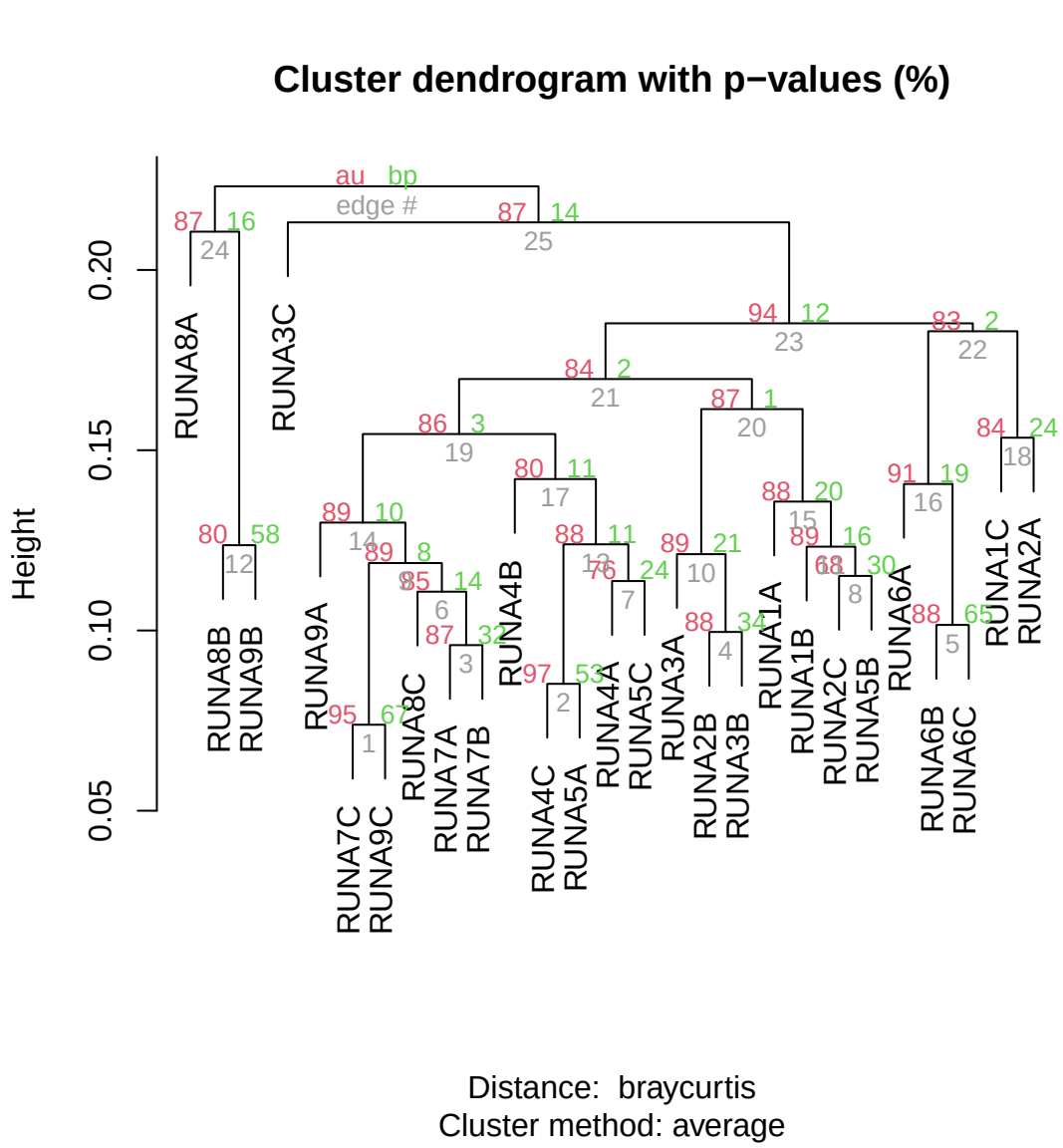
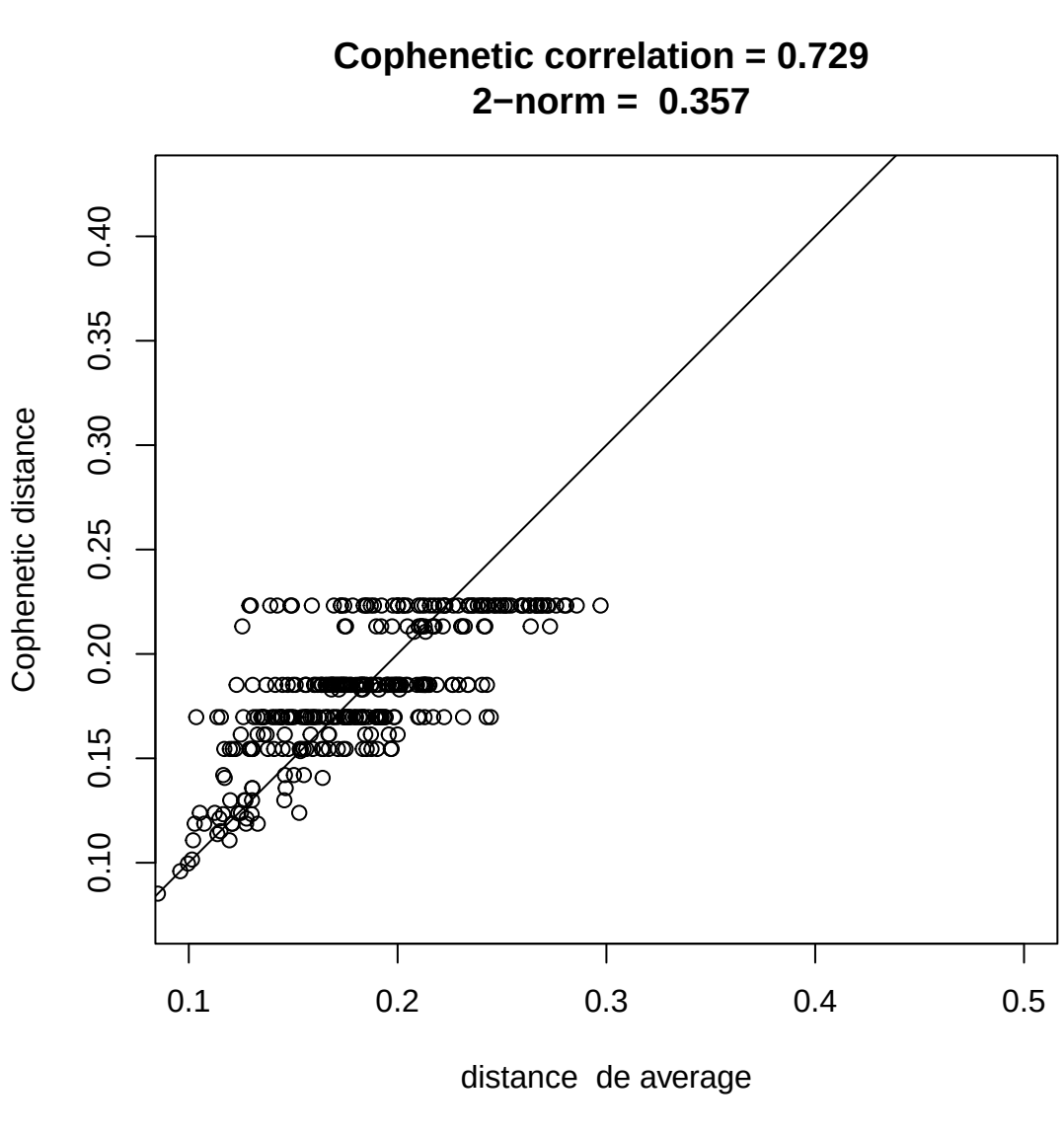

Cluster dendrogram with p−values (%)
au bp
edge #
Height
0.05
0.10
0.15
0.20
RUNA8A
RUNA3C
RUNA8B
RUNA9B
RUNA9A
RUNA8C
RUNA7A
RUNA7B
RUNA7C
RUNA9C
RUNA4B
RUNA4C
RUNA5A
RUNA4A
RUNA5C
RUNA3A
RUNA2B
RUNA3B
RUNA1A
RUNA1B
RUNA2C
RUNA5B
RUNA6A
RUNA6B
RUNA6C
RUNA1C
RUNA2A
Distance: braycurtis
Cluster method: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


Cophenetic correlation = 0.729
2−norm = 0.357
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