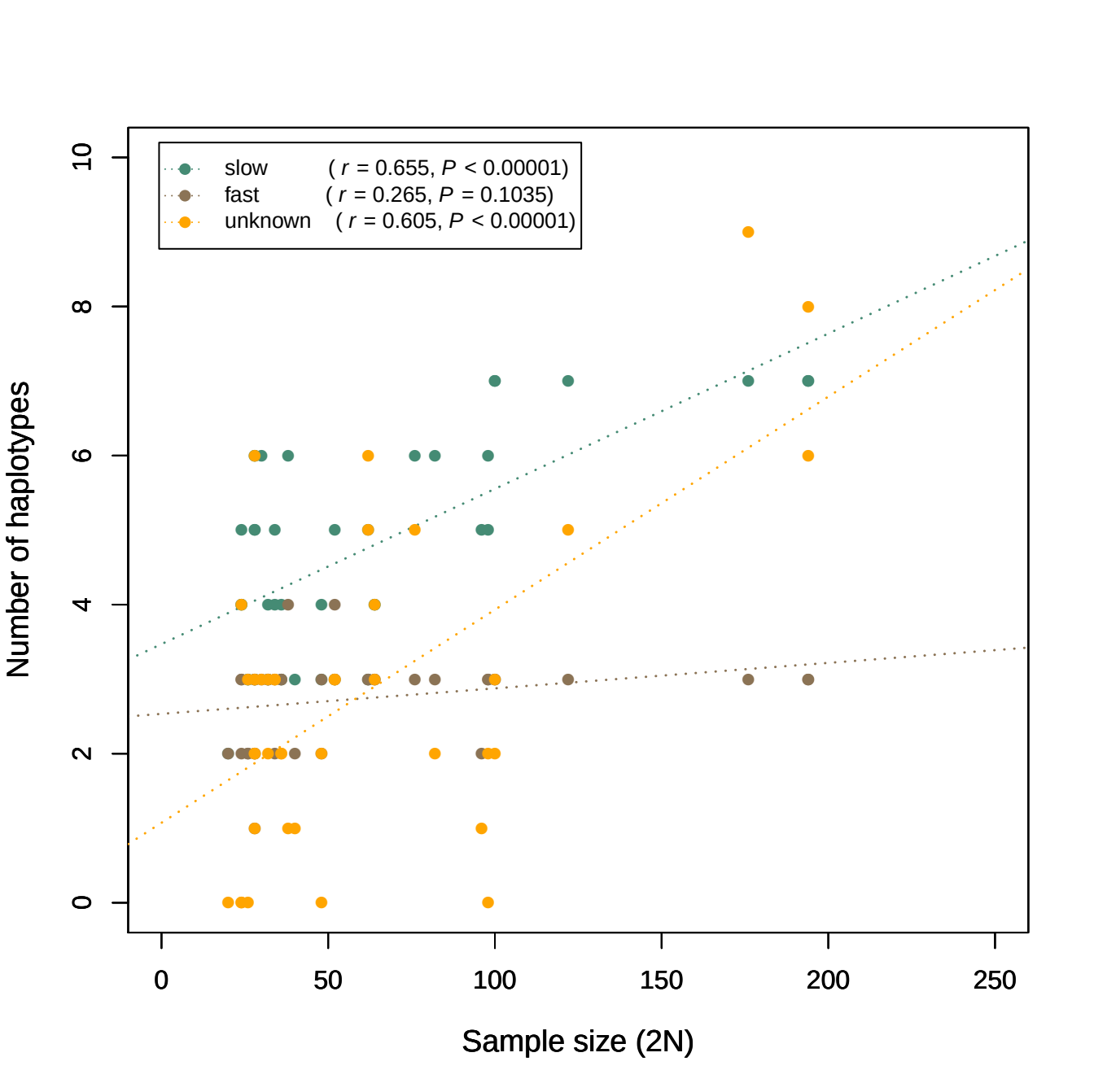

slow (r = 0.655, P < 0.00001)
fast (r = 0.265, P = 0.1035)
unknown (r = 0.605, P < 0.00001)
Number of haplotypes
0
2
4
6
8
10
0
50
100
150
200
250
Sample size (2N)