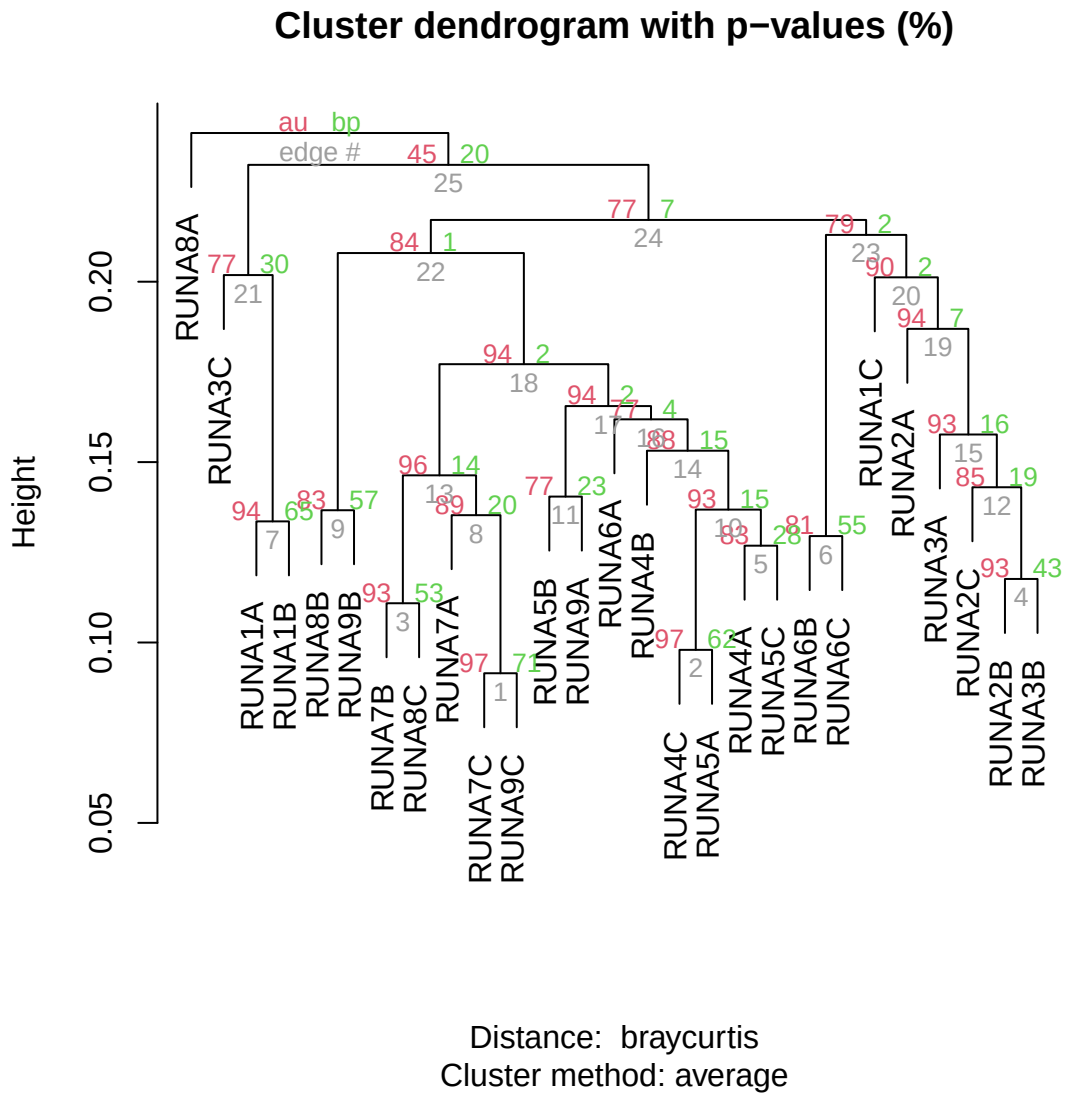
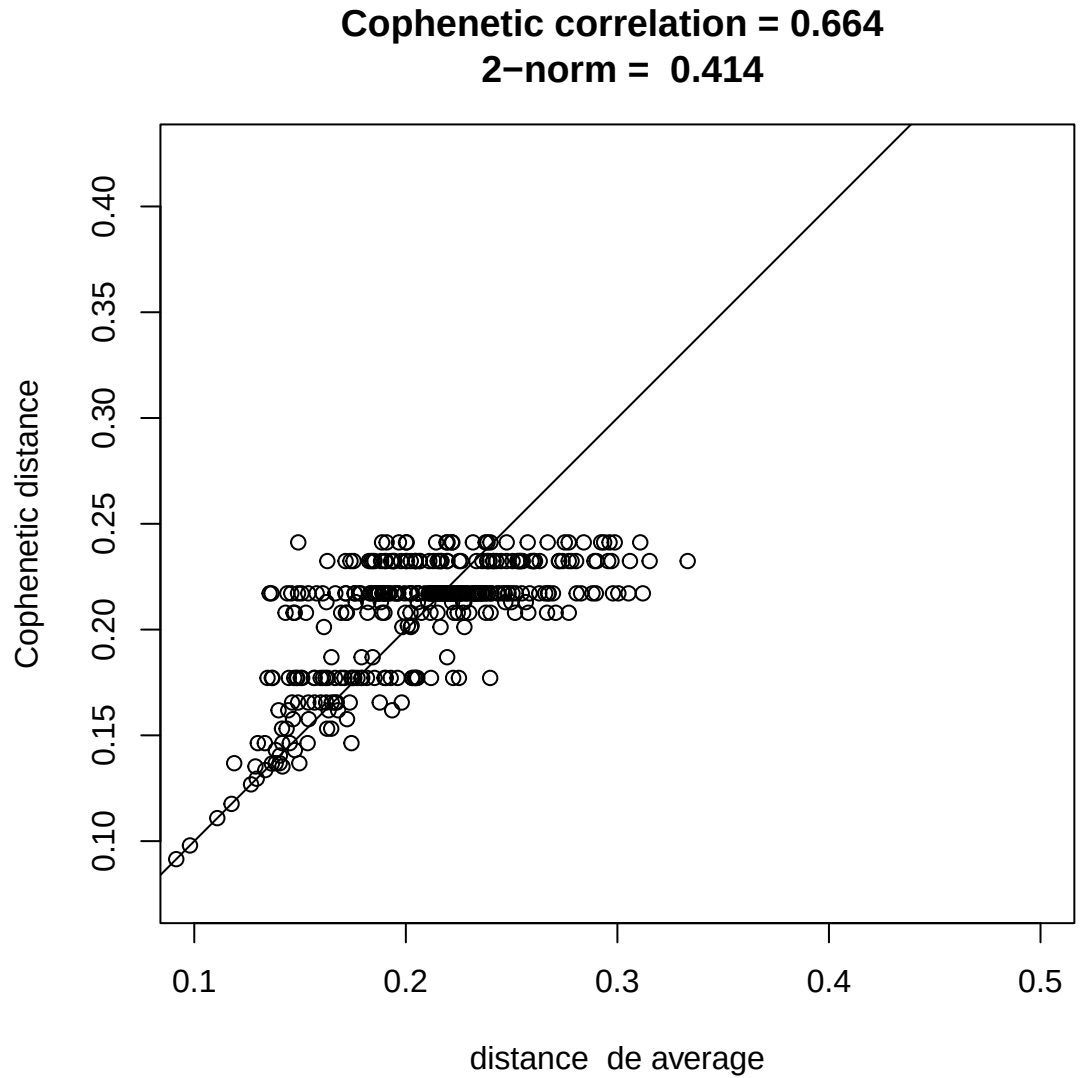

Cluster dendrogram with p−values (%)
au bp
edge #
Height
0.05
0.10
0.15
0.20
RUNA8A
RUNA3C
RUNA1A
RUNA1B
RUNA8B
RUNA9B
RUNA7B
RUNA8C
RUNA7A
RUNA7C
RUNA9C
RUNA5B
RUNA9A
RUNA6A
RUNA4B
RUNA4C
RUNA5A
RUNA4A
RUNA5C
RUNA6B
RUNA6C
RUNA1C
RUNA2A
RUNA3A
RUNA2C
RUNA2B
RUNA3B
Distance: braycurtis
Cluster method: average


Cophenetic correlation = 0.664
2−norm = 0.414
Cophenetic distance
0.10
0.15
0.20
0.25
0.30
0.35
0.40
0.1
0.2
0.3
0.4
0.5
distance de average