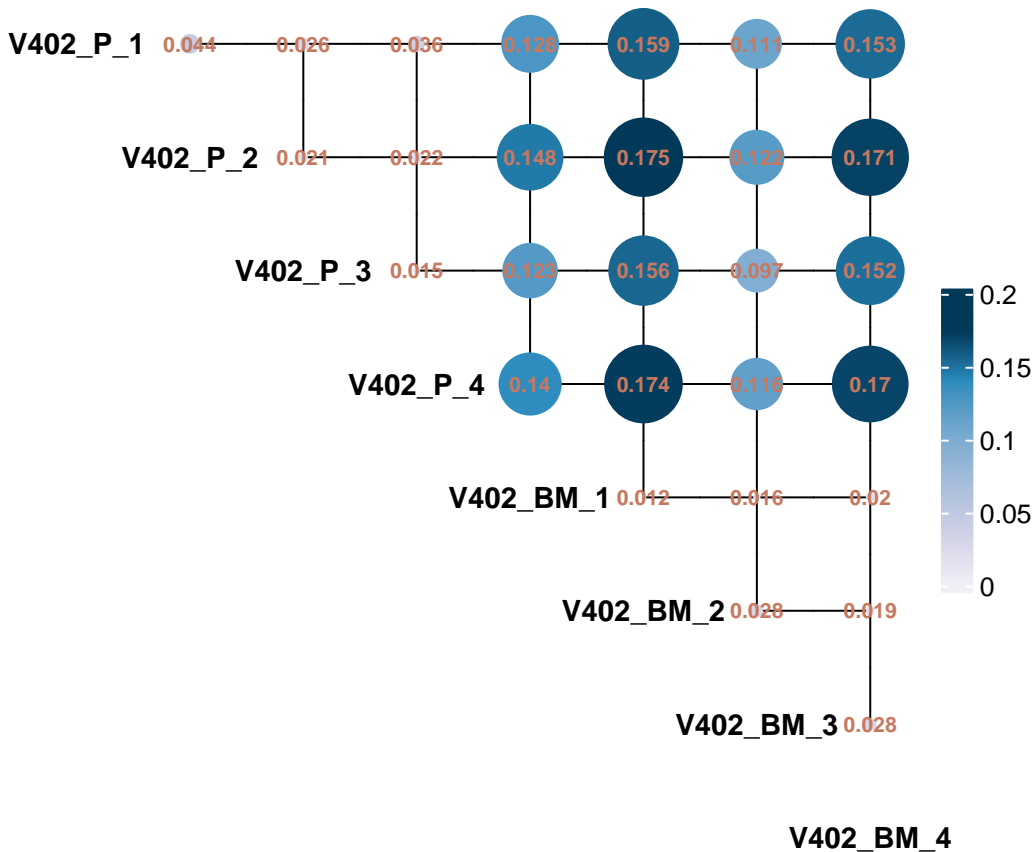
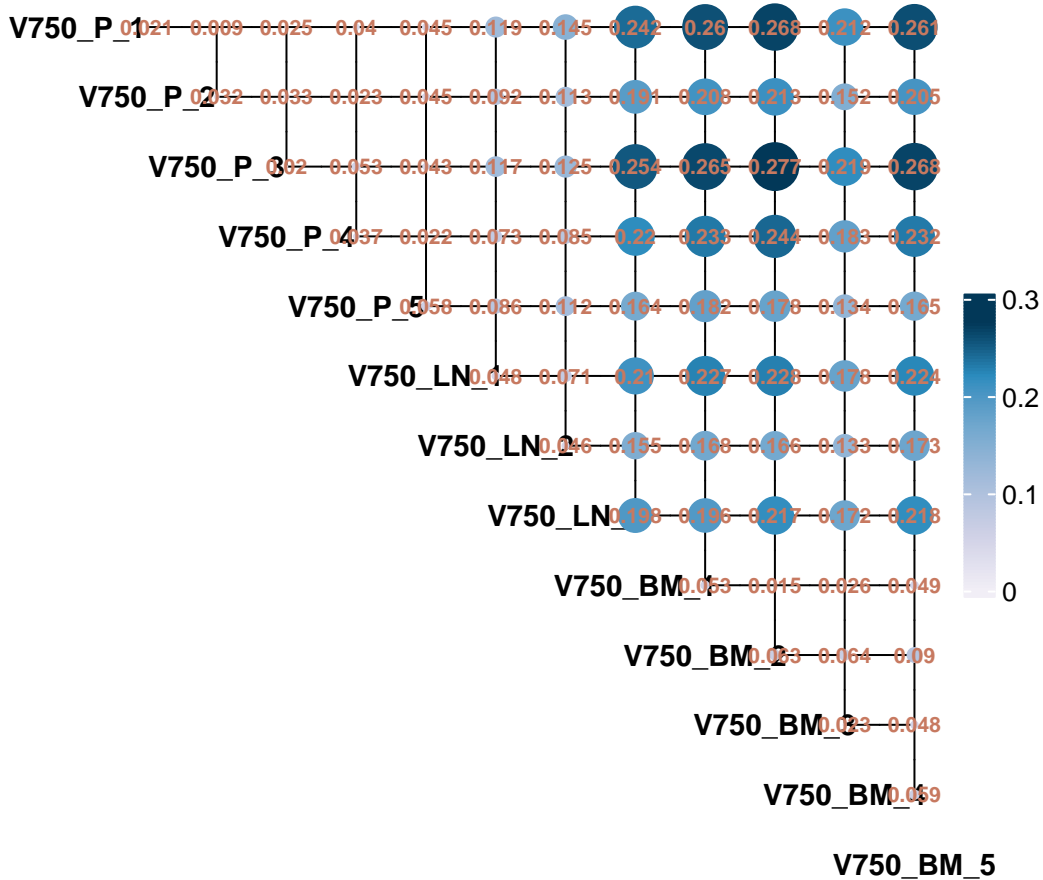


Fst of patient V402: 0.092



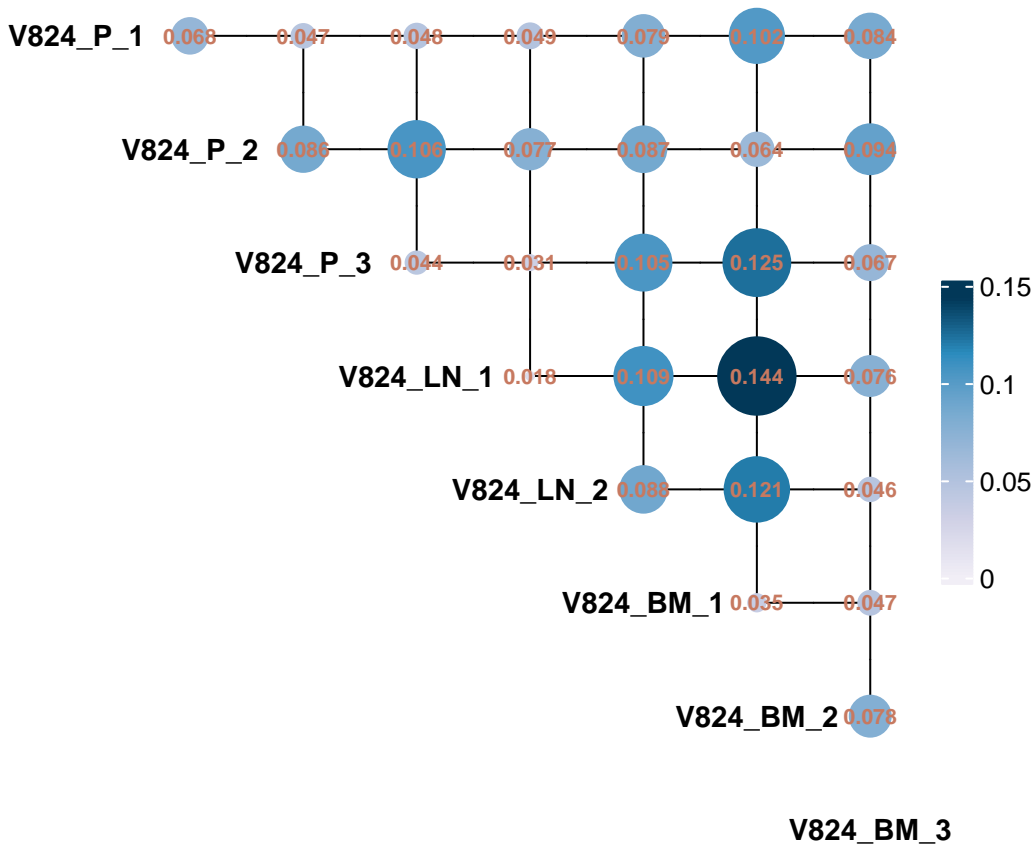
help("calEst")

Fst of patient V750: 0.1349



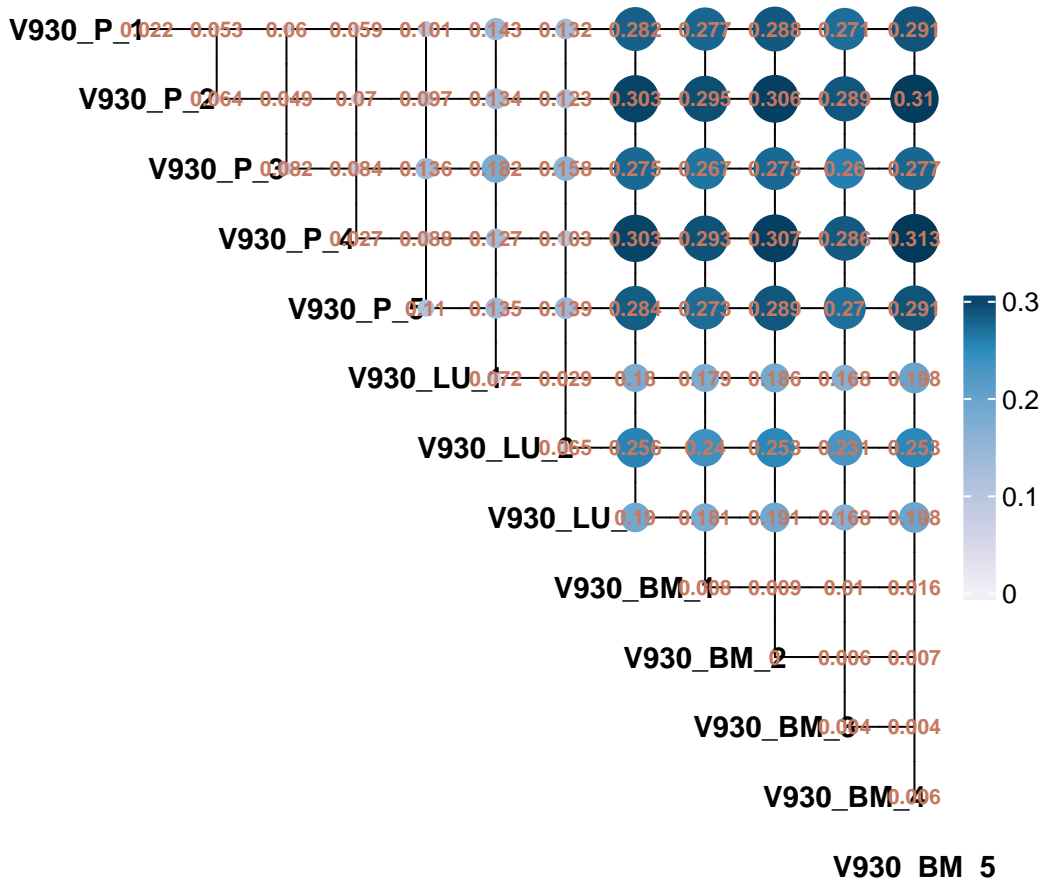
help("calEst")

Fst of patient V824: 0.076



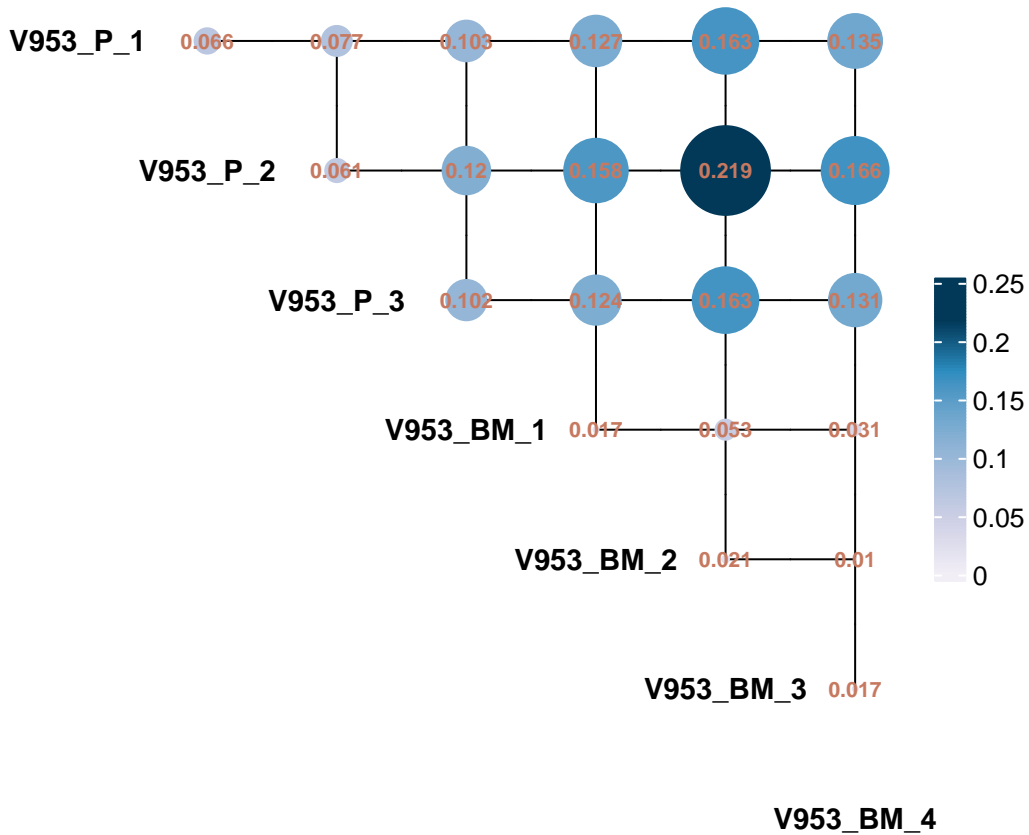
help("calEst")

Fst of patient V930: 0.166223

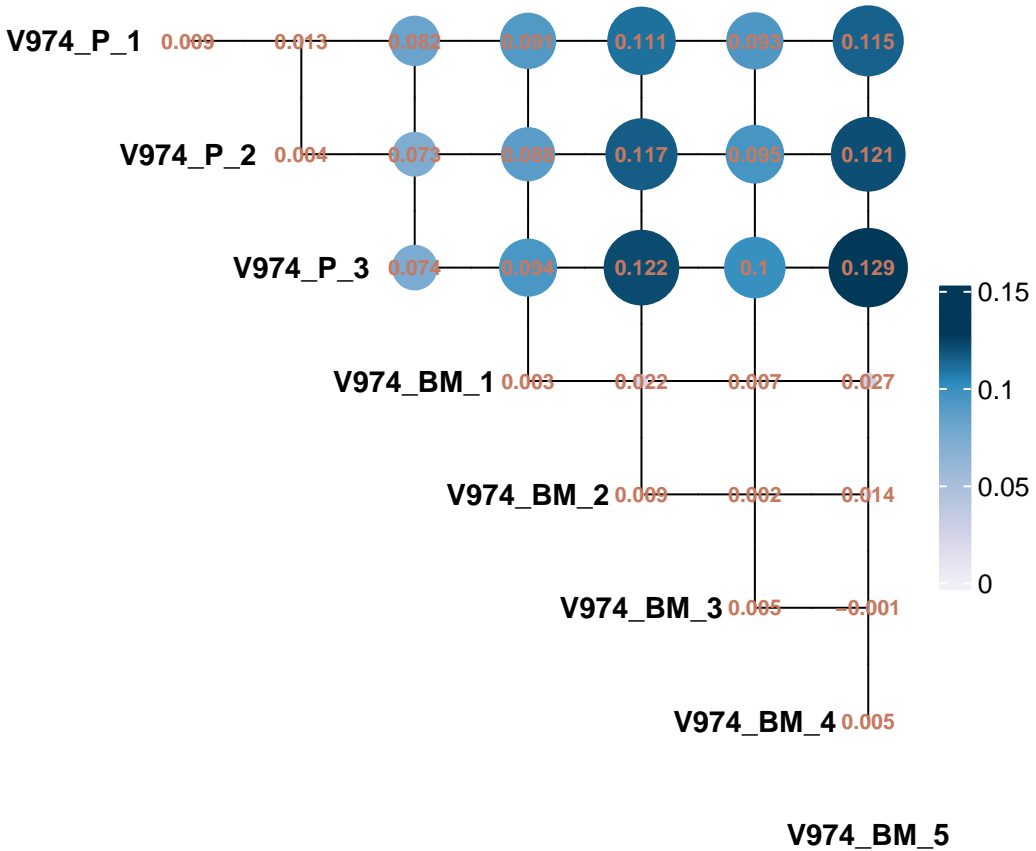


help("calEst")

Fst of patient V953: 0.0983

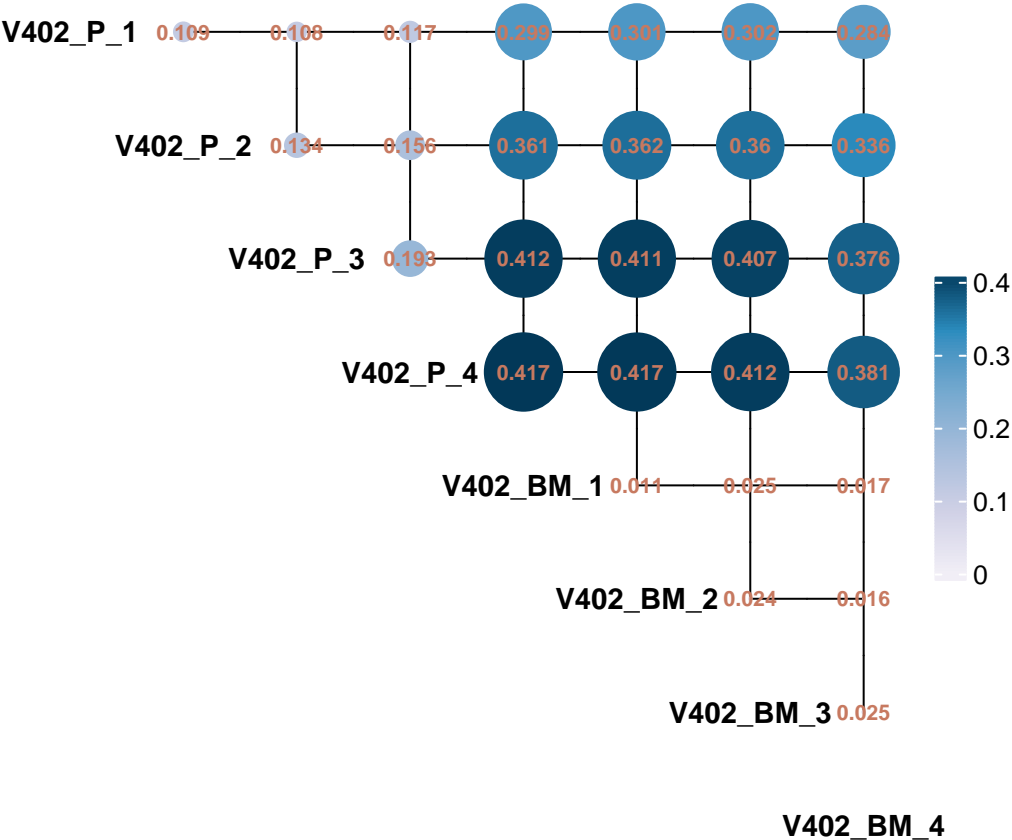


Fst of patient V974: 0.0579



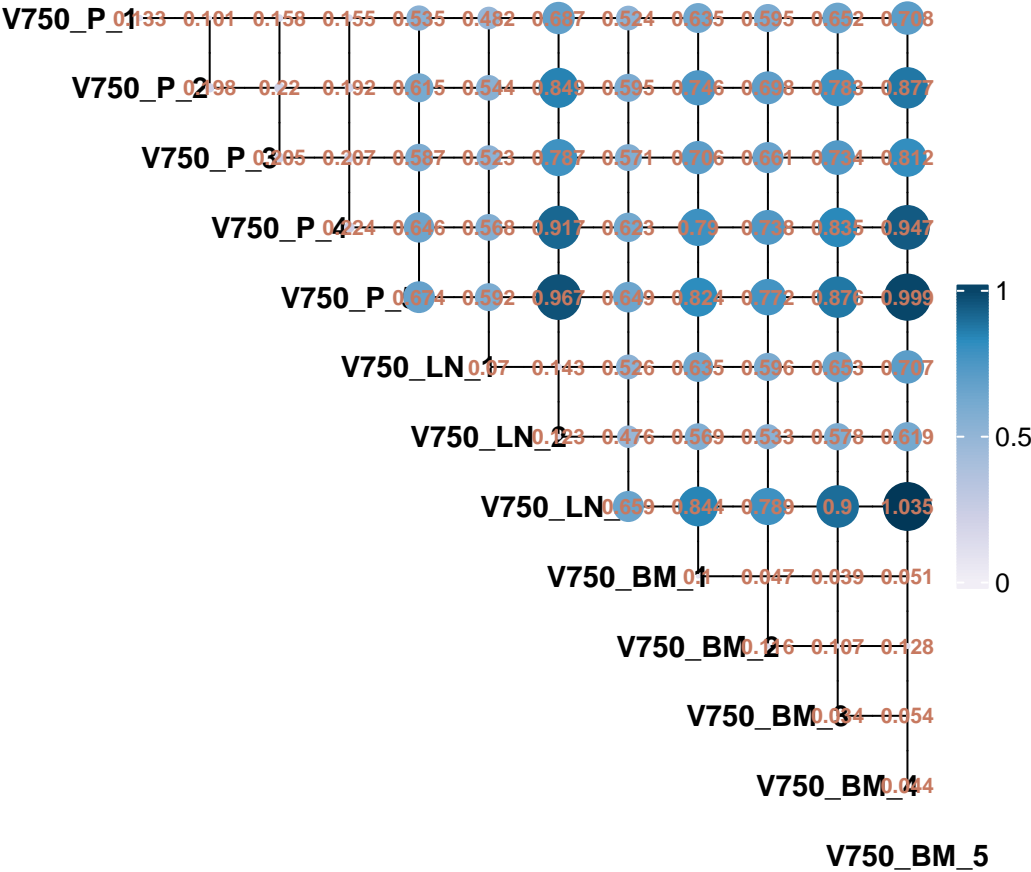
help("calEst")

