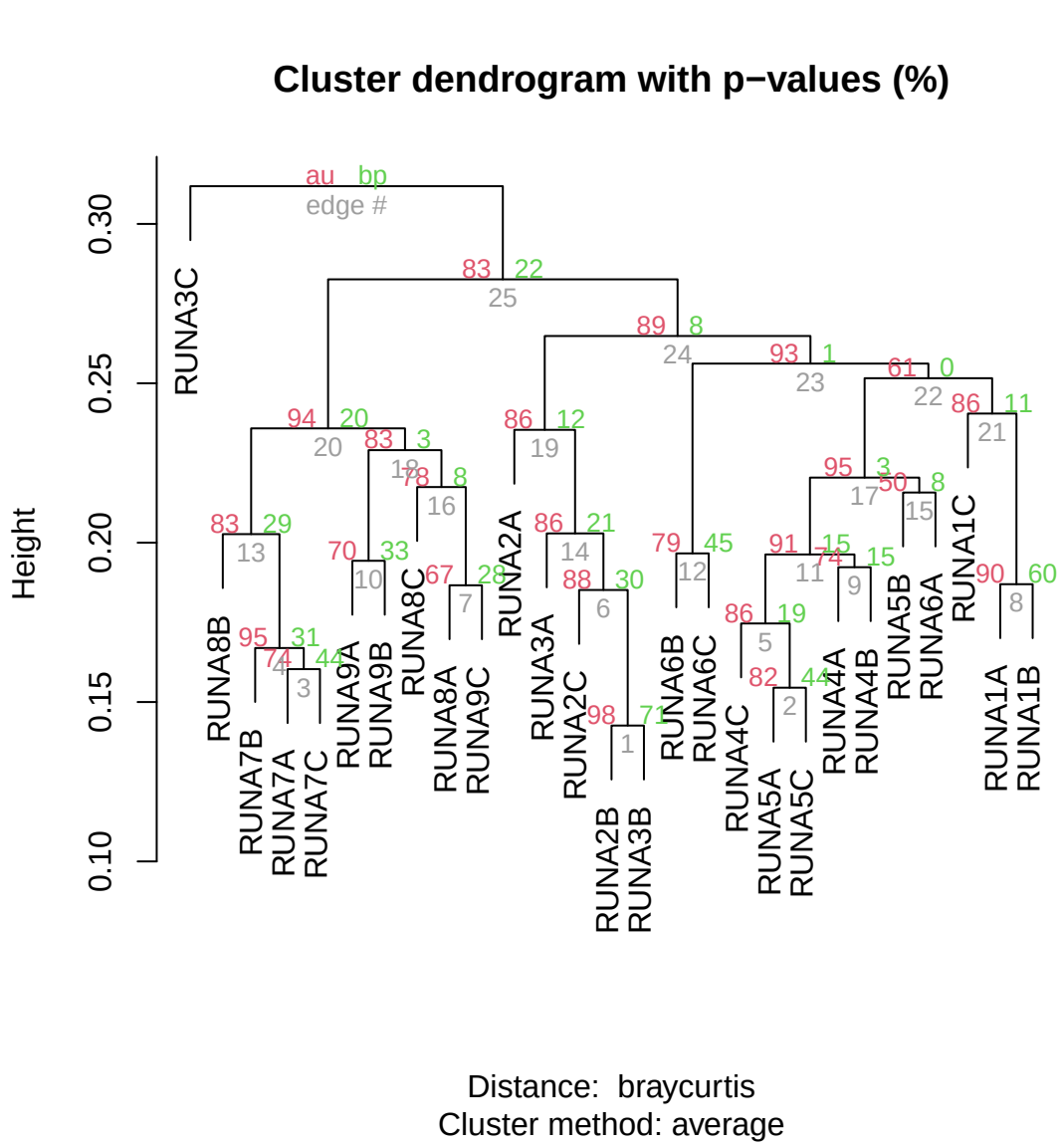
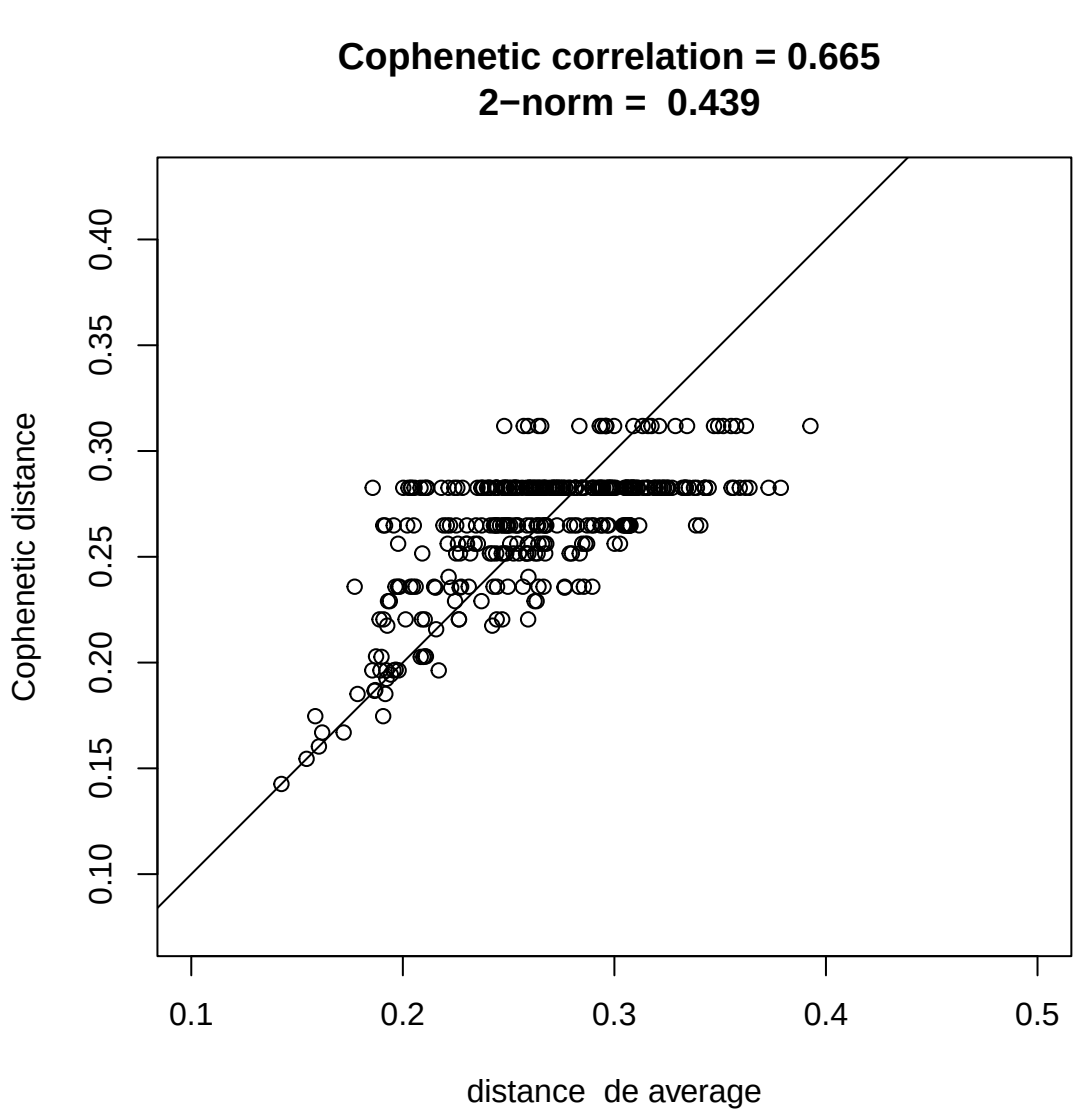

Cluster dendrogram with p−values (%)
au bp
edge #
Height
0.30
0.25
0.20
0.15
0.10
RUNA3C
RUNA8B
RUNA7B
RUNA7A
RUNA7C
RUNA9A
RUNA9B
RUNA8C
RUNA8A
RUNA9C
RUNA2A
RUNA3A
RUNA2C
RUNA2B
RUNA3B
RUNA6B
RUNA6C
RUNA4C
RUNA5A
RUNA5C
RUNA4A
RUNA4B
RUNA5B
RUNA6A
RUNA1C
RUNA1A
RUNA1B
Distance: braycurtis
Cluster method: average


Cophenetic correlation = 0.665
2−norm = 0.439
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