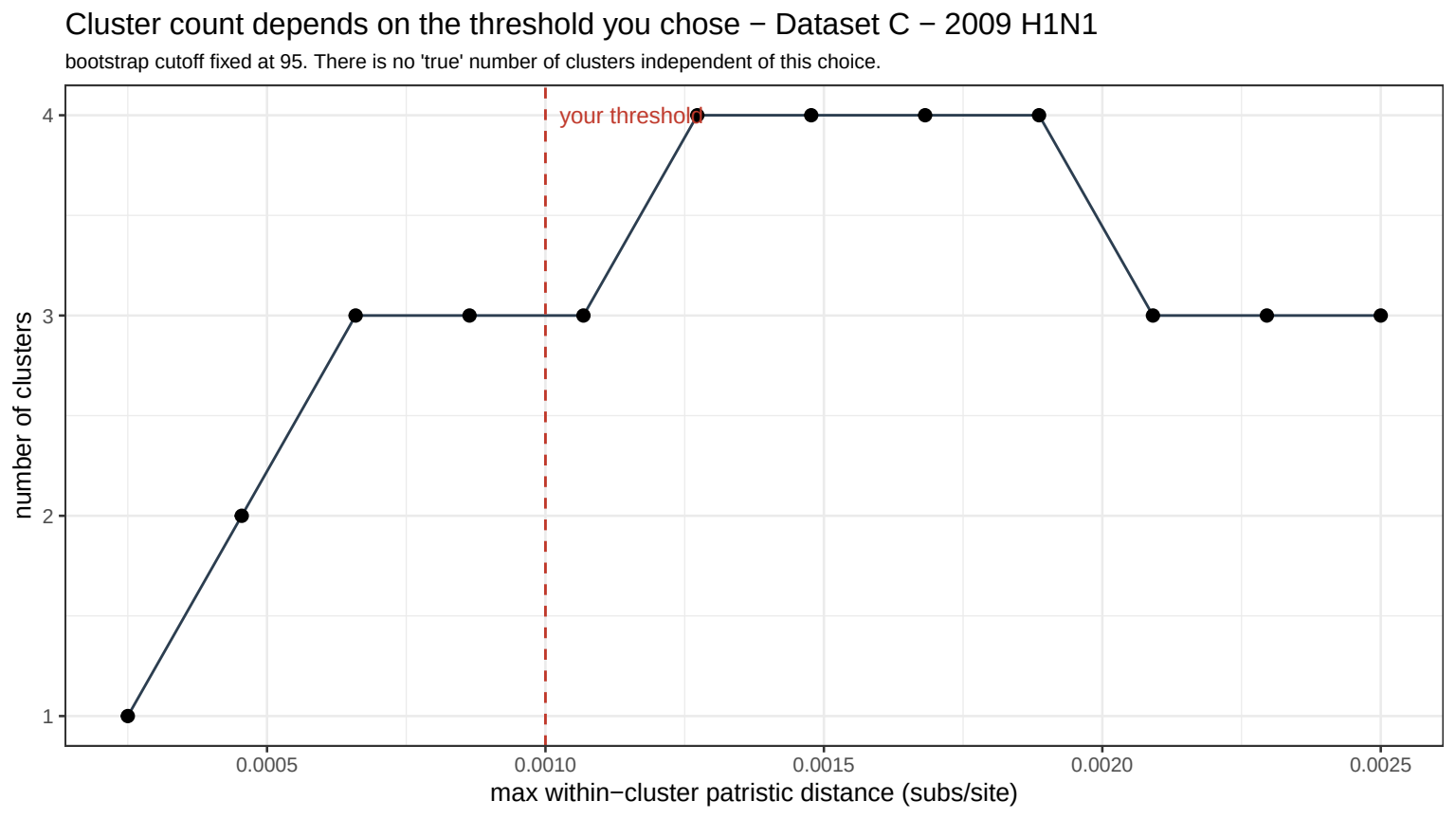

Cluster count depends on the threshold you chose − Dataset C − 2009 H1N1
bootstrap cutoff fixed at 95. There is no 'true' number of clusters independent of this choice.
your threshold
number of clusters
1
2
3
4
0.0005
0.0010
0.0015
0.0020
0.0025
max within−cluster patristic distance (subs/site)