Nei's distance of patient V402: 0.242



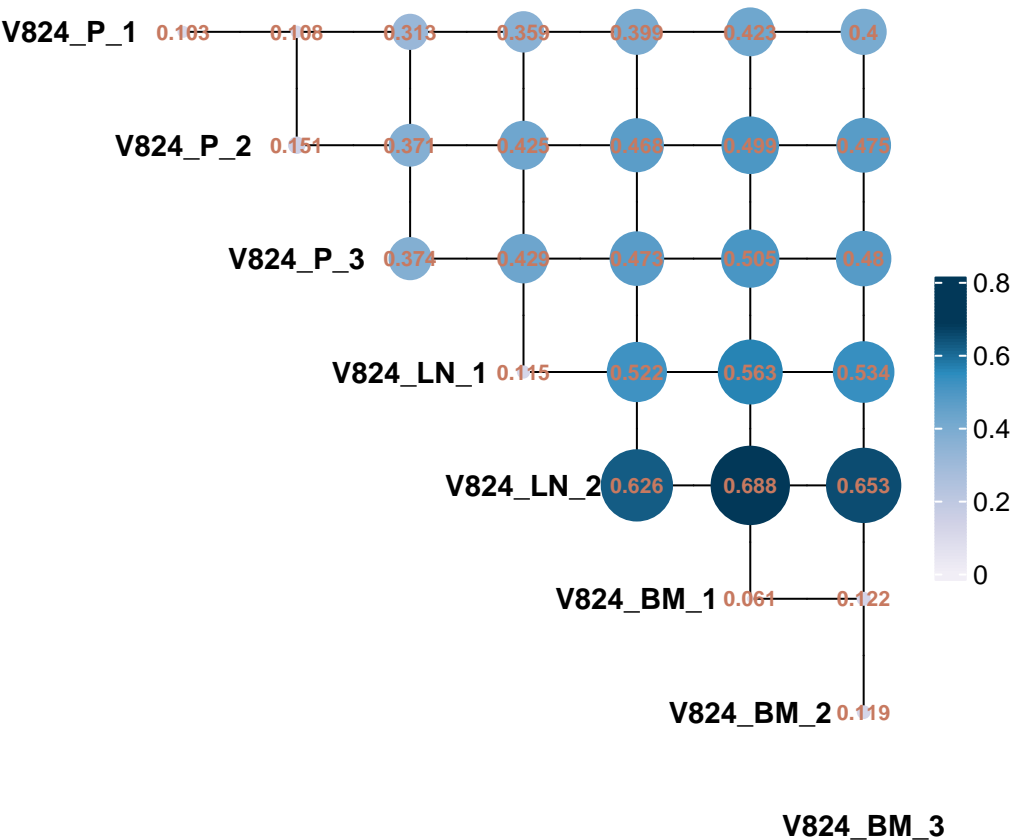
help("calNeiDist")

Nei's distance of patient V750: 0.529



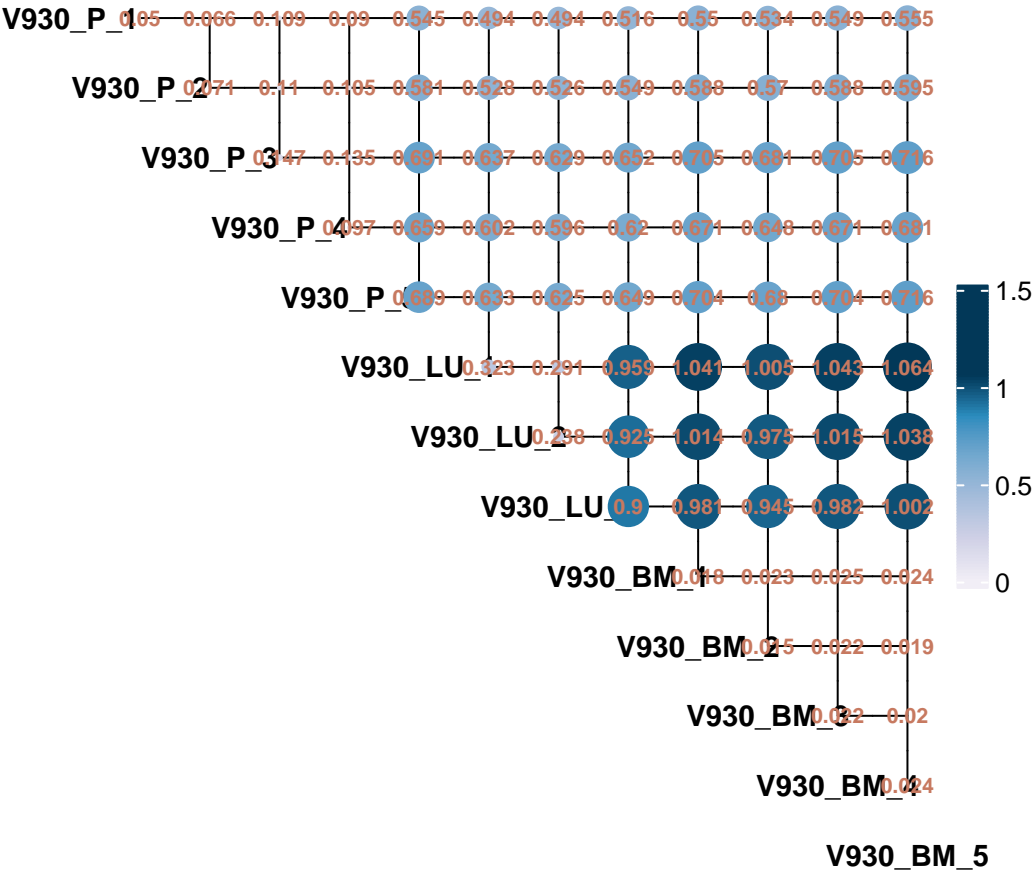
help("calNeiDist")

Nei's distance of patient V824: 0.384



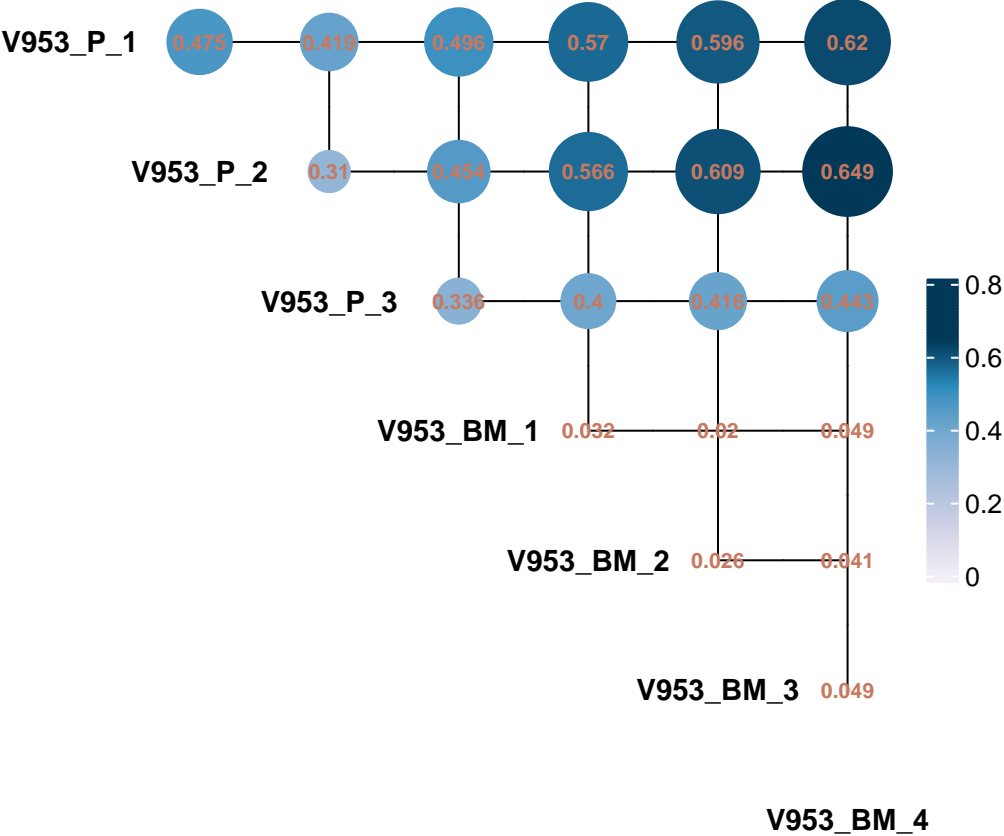
help("calNeiDist")

Nei's distance of patient V930: 0.534

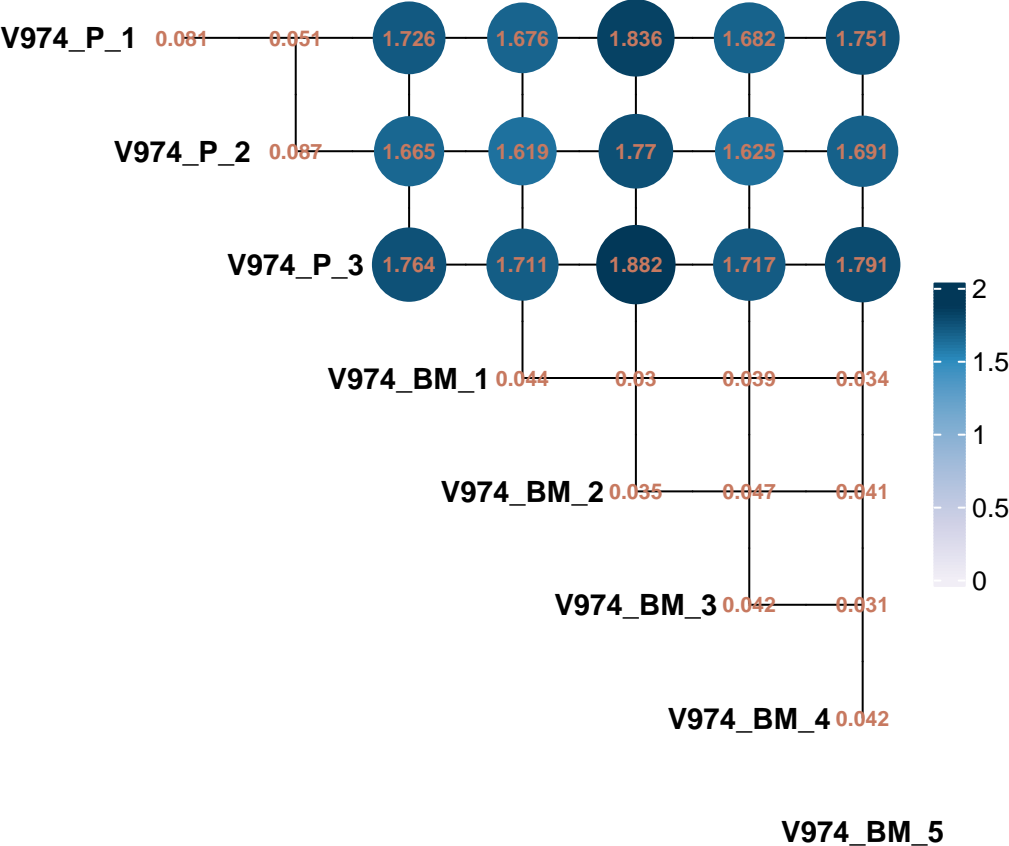


help("calNeiDist")

