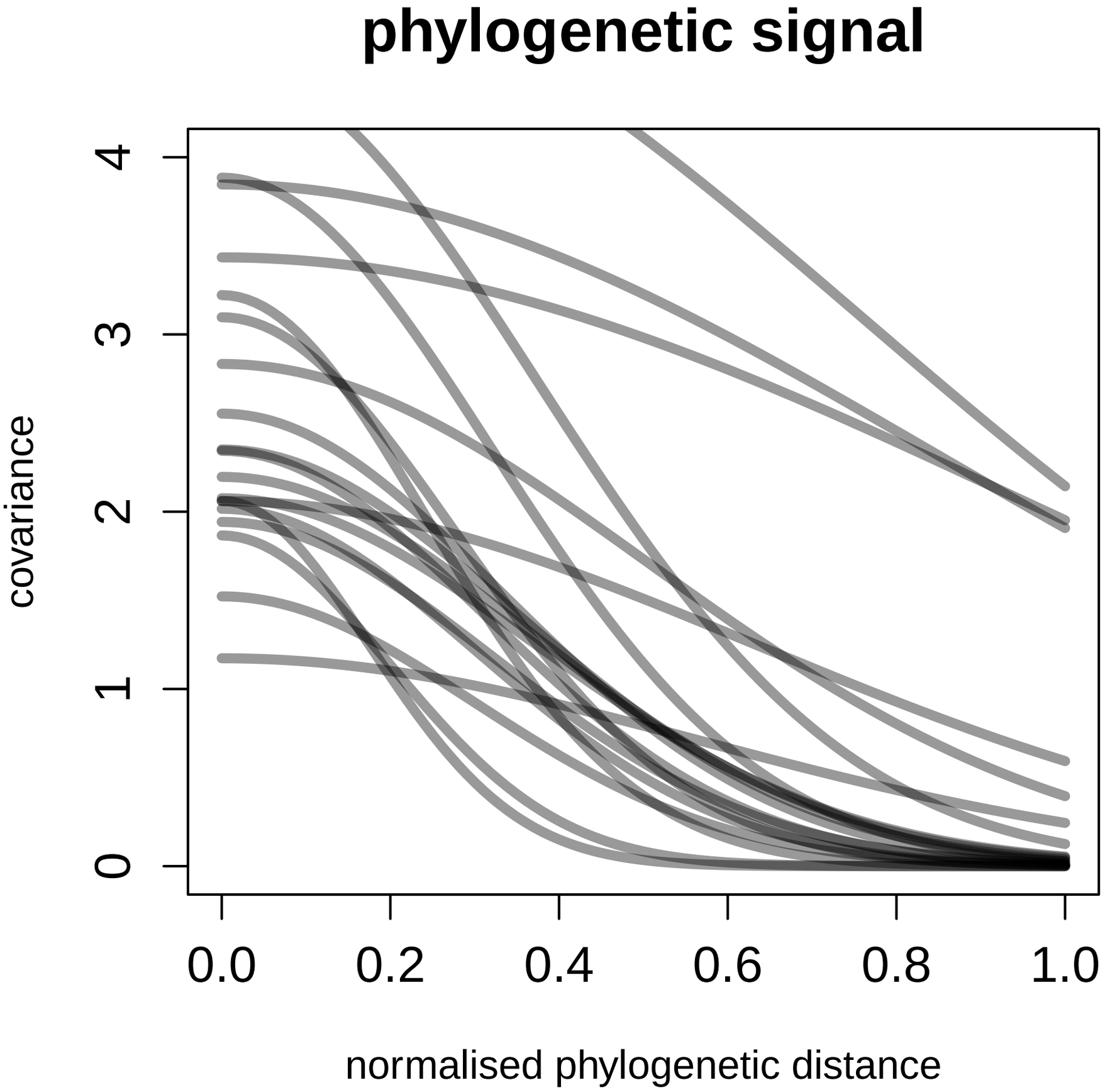
phylogenetic signal
covariance
0
1
2
3
4
0.0
0.2
0.4
0.6
0.8
1.0
normalised phylogenetic distance