Clustering avec 62 sp (thresh = 99 ; method = UPGMA)

Height

RUNA8A
RUNA8B
RUNA7B
RUNA7A
RUNA7C
RUNA8C
RUNA9B
RUNA9A
RUNA9C
RUNA3C
RUNA2A
RUNA5B
RUNA1A
RUNA1B
RUNA3A
RUNA2C
RUNA2B
RUNA3B
RUNA6C
RUNA1C
RUNA6A
RUNA6B
RUNA5A
RUNA5C
RUNA4A
RUNA4B
RUNA4C

matrix.dist
hclust (*, "average")
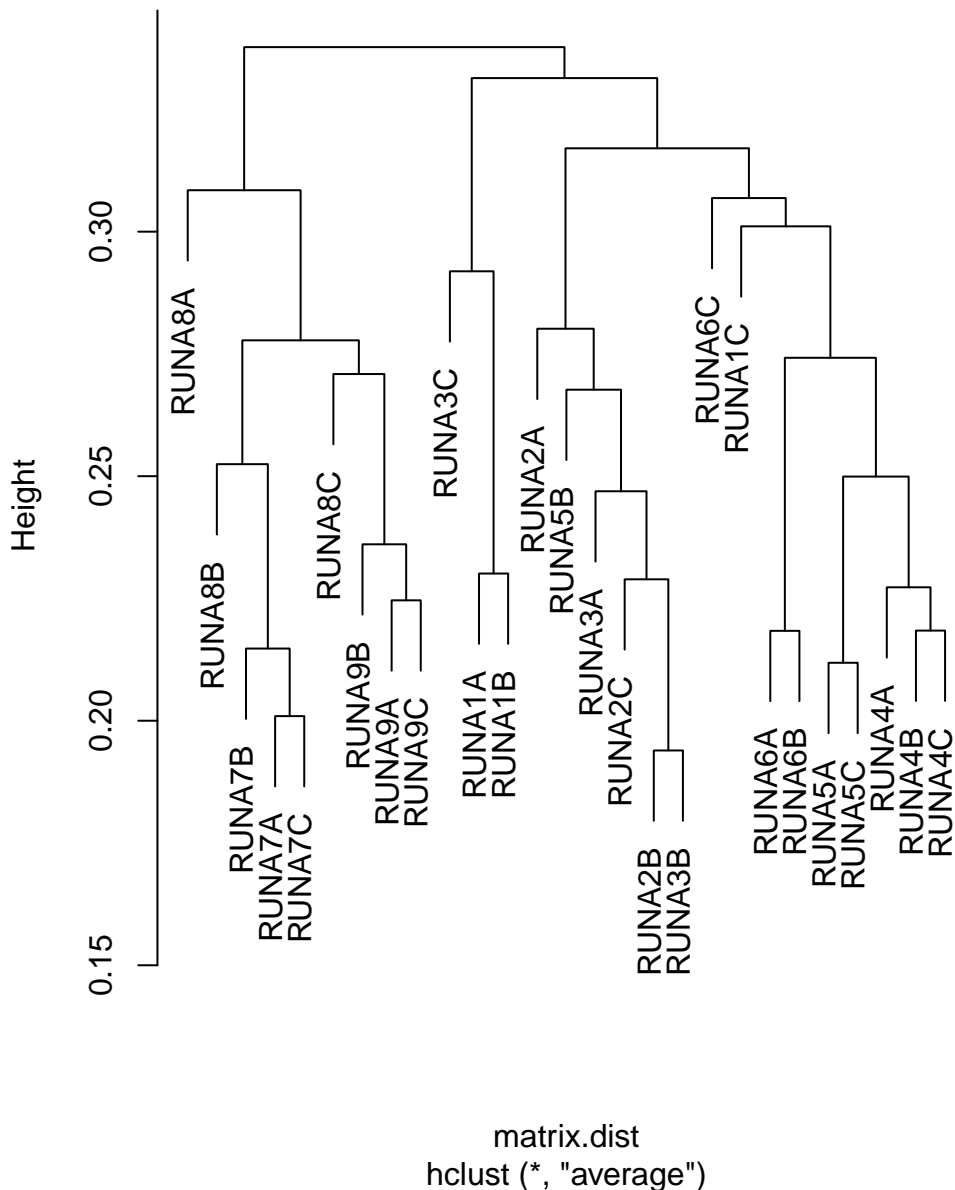
Cophenetic correlation = 0.665
2−norm = 0.454

Cophenetic distance

UPGMA distance