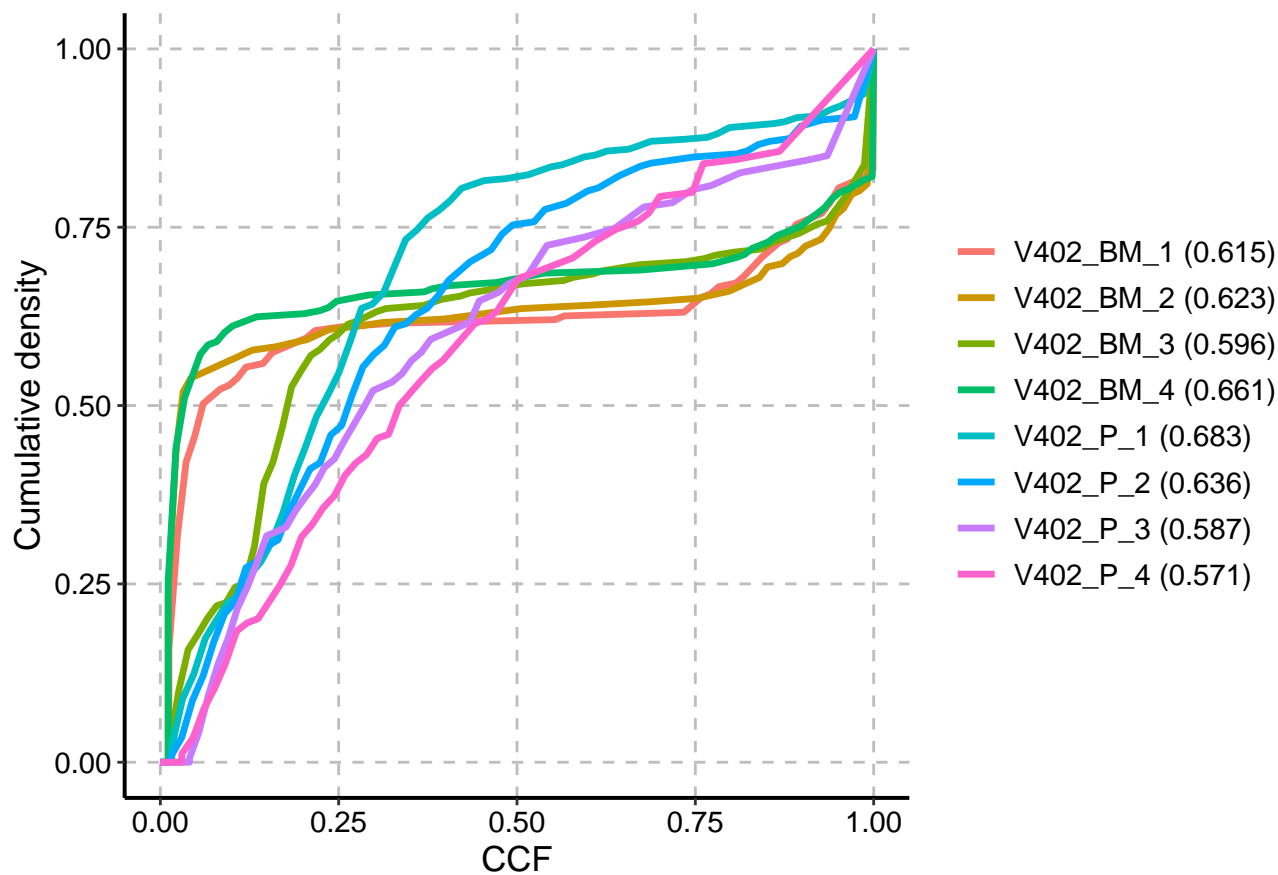
Nei's distance of patient V953: 0.361

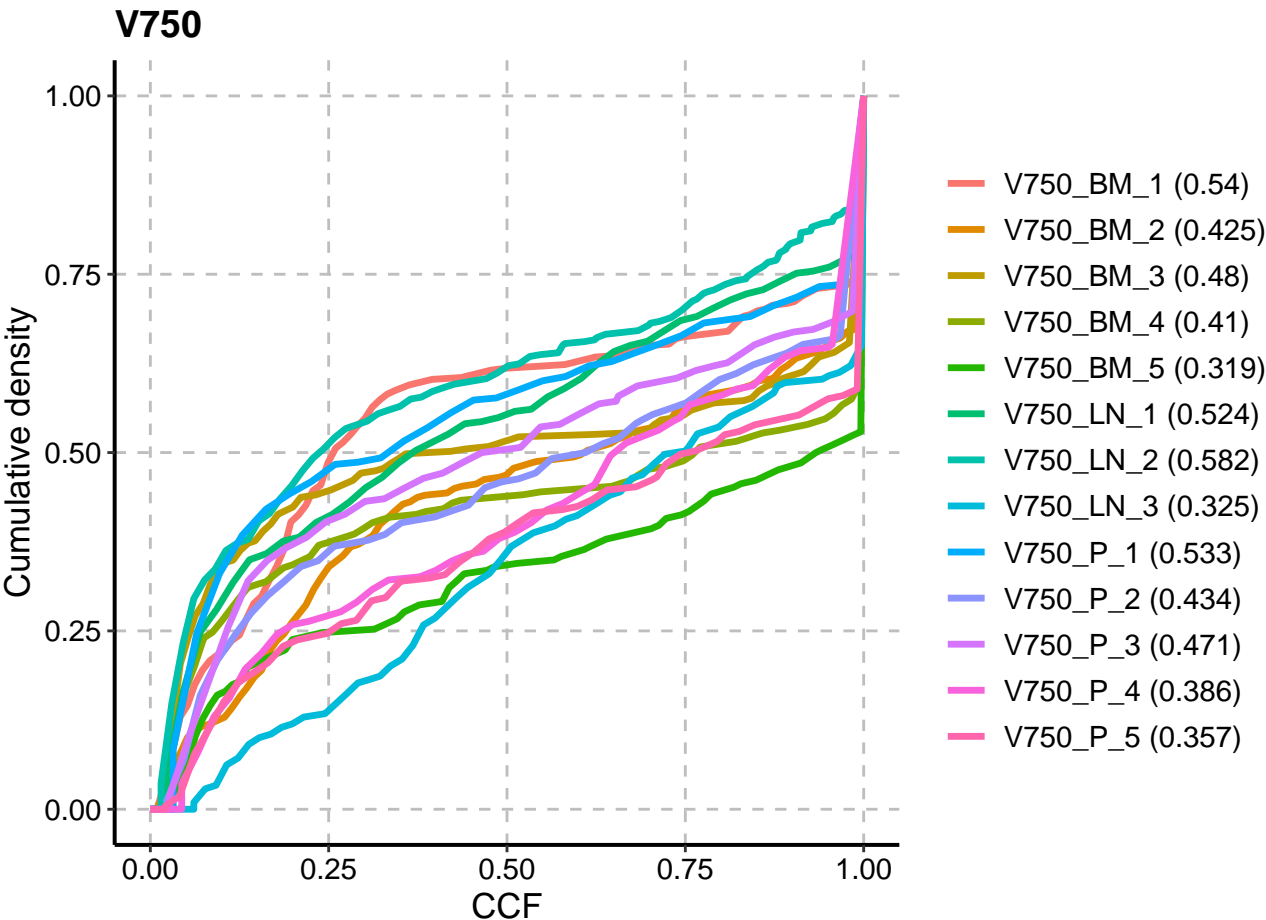


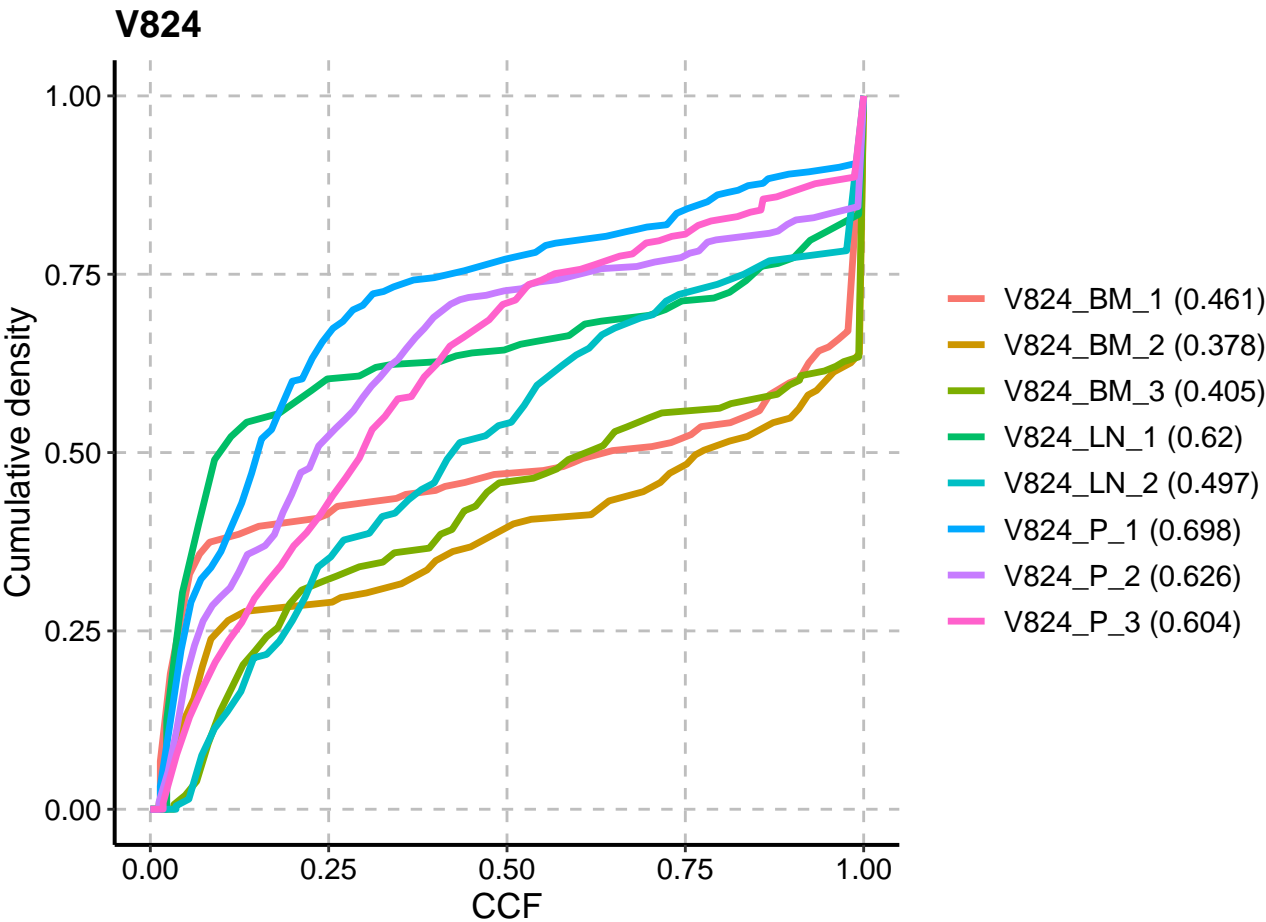
Nei's distance of patient V974: 0.947

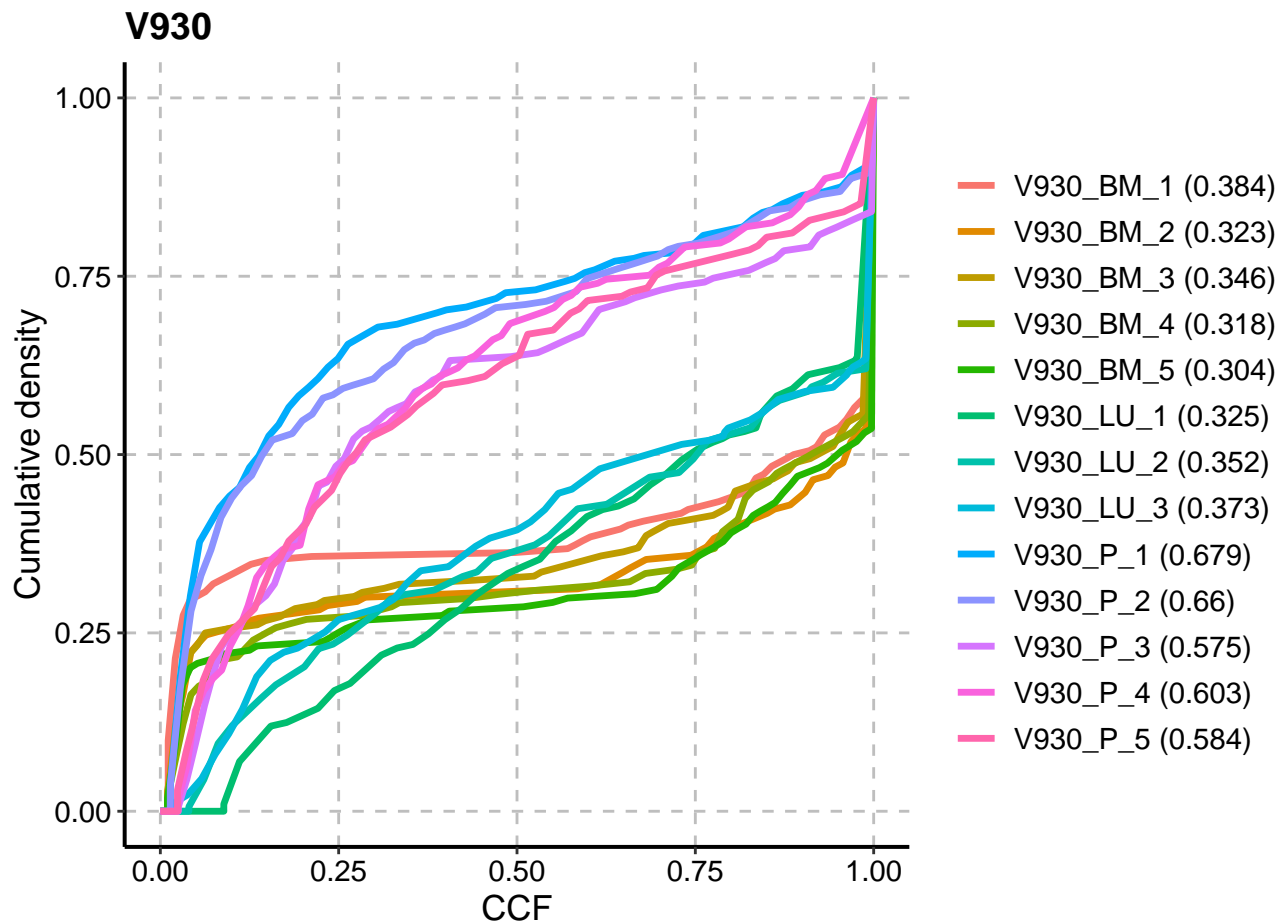


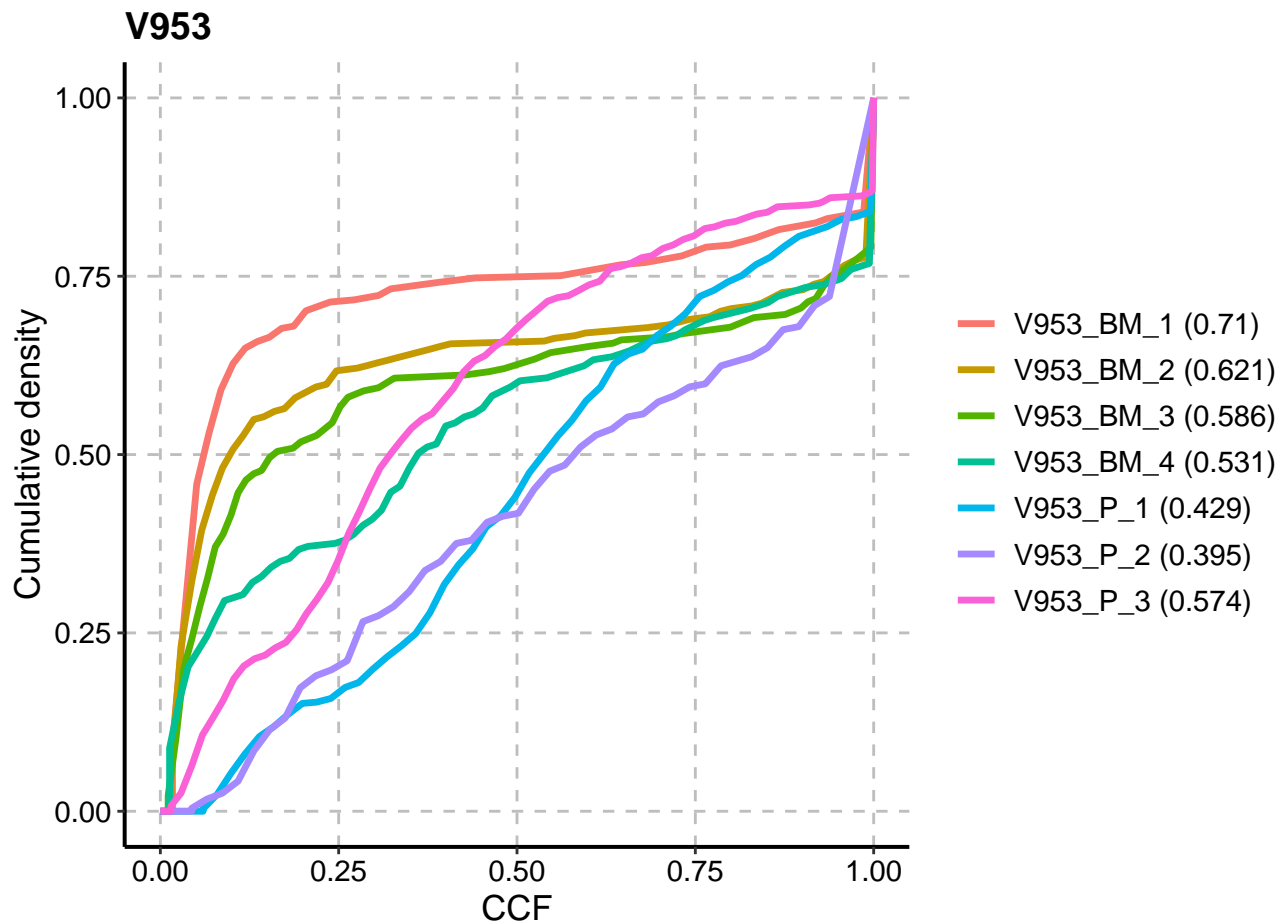
V402



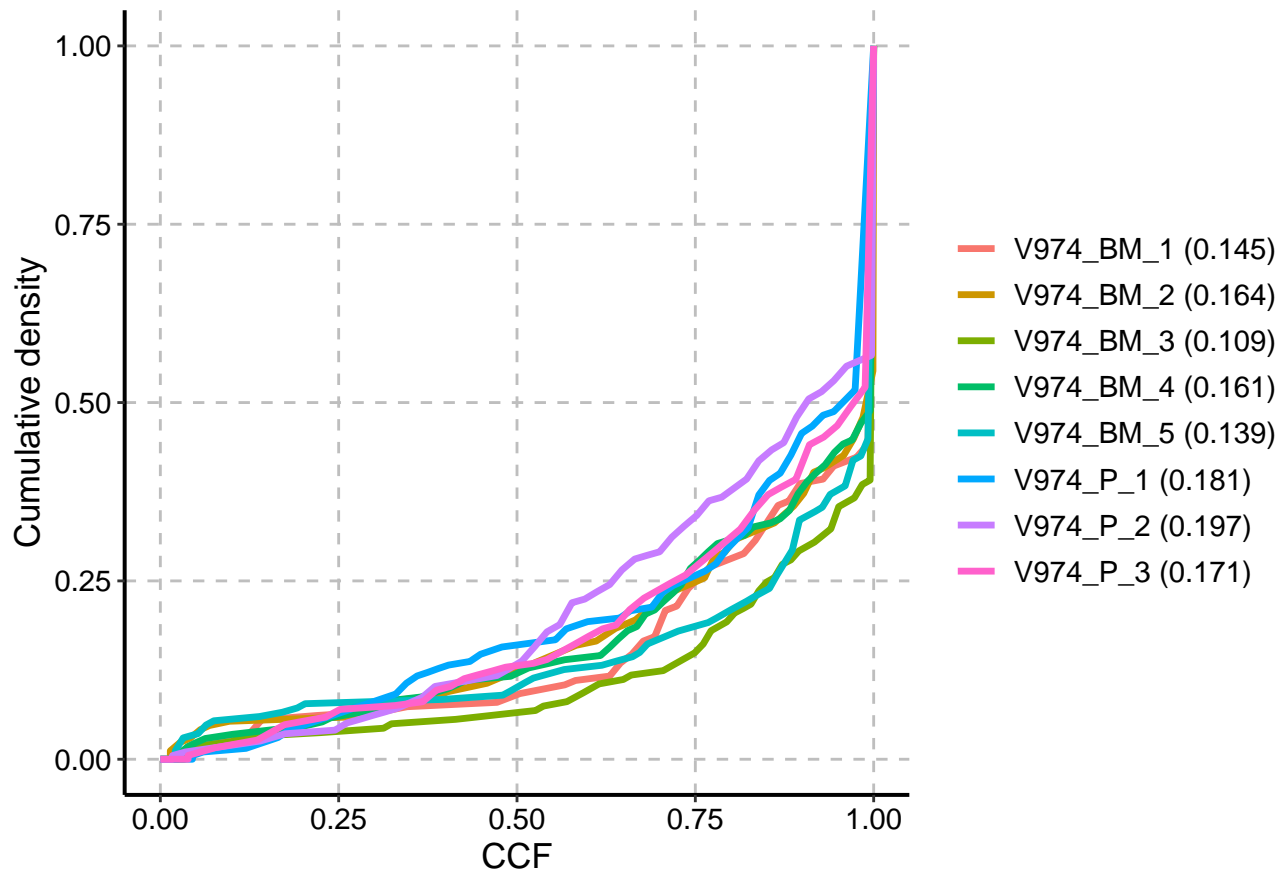




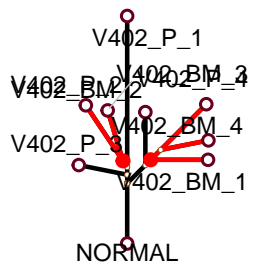




V974

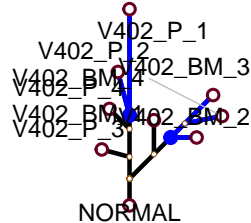


