

Height

0.00 0.05 0.10 0.15 0.20 0.25 0.30 0.35

Time0 AS1

A

B

A2

B2

AS2 #2 gMed=196 rho12=0.25

AS3 #3 gMed=314 rho12=0.30

AS4 #4 gMed=182 rho12=0.32

AS5 #5 gMed=216 rho12=0.33

as.dist(1 - cor(log2(1 + fit\$gN[fit\$gN\$locusId %in% fit\$genesUsed,
hclust(*, "complete")