Reordered dendrogram from
hclust(d = spe.ch, method = "ward.D2")

Height

3.0
1.5
0.0

Cluster 1
Cluster 2
Cluster 3
Cluster 4

30 29 28 27 26 22 21 20 25 24 23 9 5 19 18 17 16 15 10 6 4 7 3 2 12 11 13 14 8 1

30 sites
4 clusters