tree1: V402–NJ

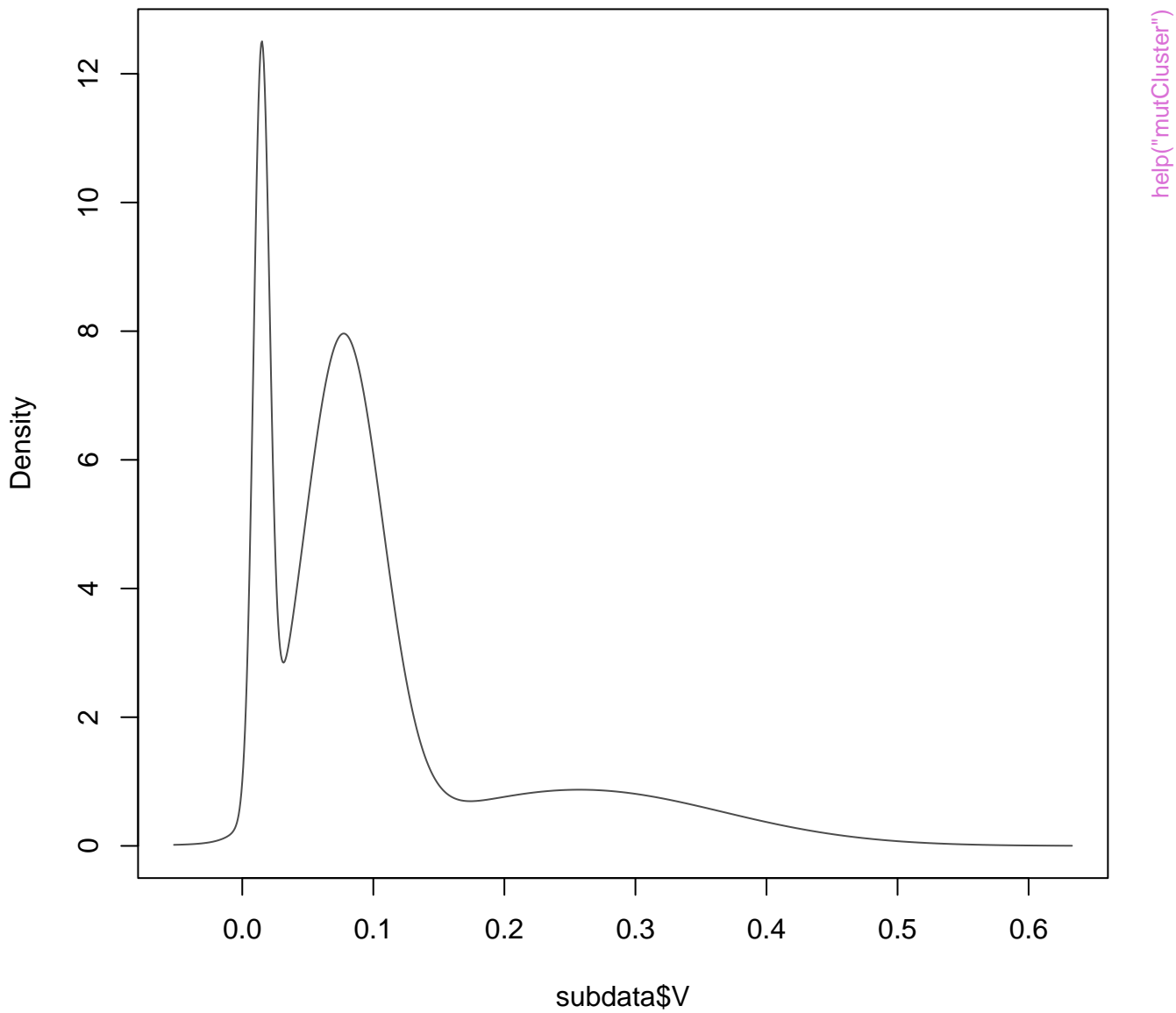


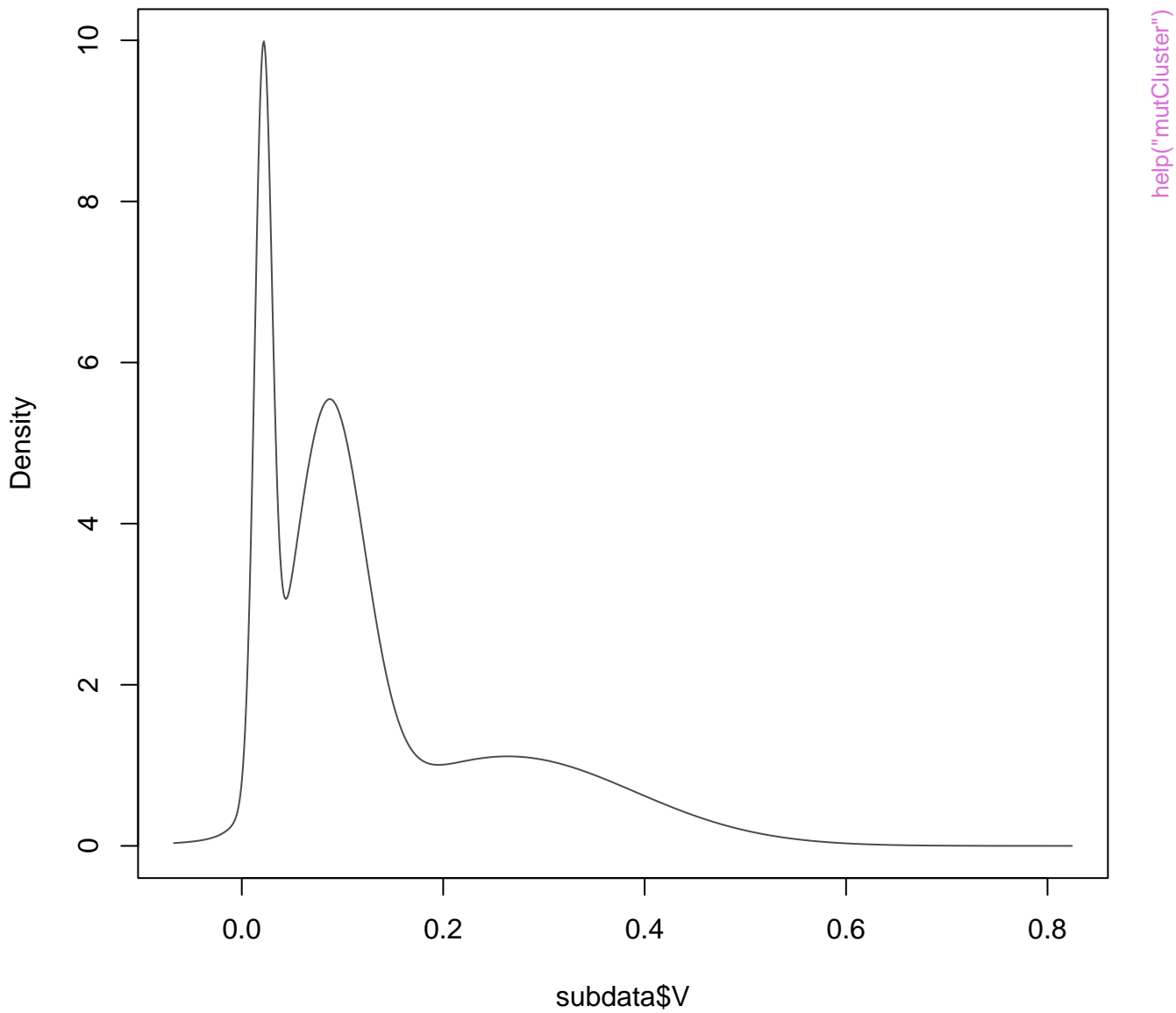
Clades absent in tree2

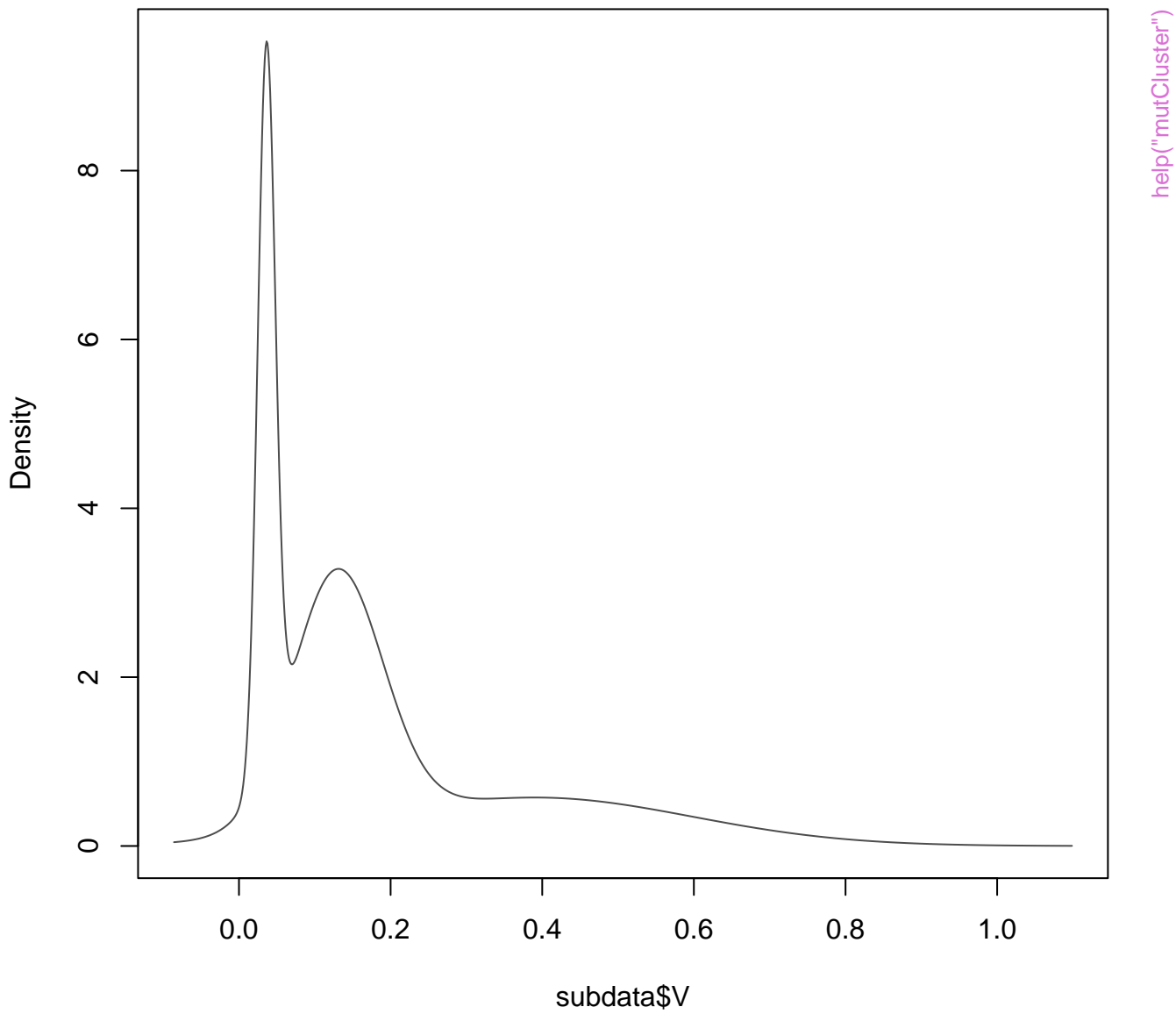
tree2: V402–MP

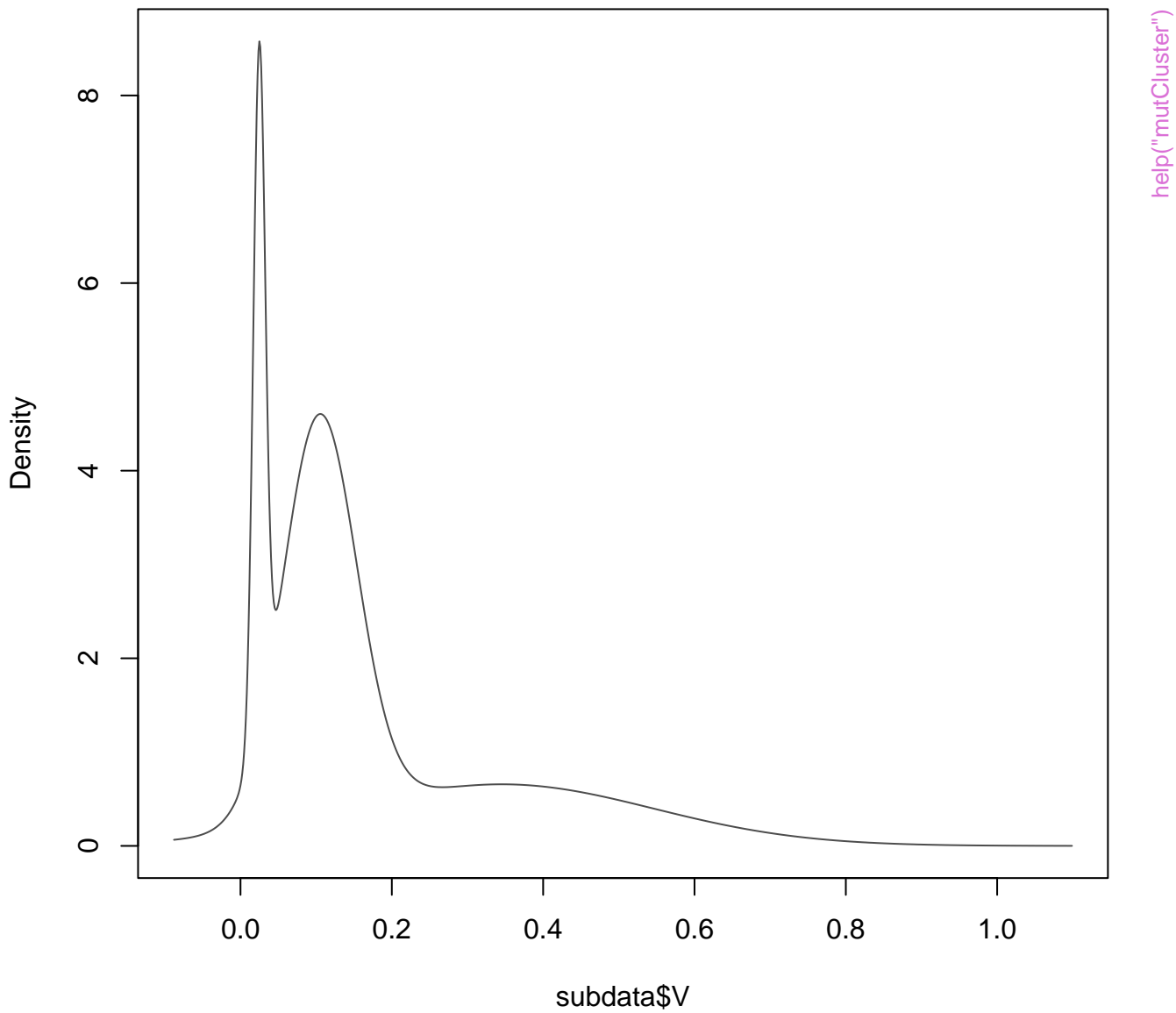


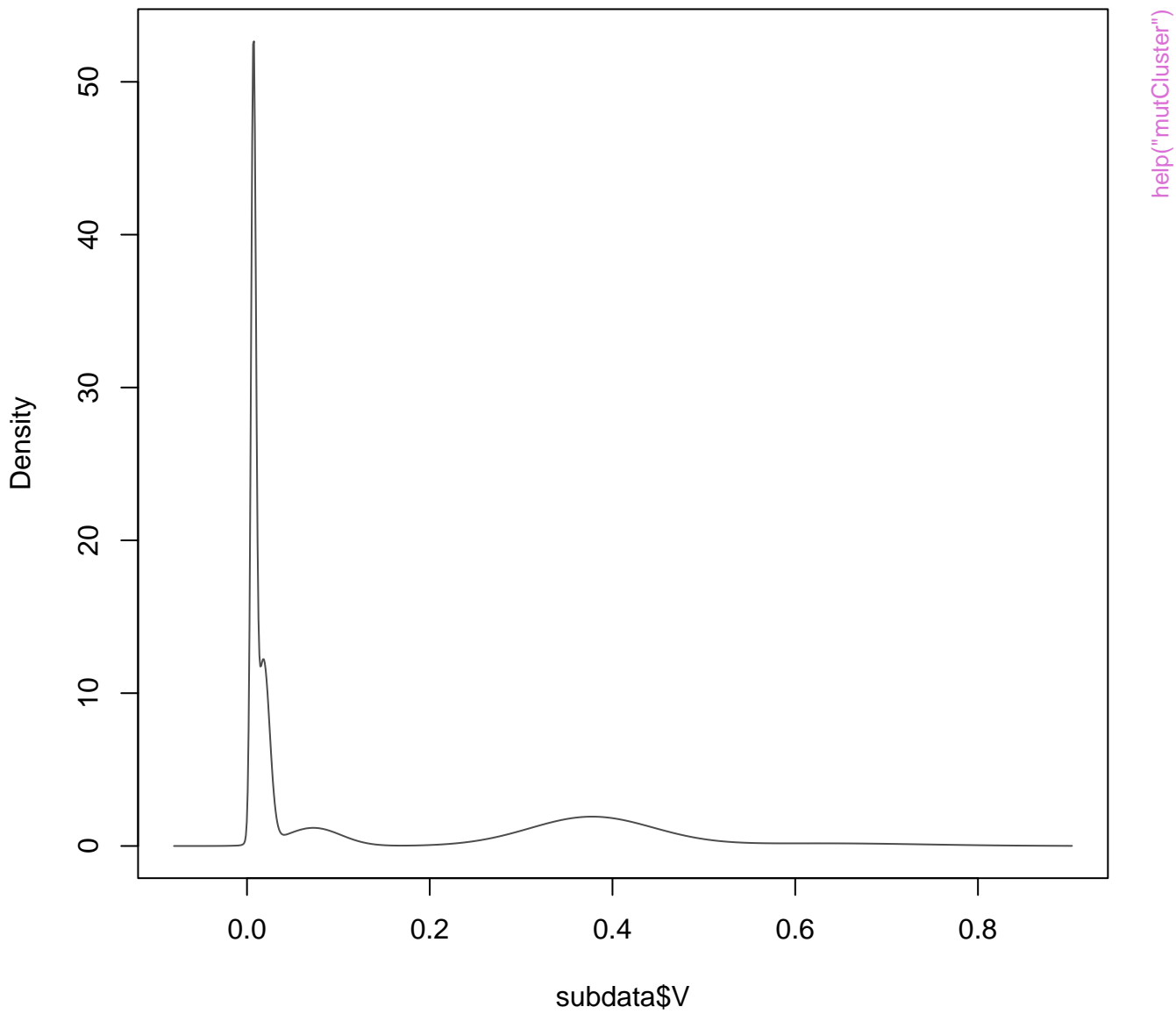
Clades absent in tree1

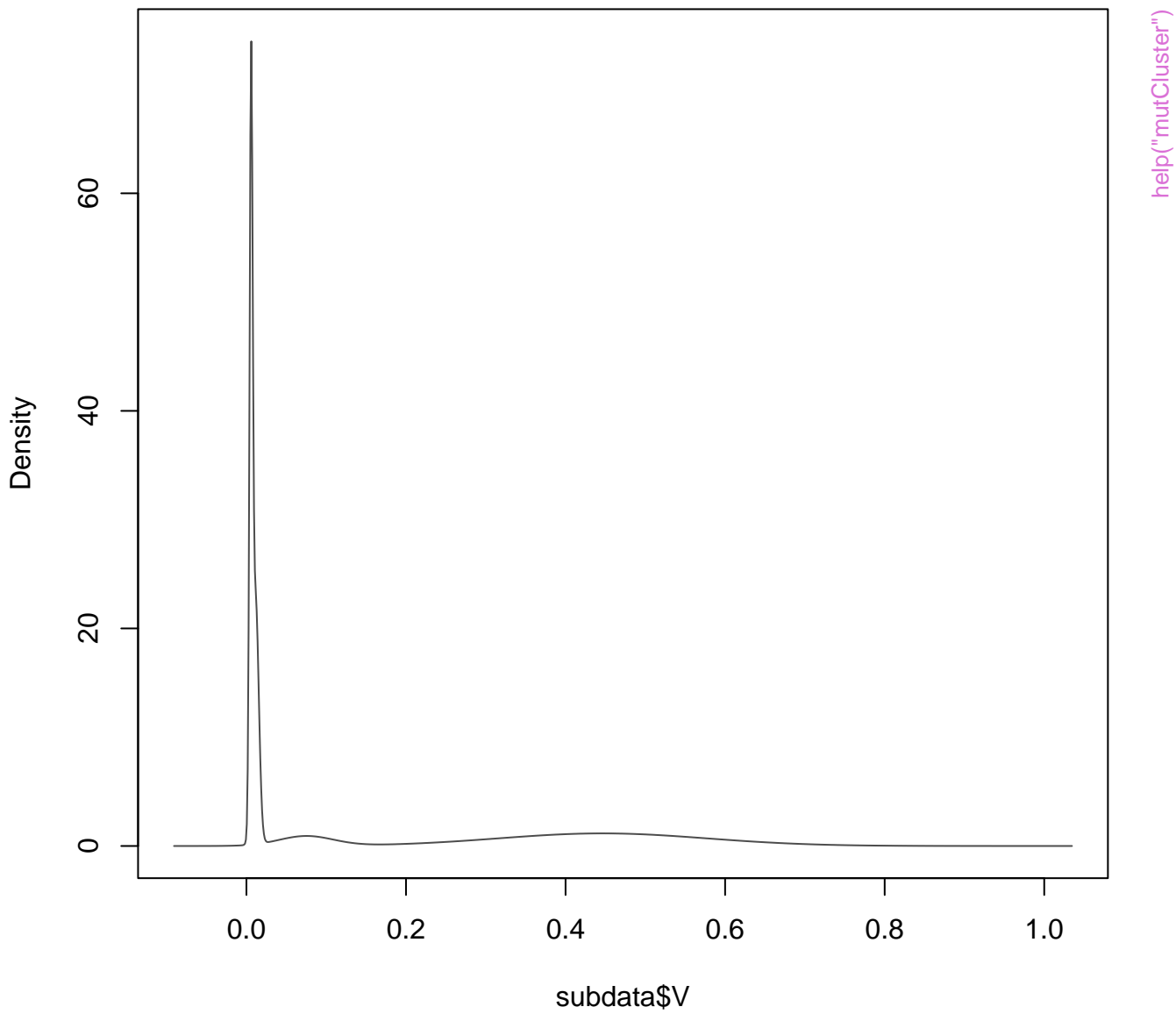


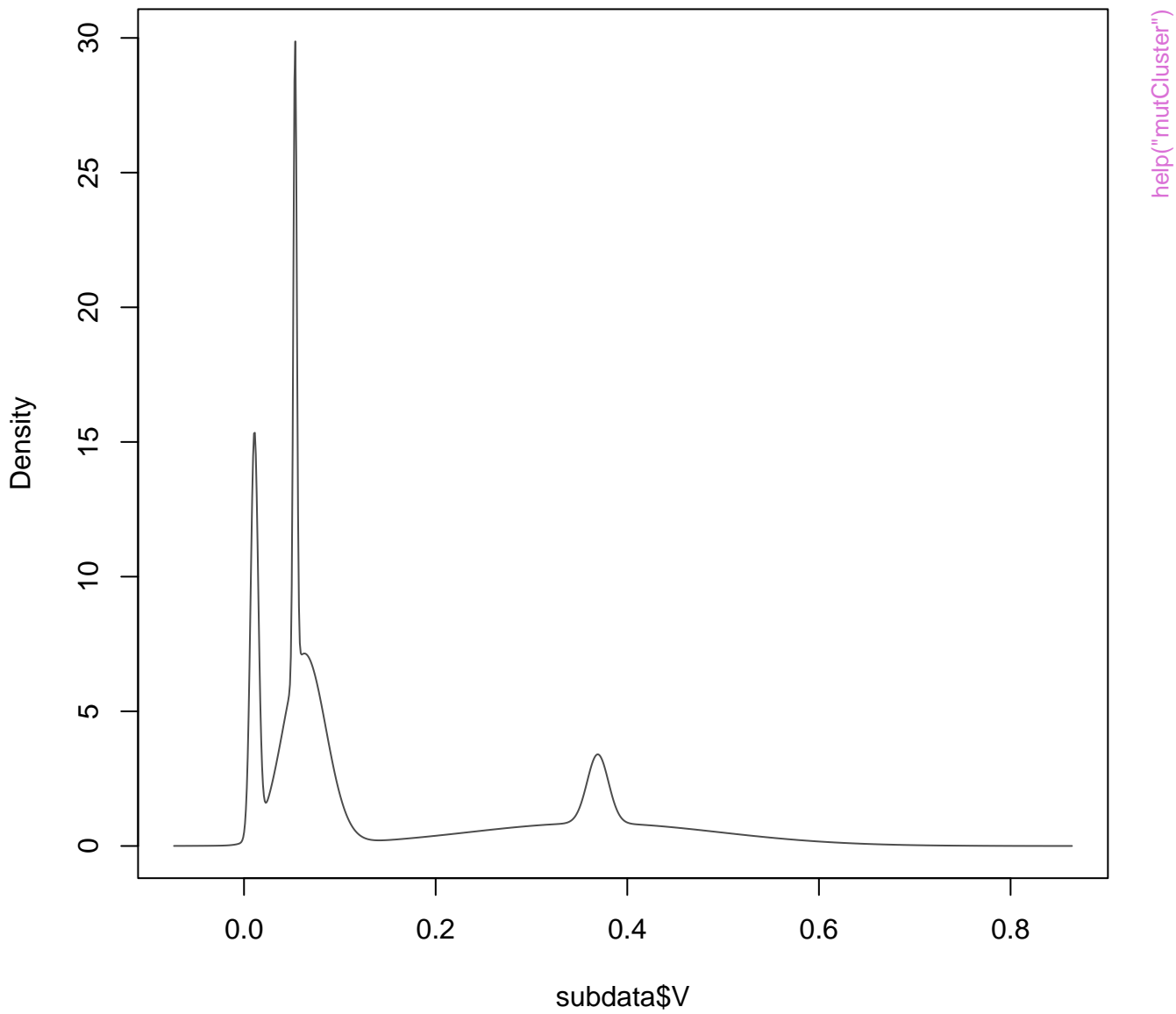


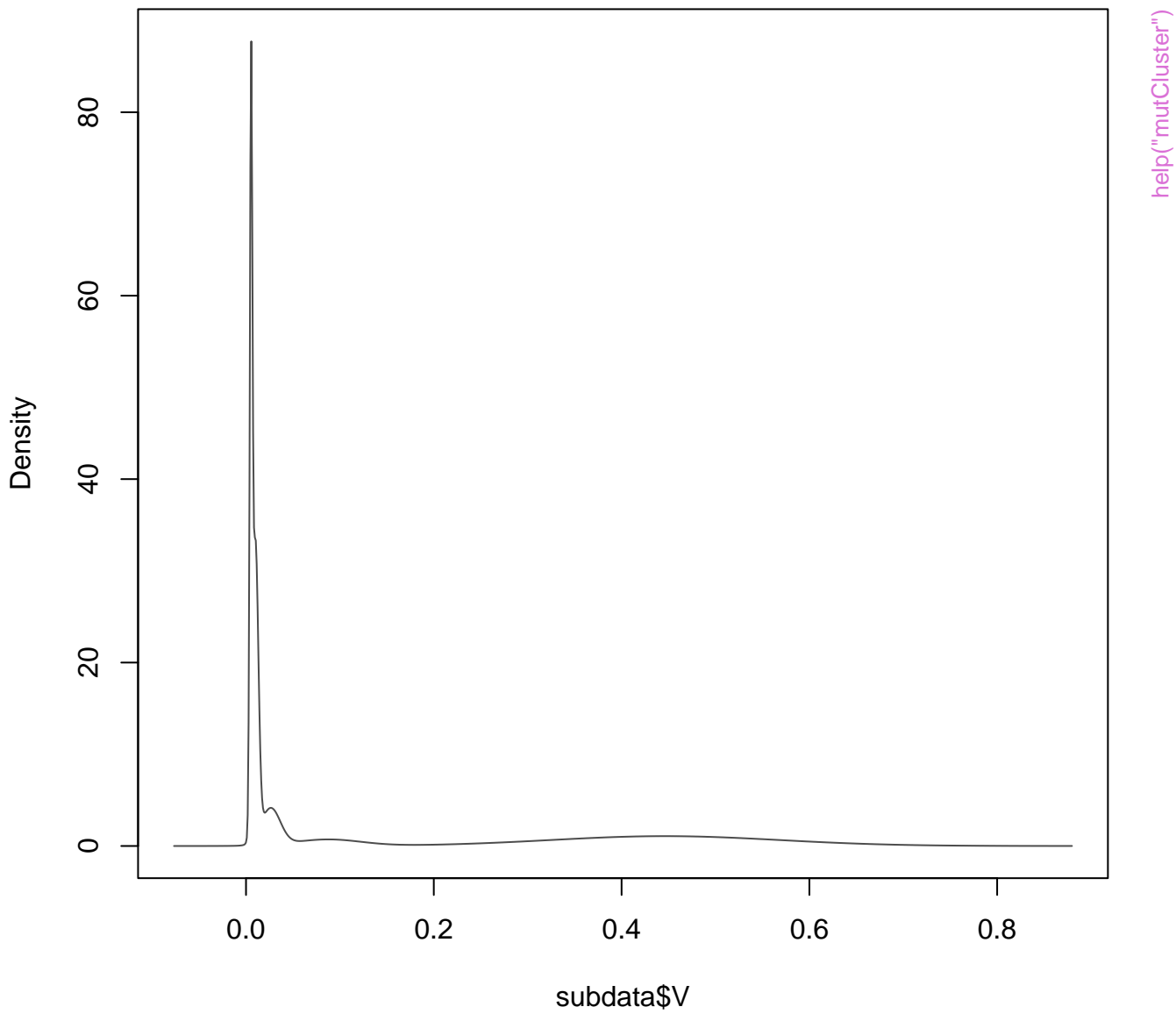




