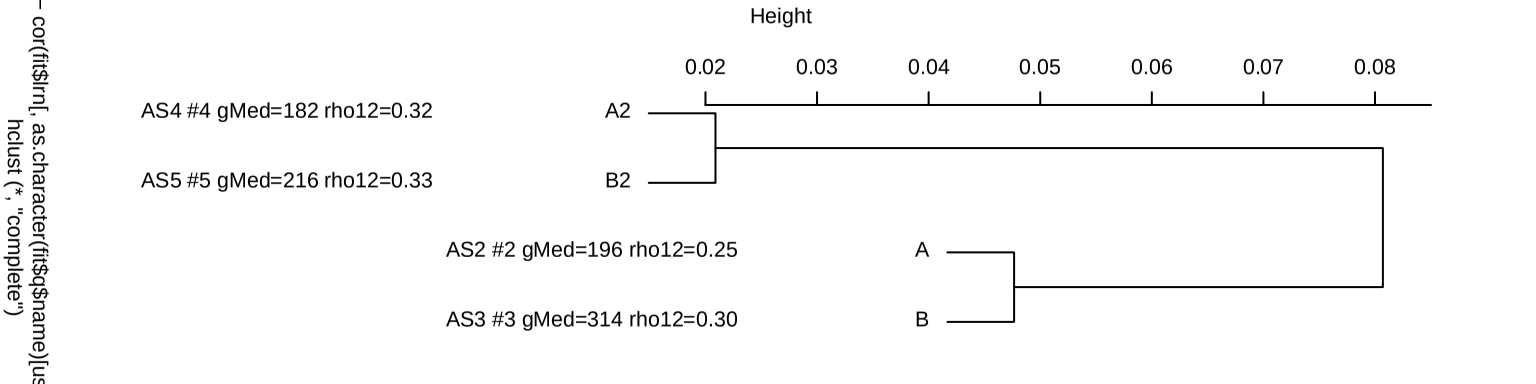
cor(fit$lrn[, as.character(fit$qs$name)[us
hclust (*, "complete")
Height
0.02
0.03
0.04
0.05
0.06
0.07
0.08
AS4 #4 gMed=182 rho12=0.32
AS5 #5 gMed=216 rho12=0.33
AS2 #2 gMed=196 rho12=0.25
AS3 #3 gMed=314 rho12=0.30
A2
B2
A
B