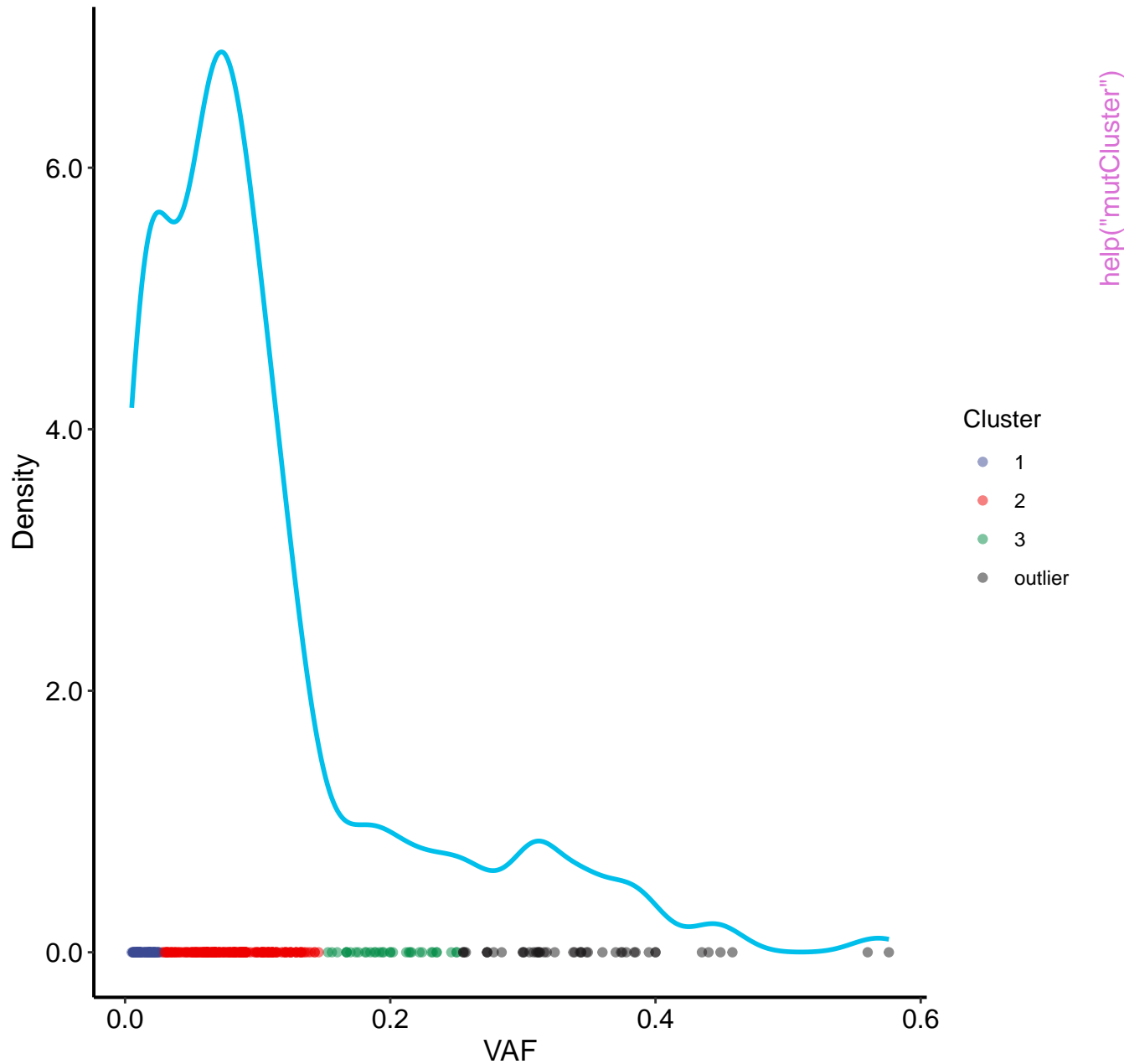




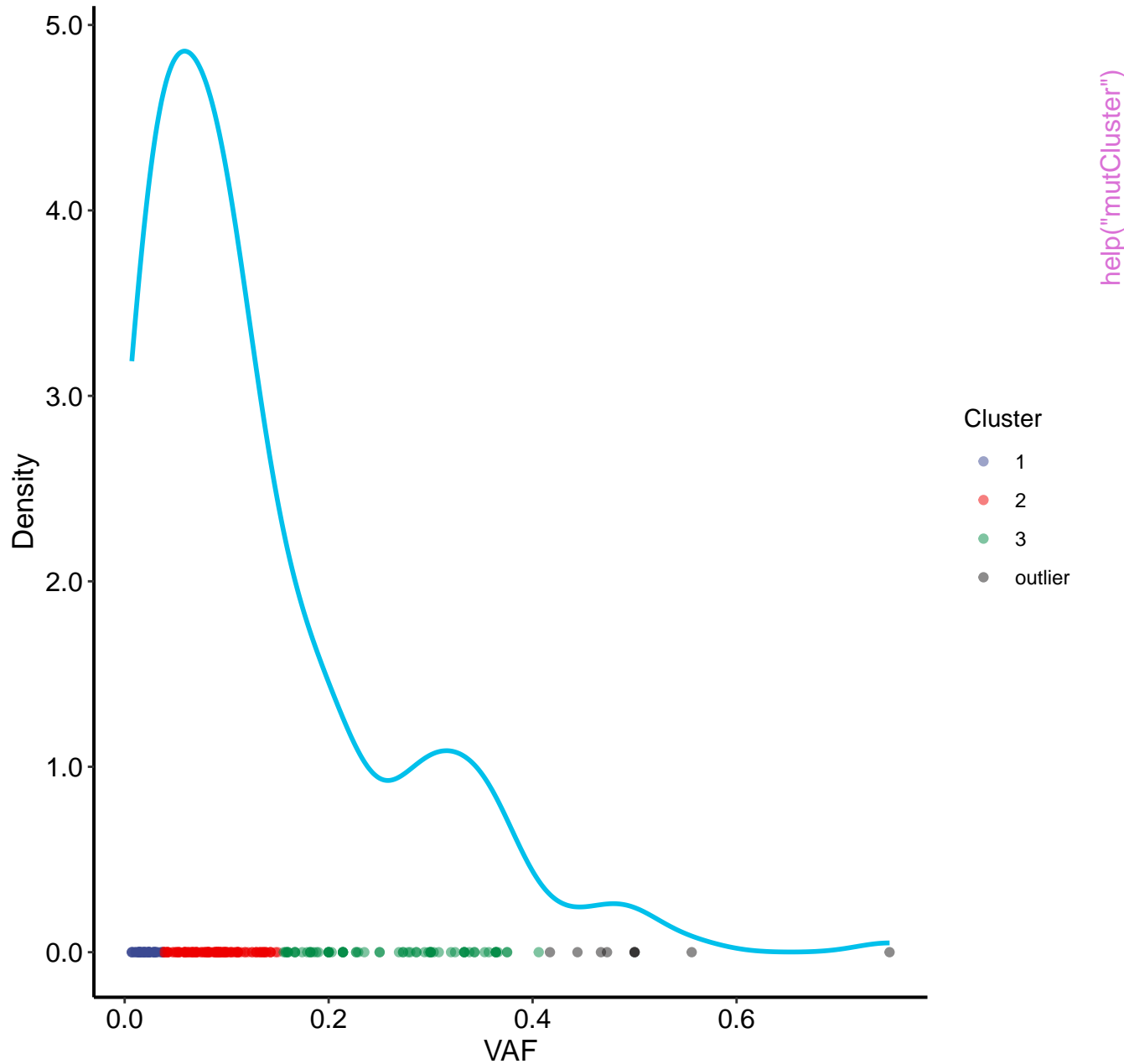




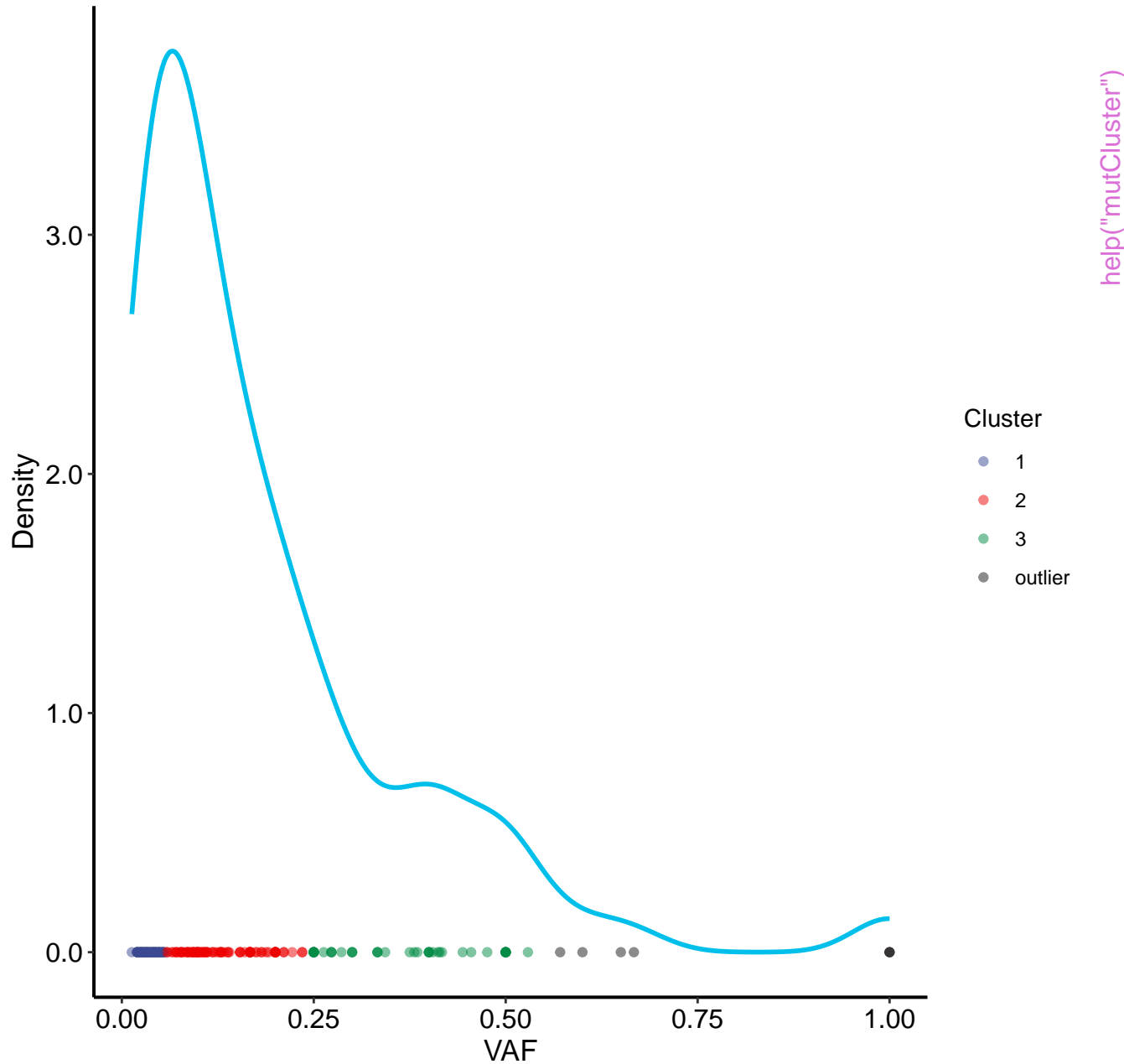
V402: V402_P_1



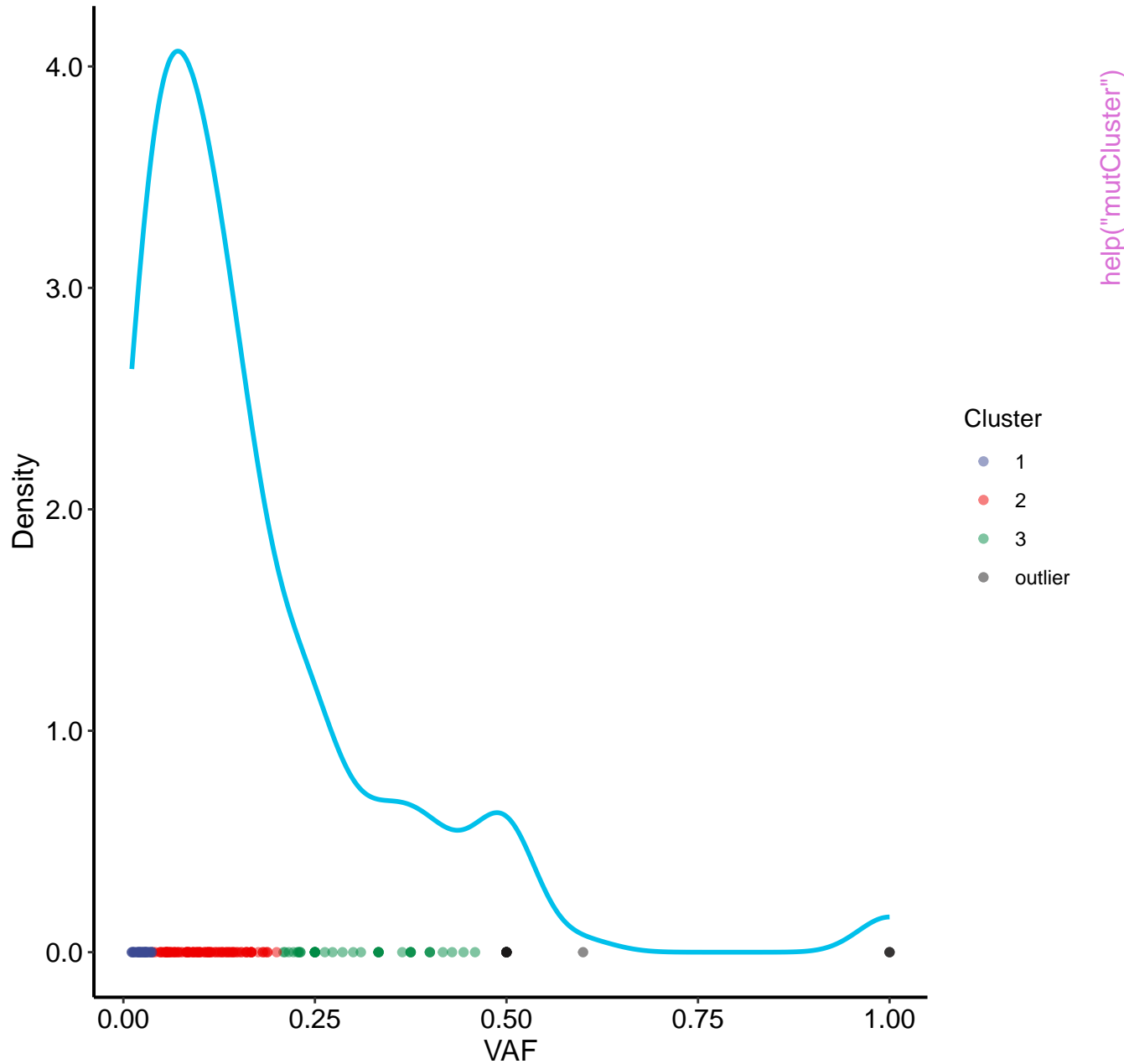
V402: V402_P_2



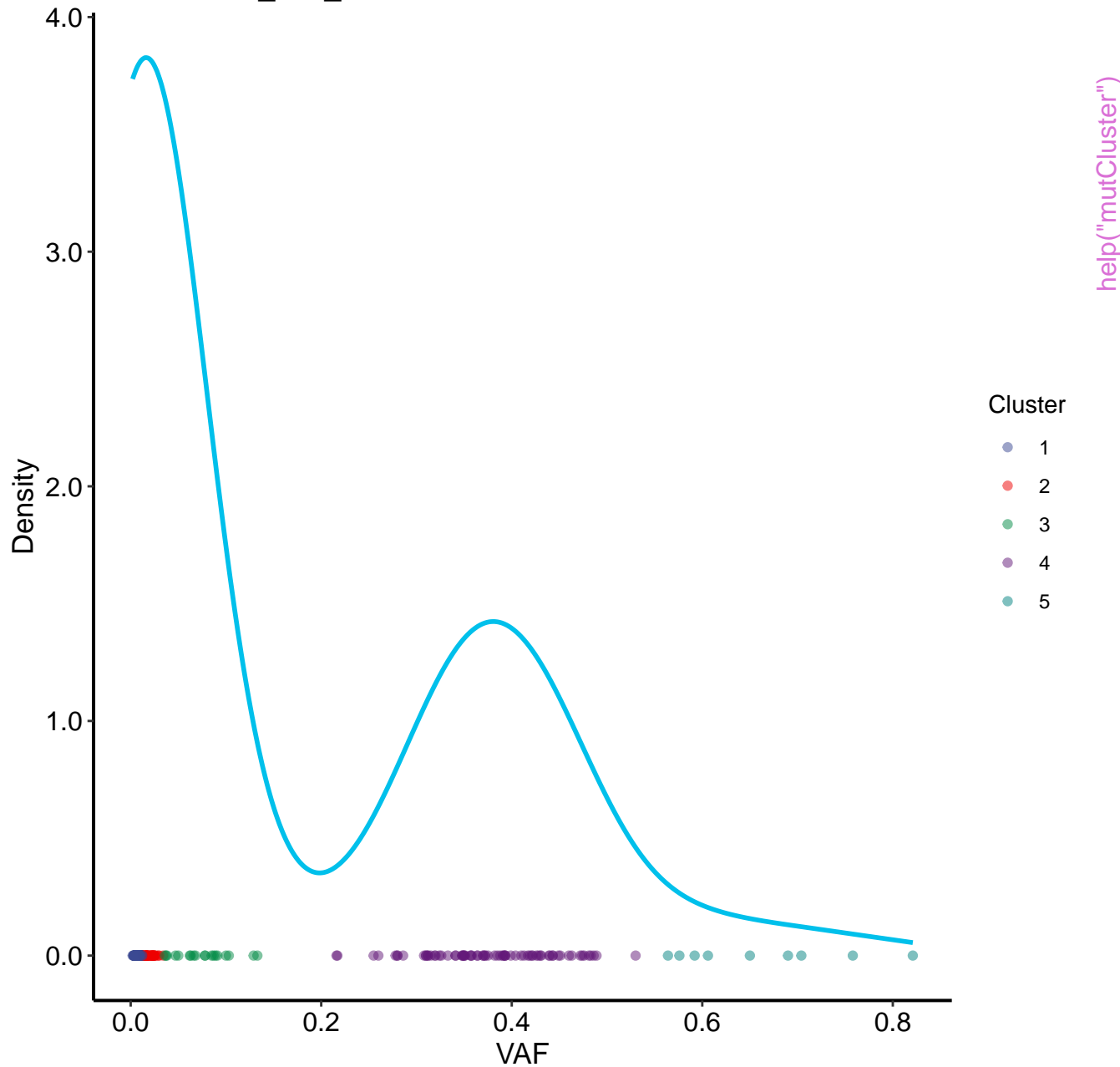
V402: V402_P_3



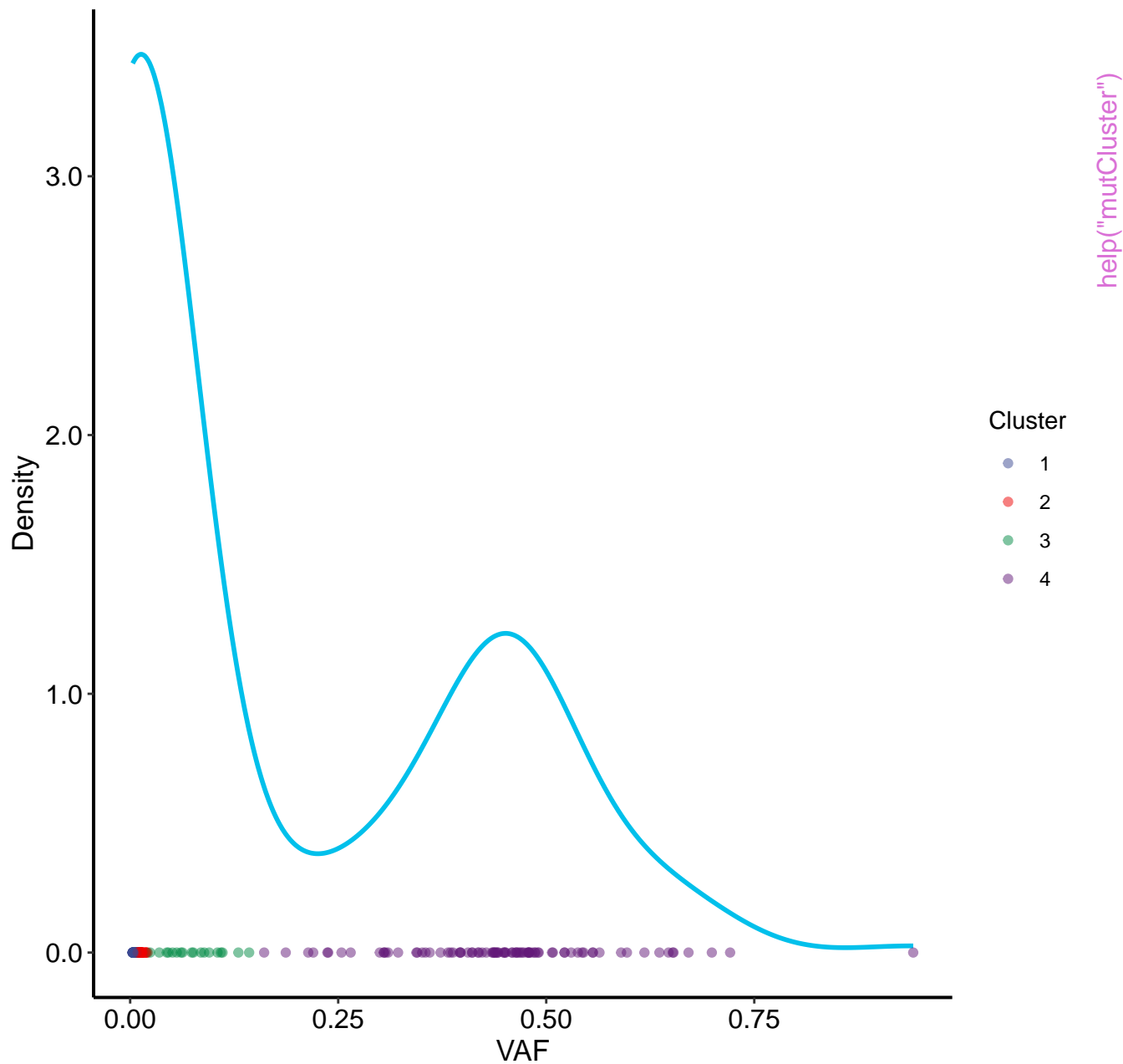
V402: V402_P_4



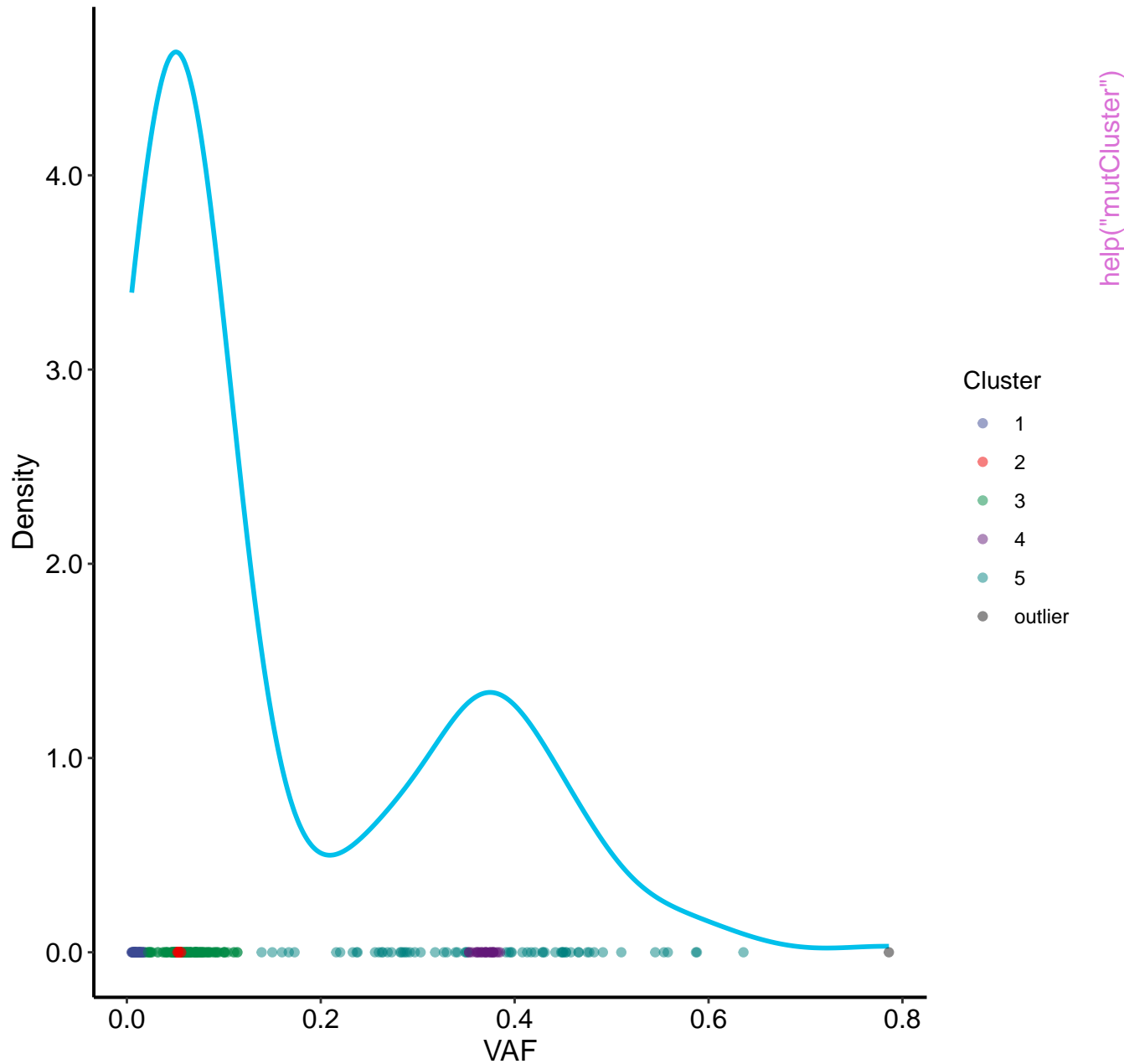
V402: V402_BM_1



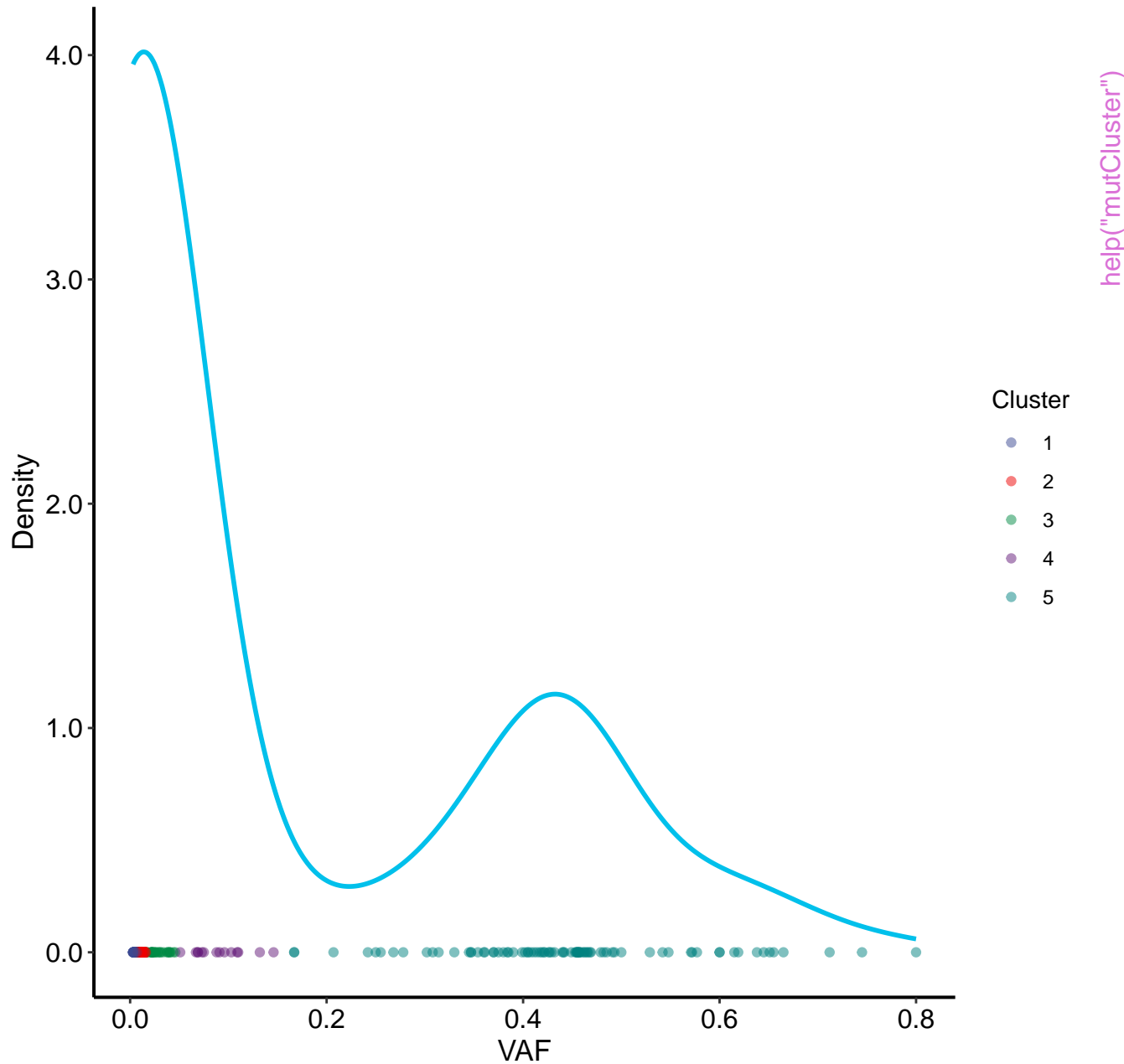
V402: V402_BM_2



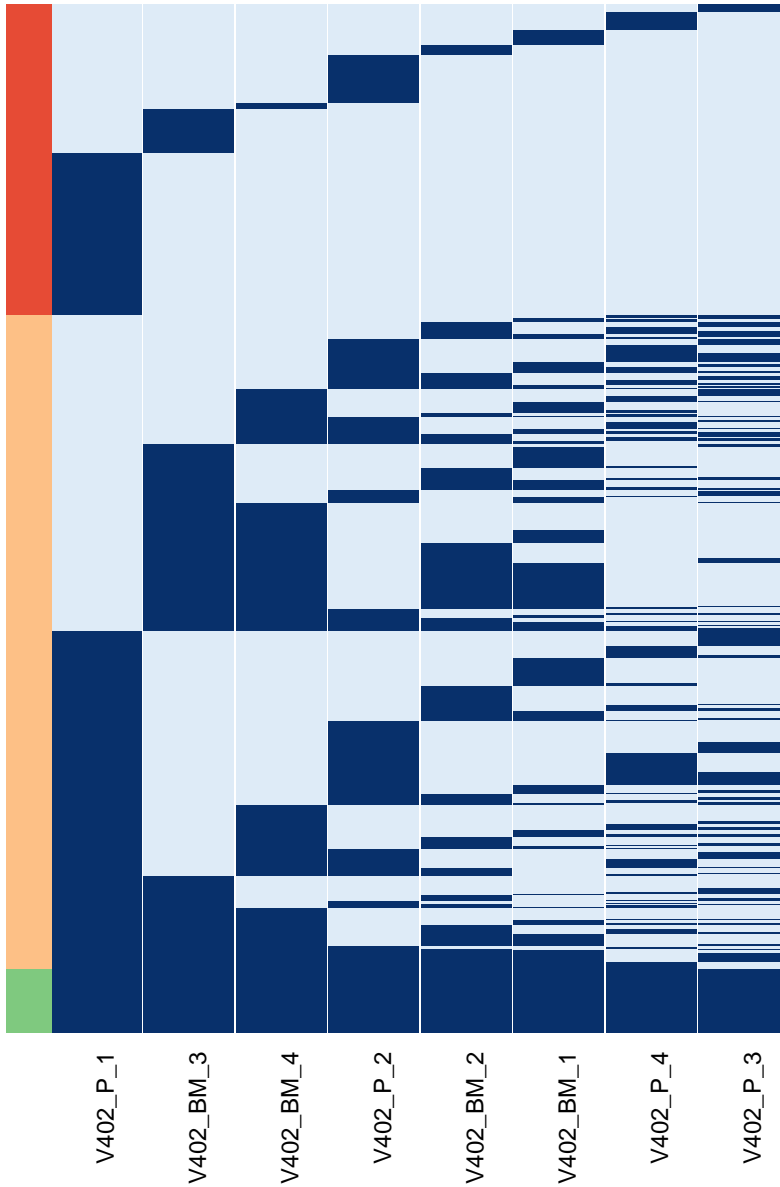
V402: V402_BM_3



V402: V402_BM_4

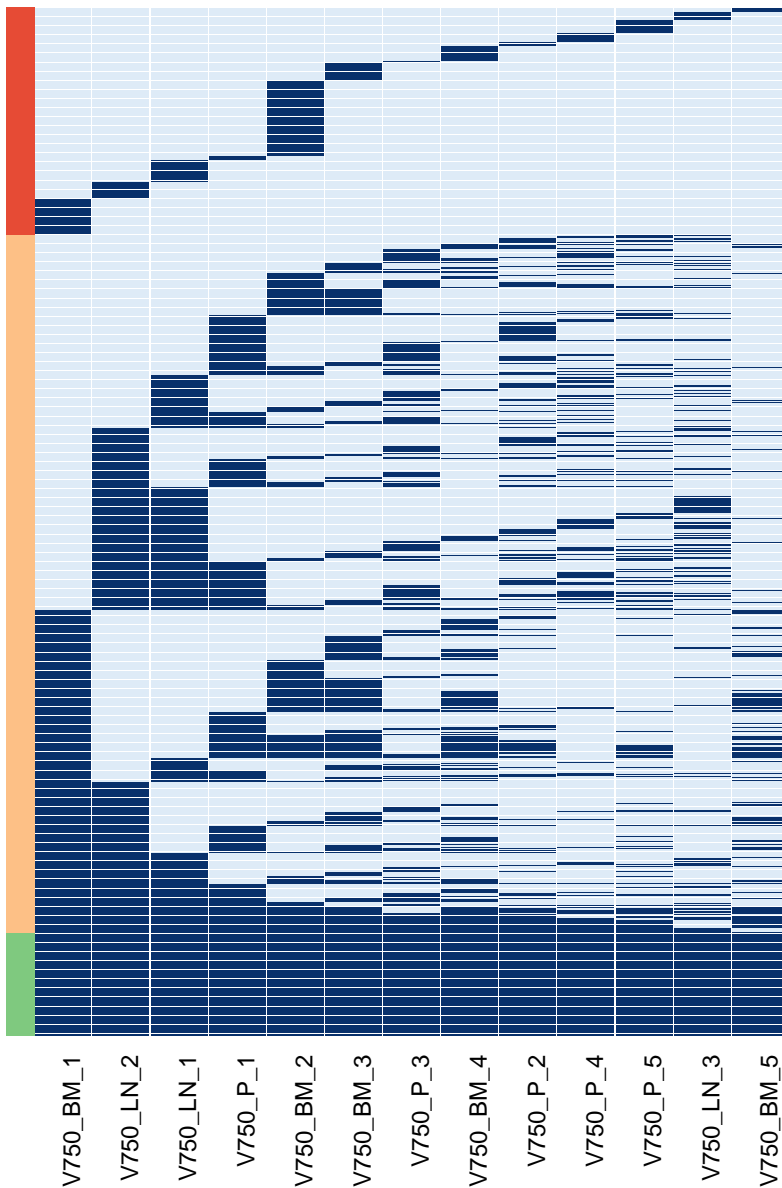


V402 (n=693)



help("mutHeatmap")

V750 (n=915)



Mutation

0

1

Type

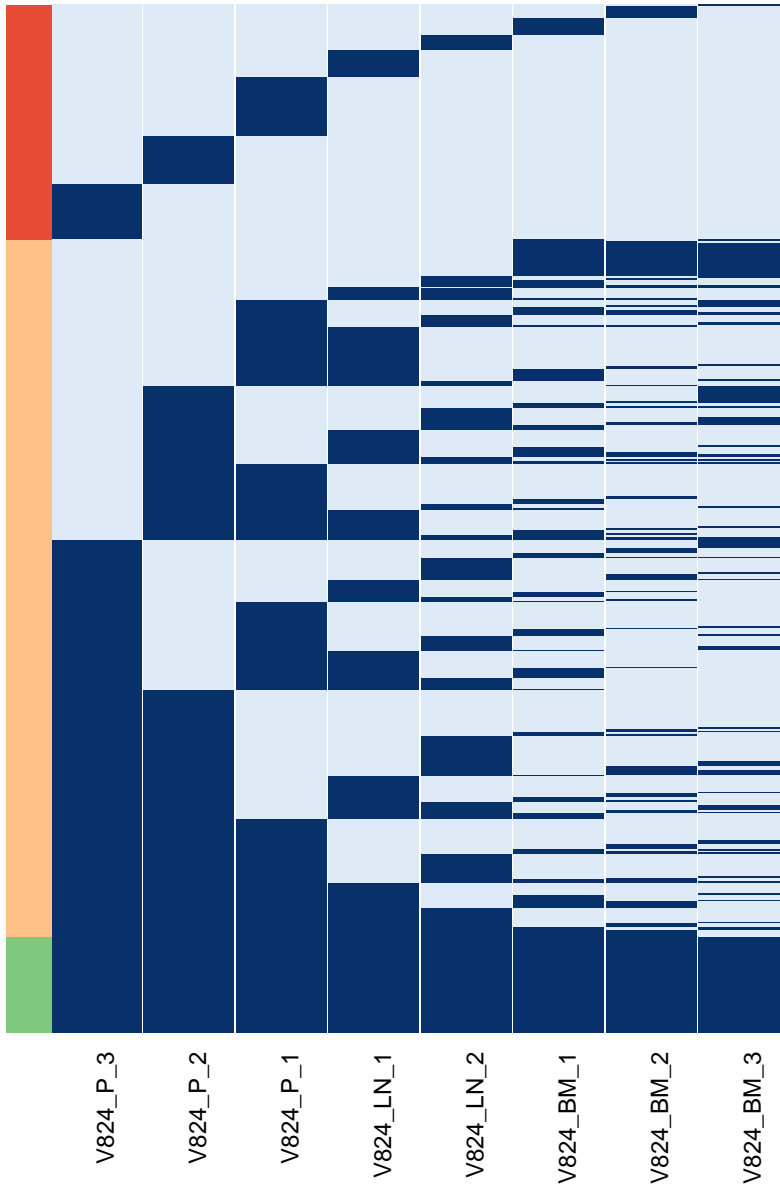
Public (10.1%)

Shared (67.9%)

Private (22.1%)

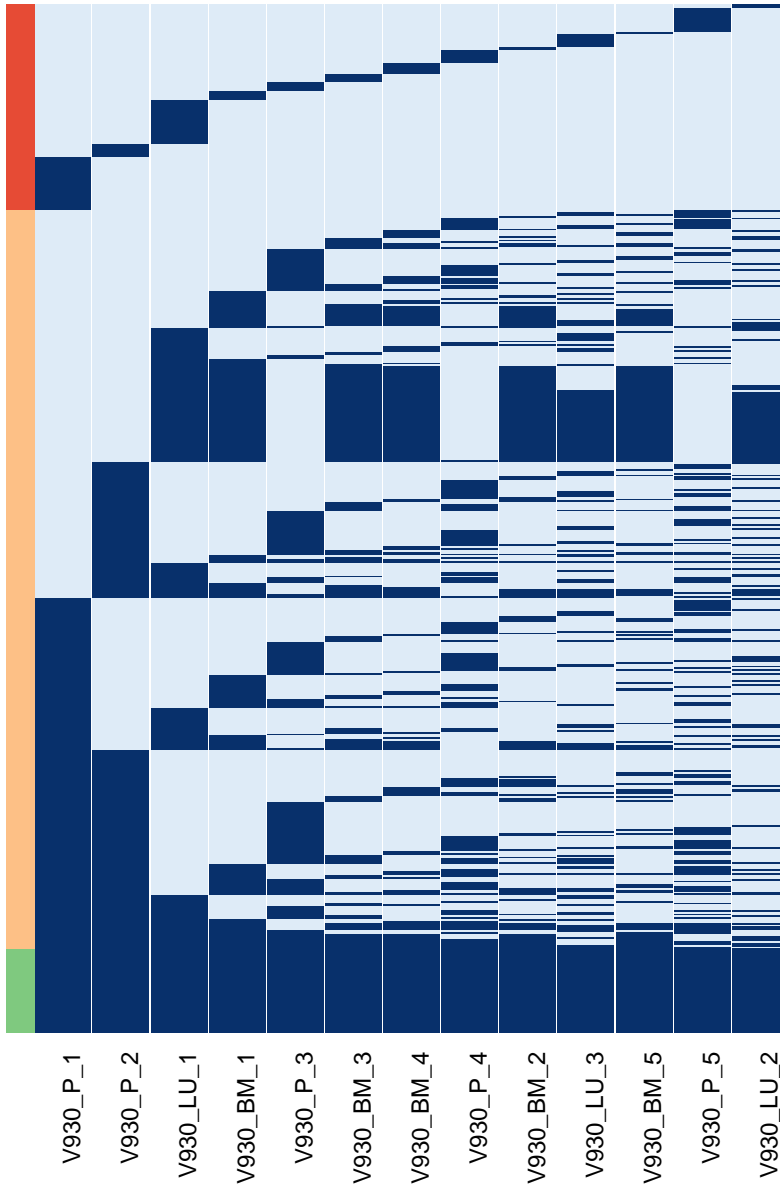
```
help("mutHeatmap")
```

V824 (n=609)



help("mutHeatmap")

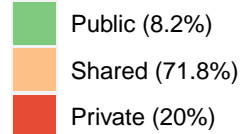
V930 (n=560)



Mutation

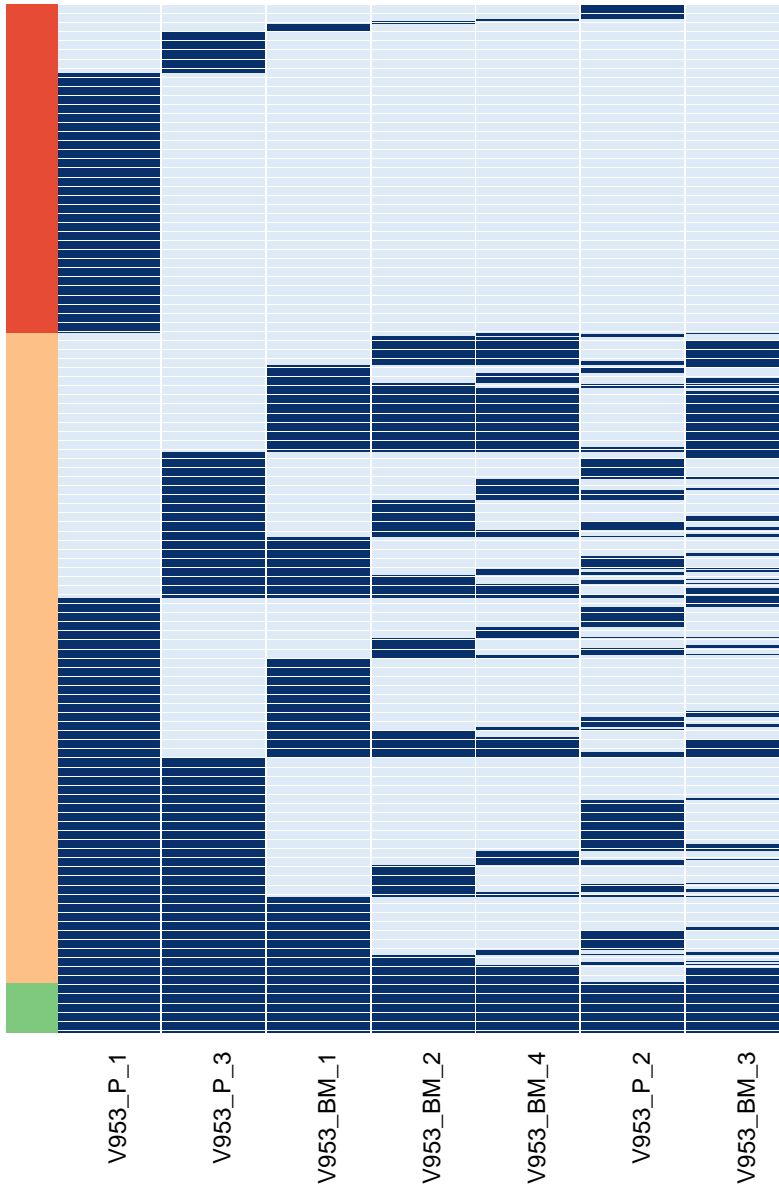


Type



help("mutHeatmap")

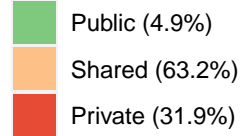
V953 (n=915)



Mutation

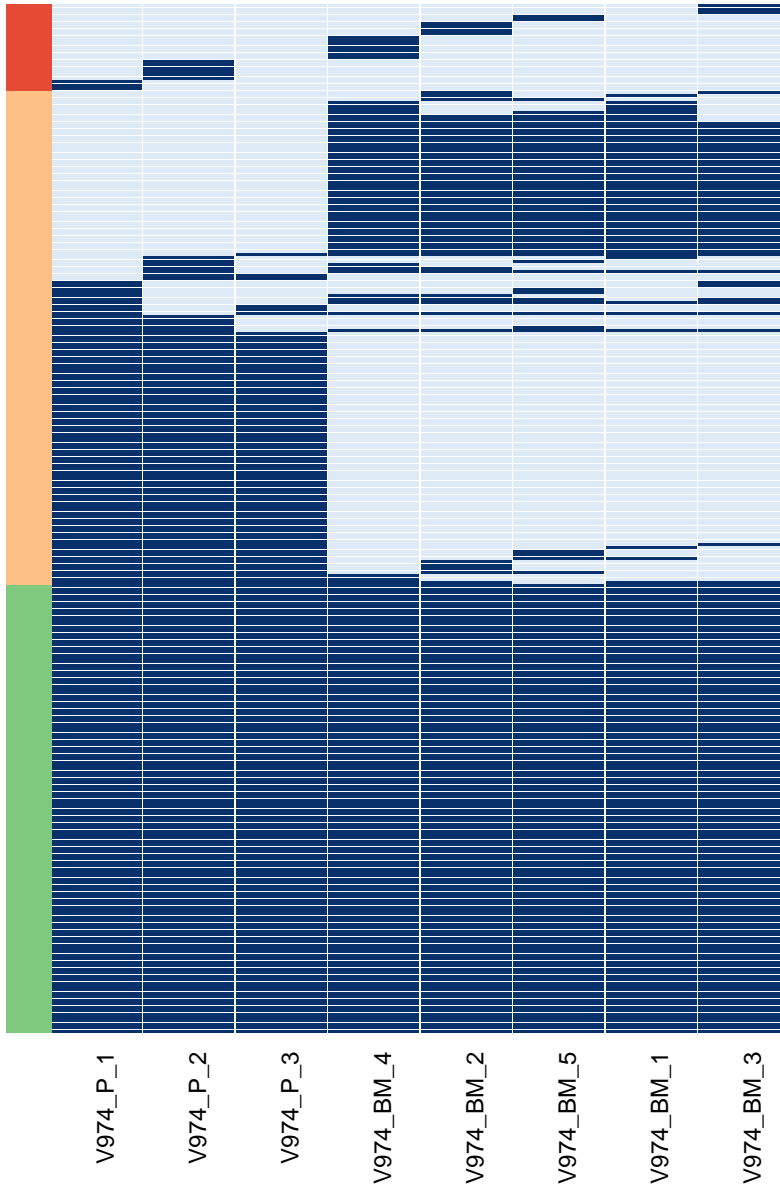


Type



help("mutHeatmap")

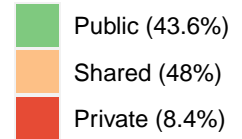
V974 (n=298)



Mutation



Type



help("mutHeatmap")

V402

43

267

Proportion

1.00
0.75
0.50
0.25
0.00

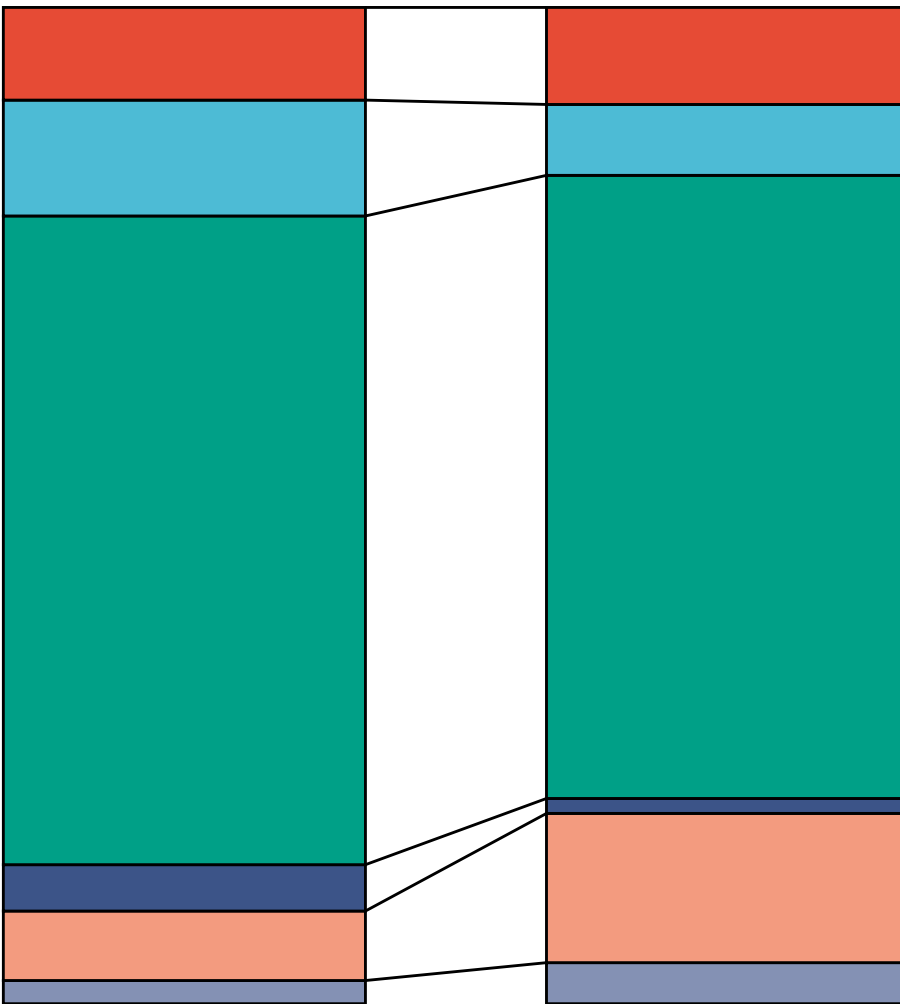
Trunk

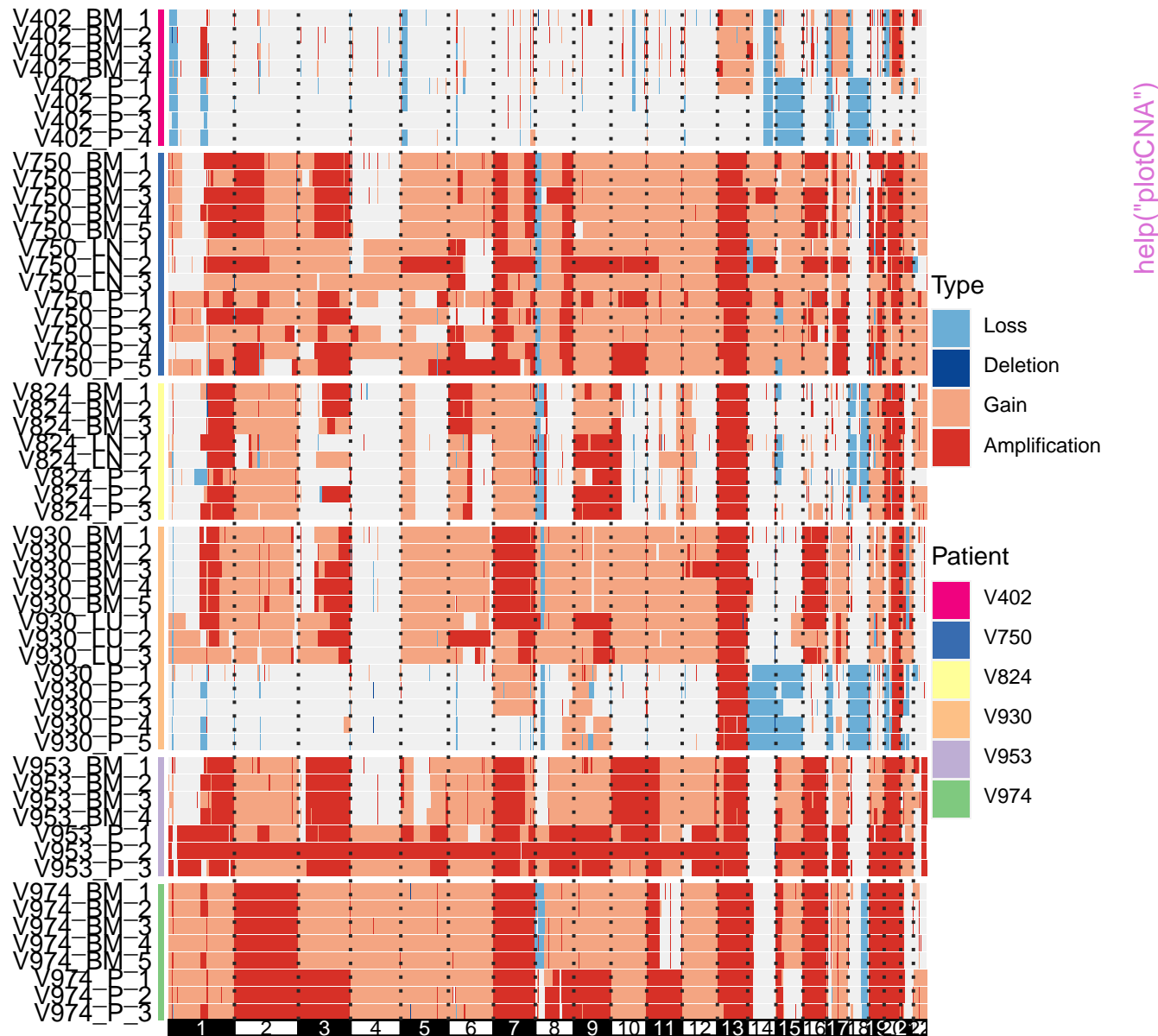
Branches

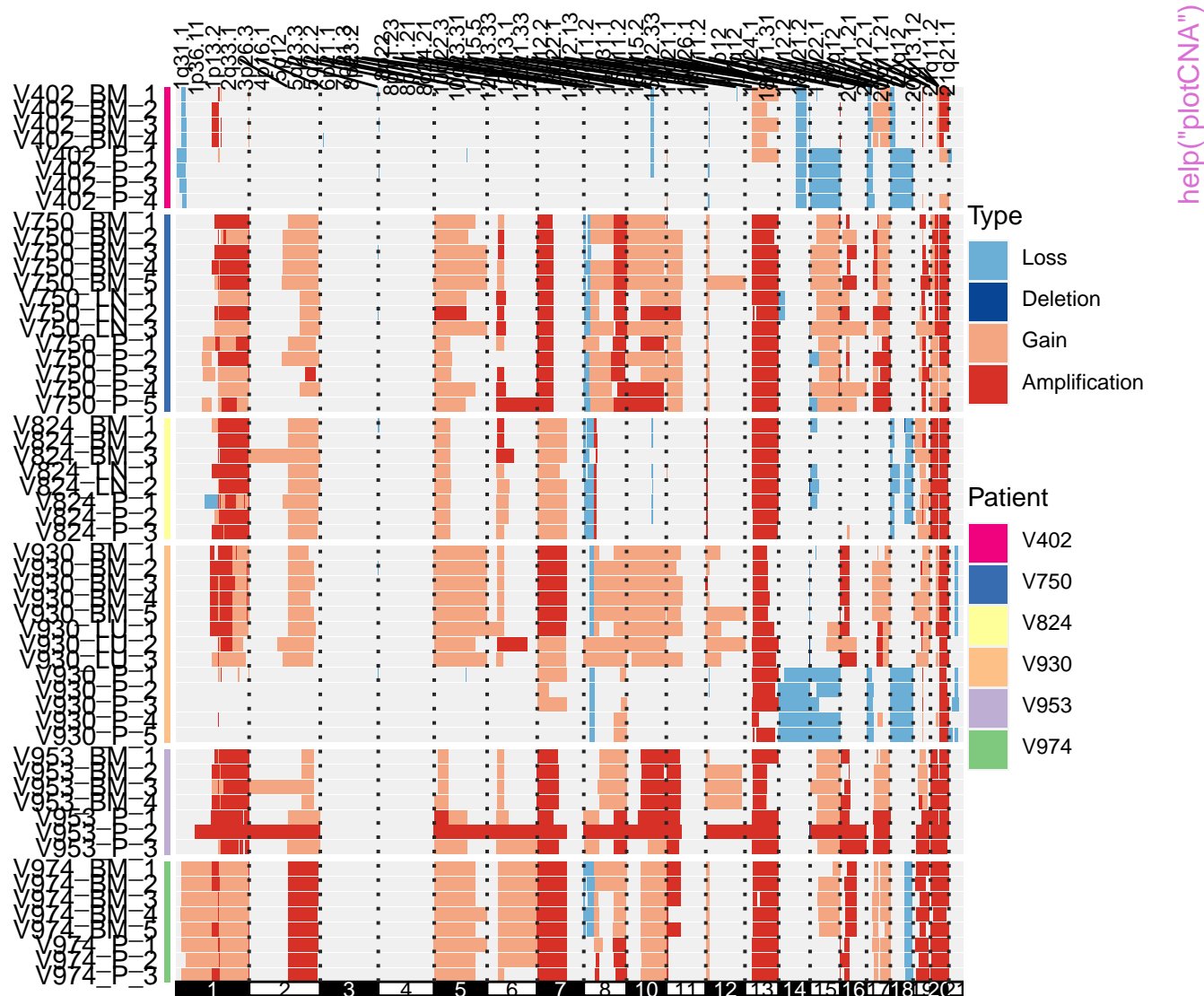
Group

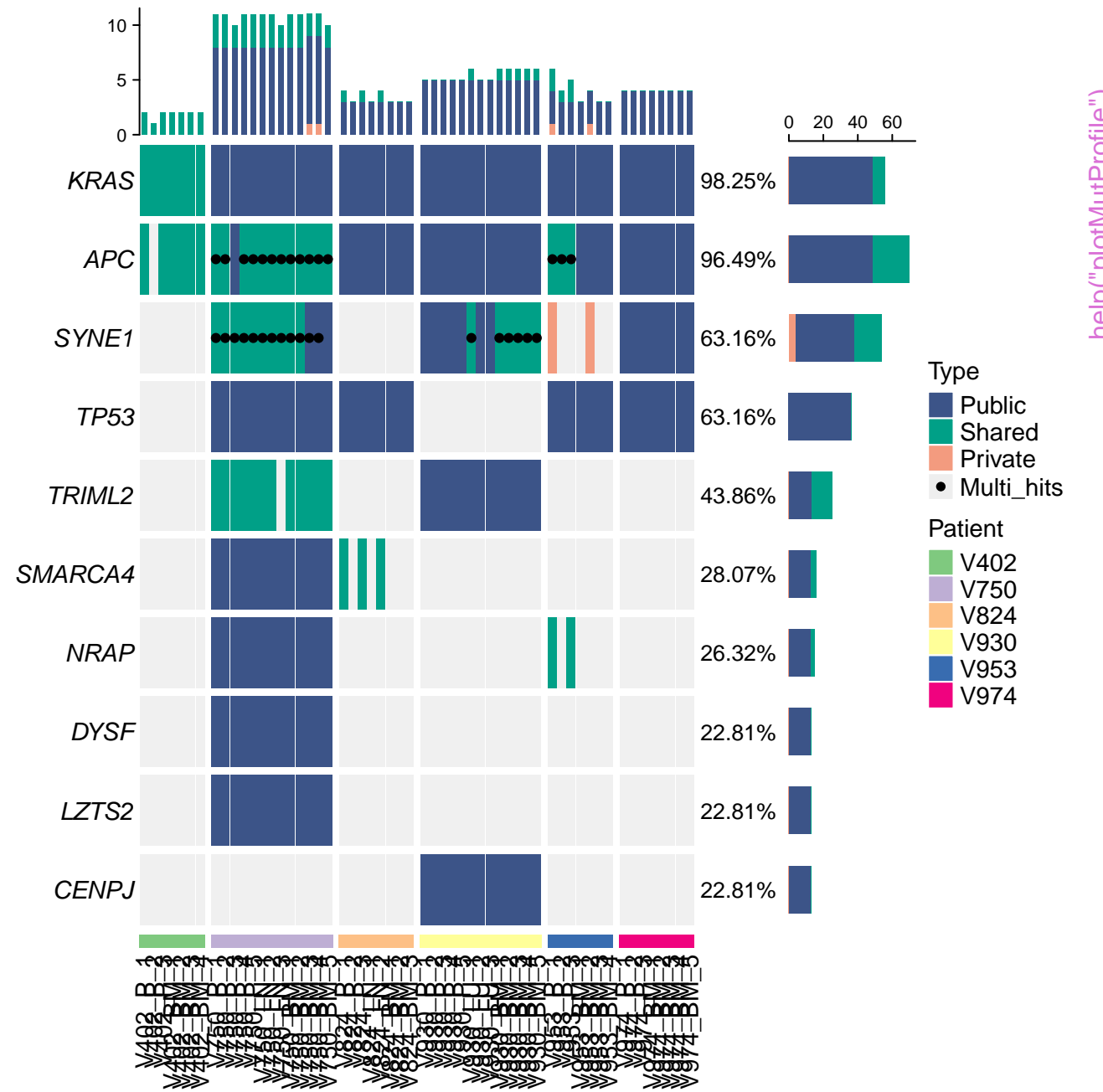


help("mutTrunkBranch")





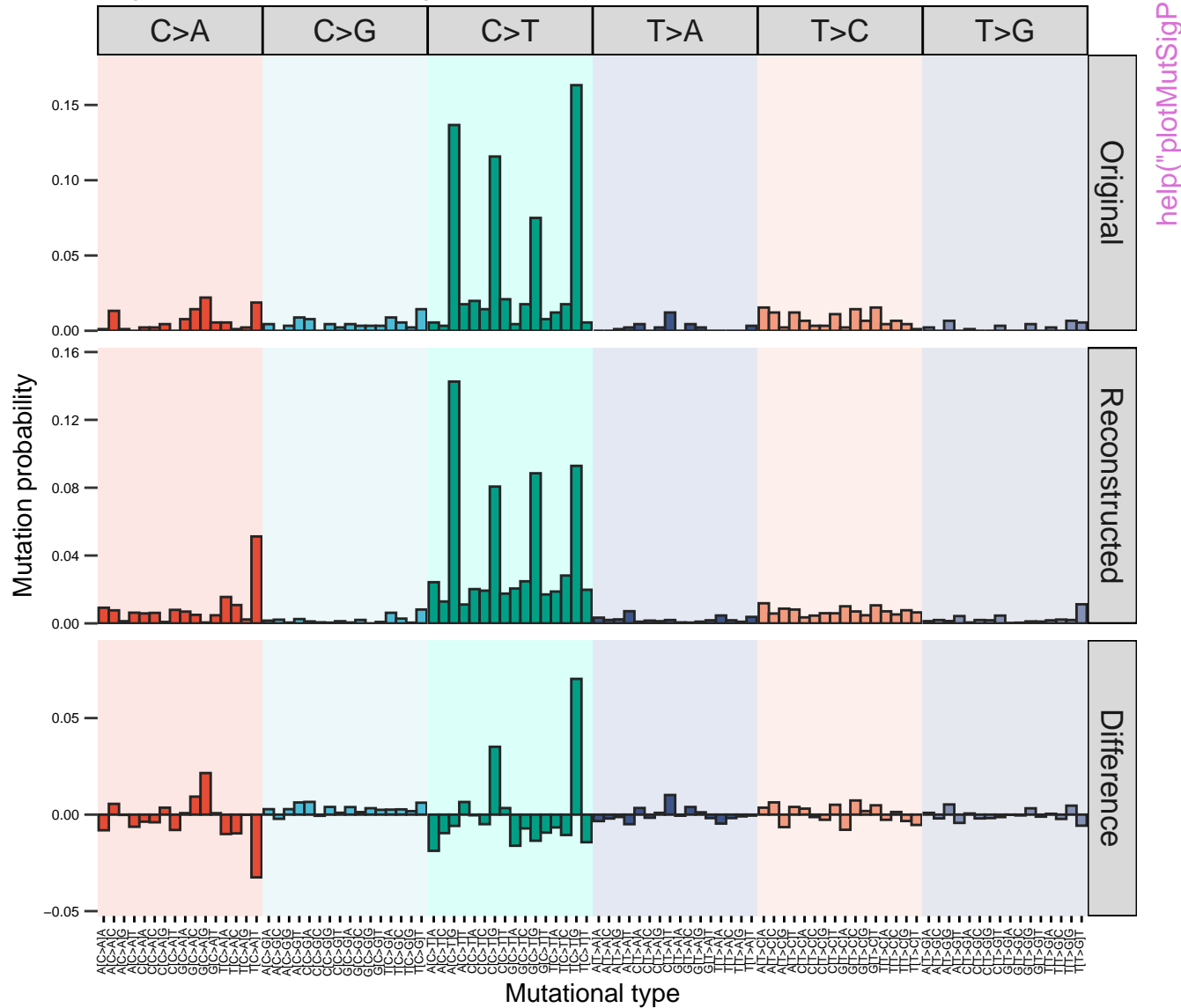




V402: BM

RSS = 0.011; Cosine similarity = 0.923

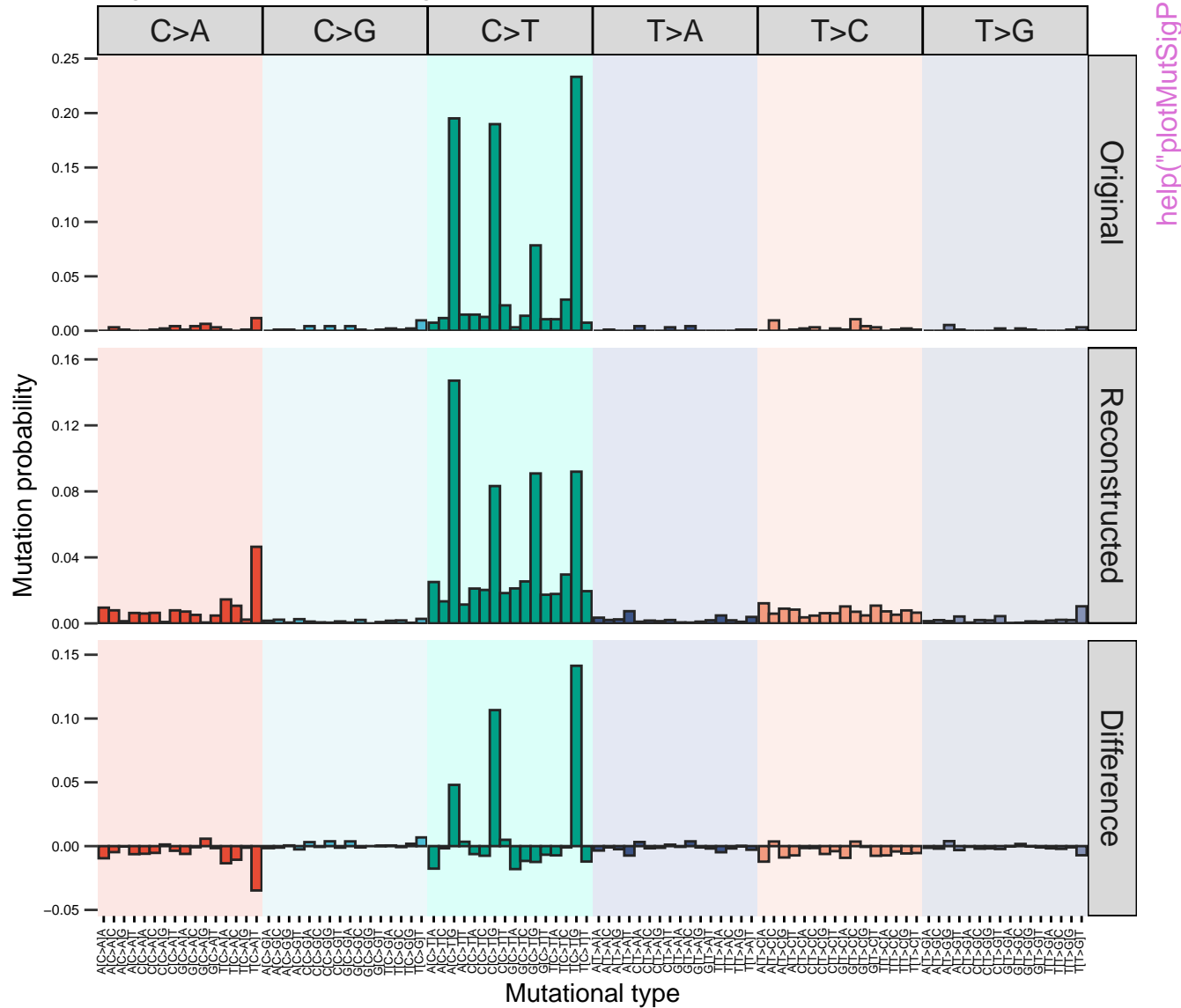
Signature 1: 0.82 & Signature 10: 0.12



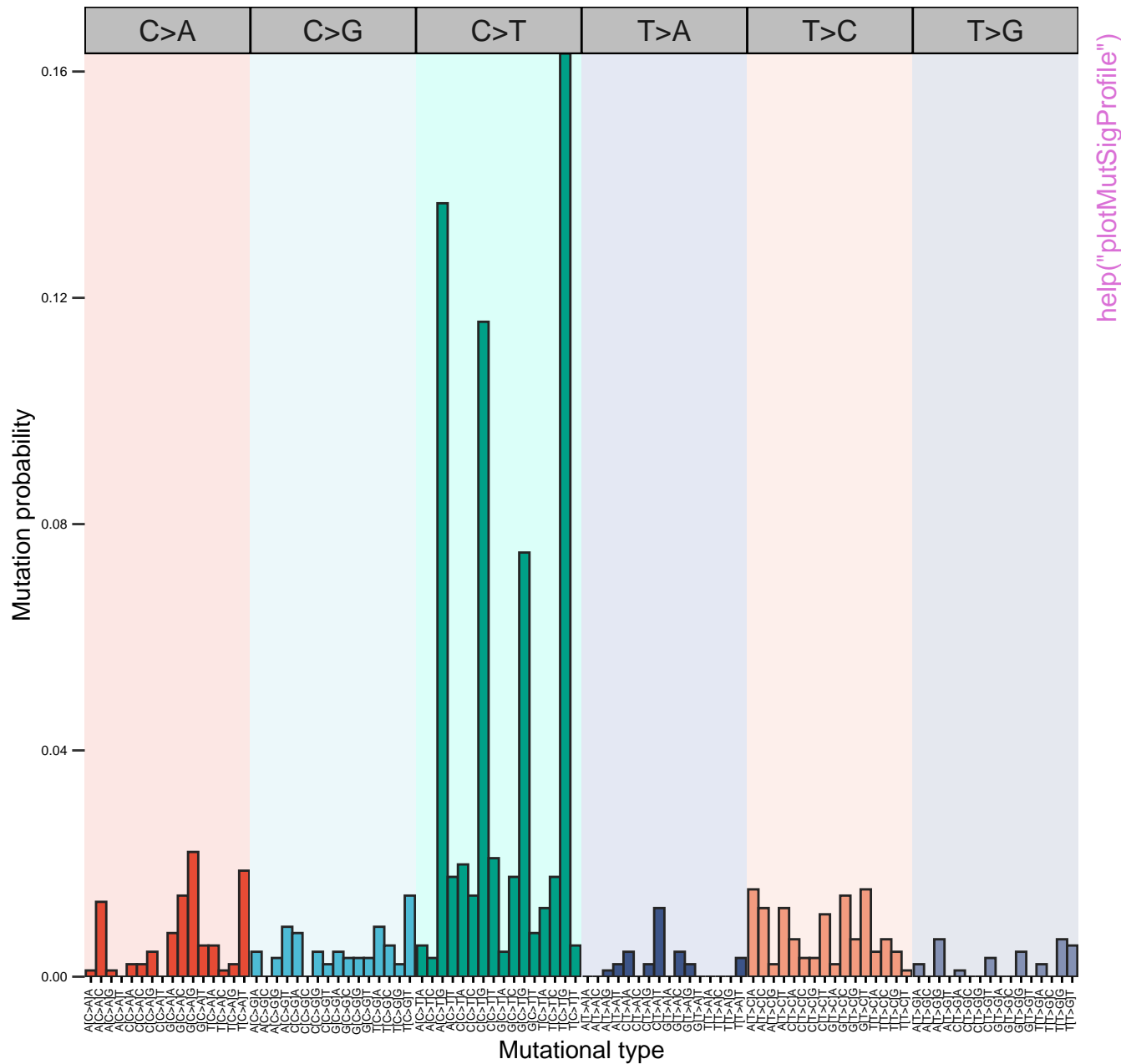
V402: P

RSS = 0.038; Cosine similarity = 0.892

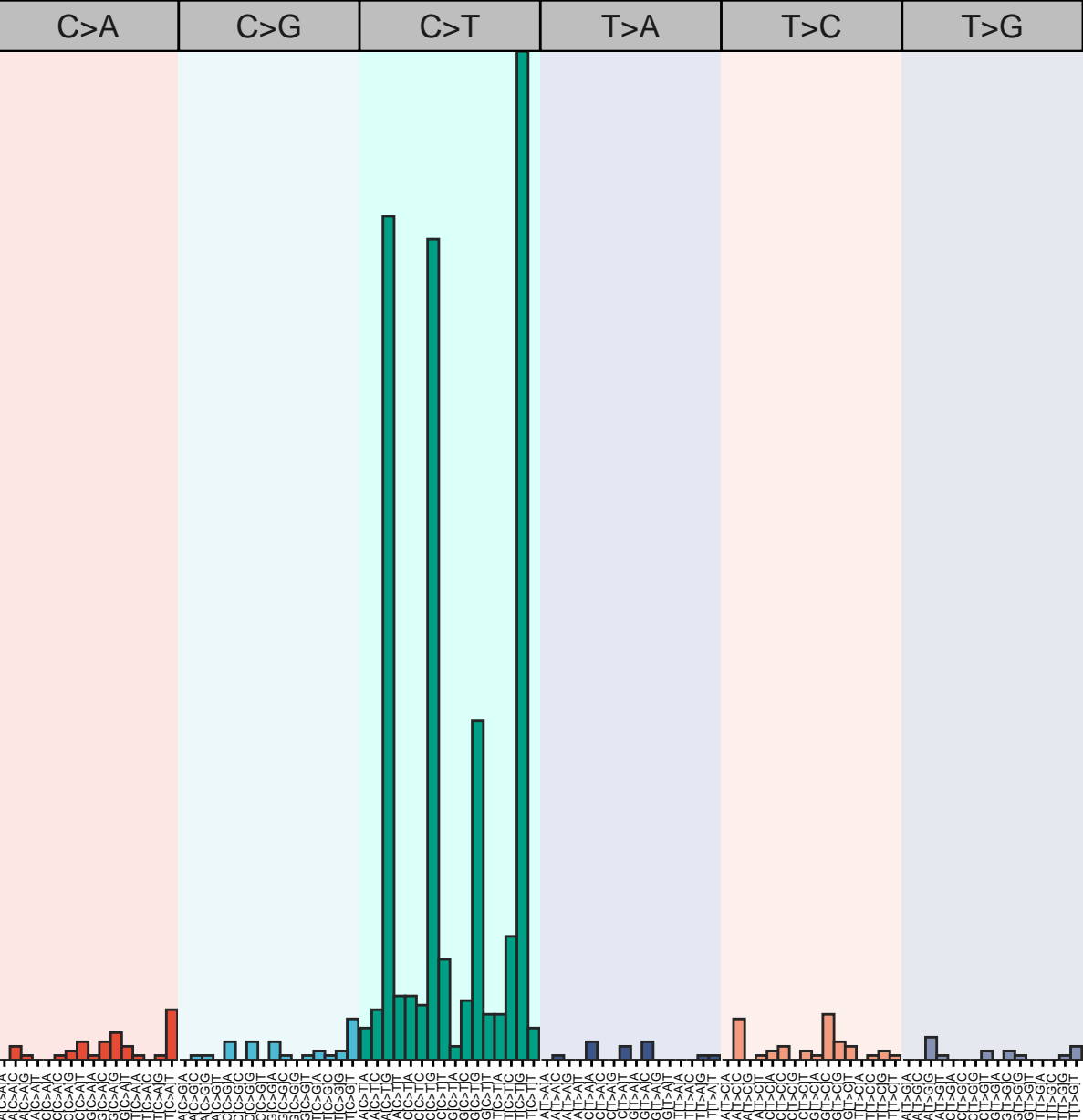
Signature 1: 0.85 & Signature 10: 0.11



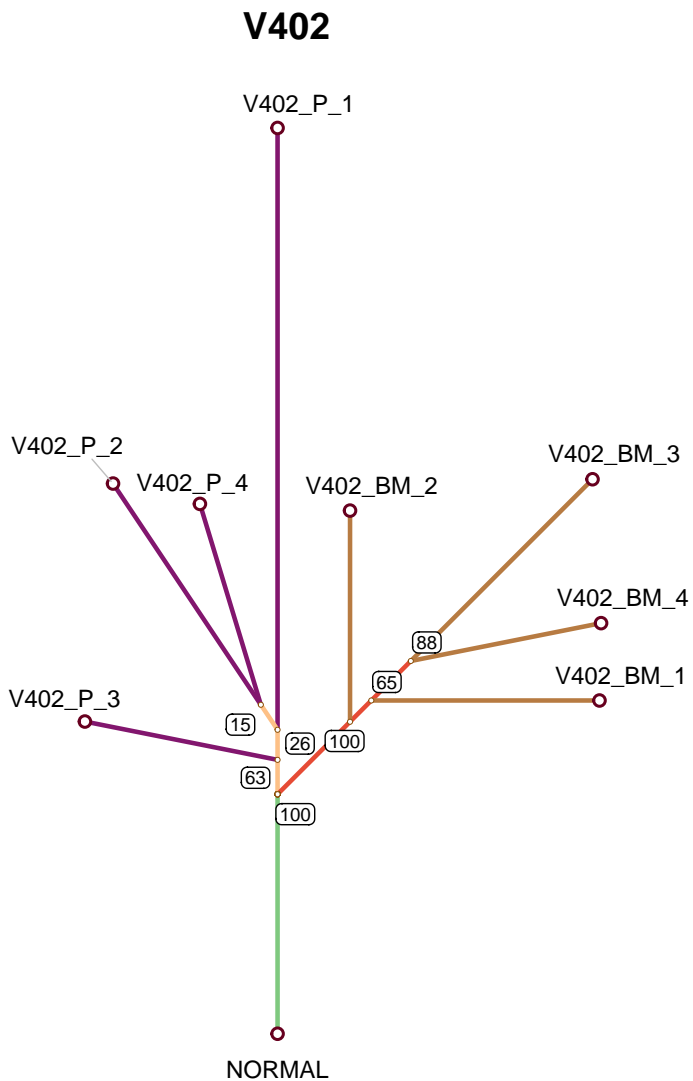
V402: BM



V402: P

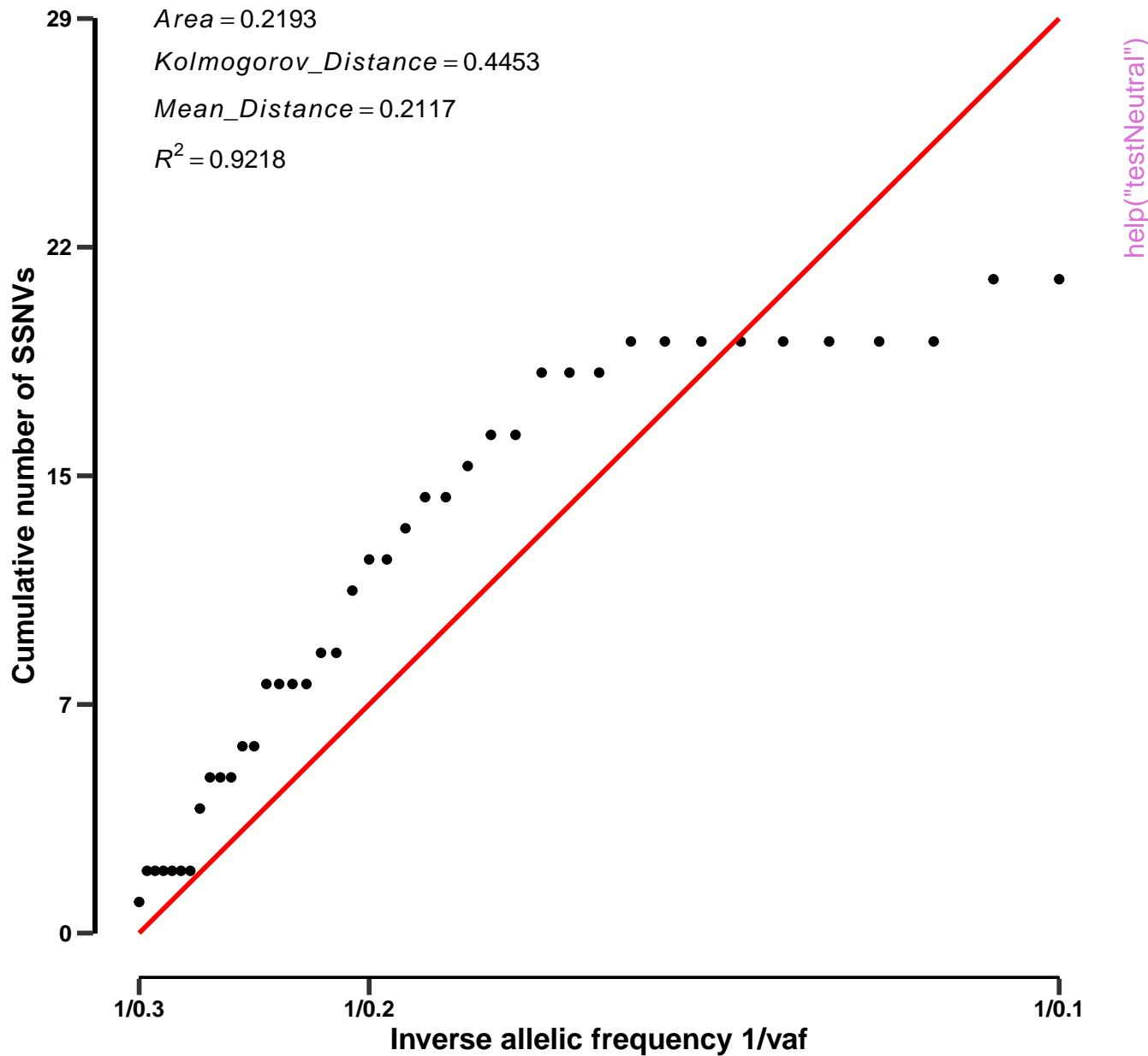


help("plotMutSigProfile")

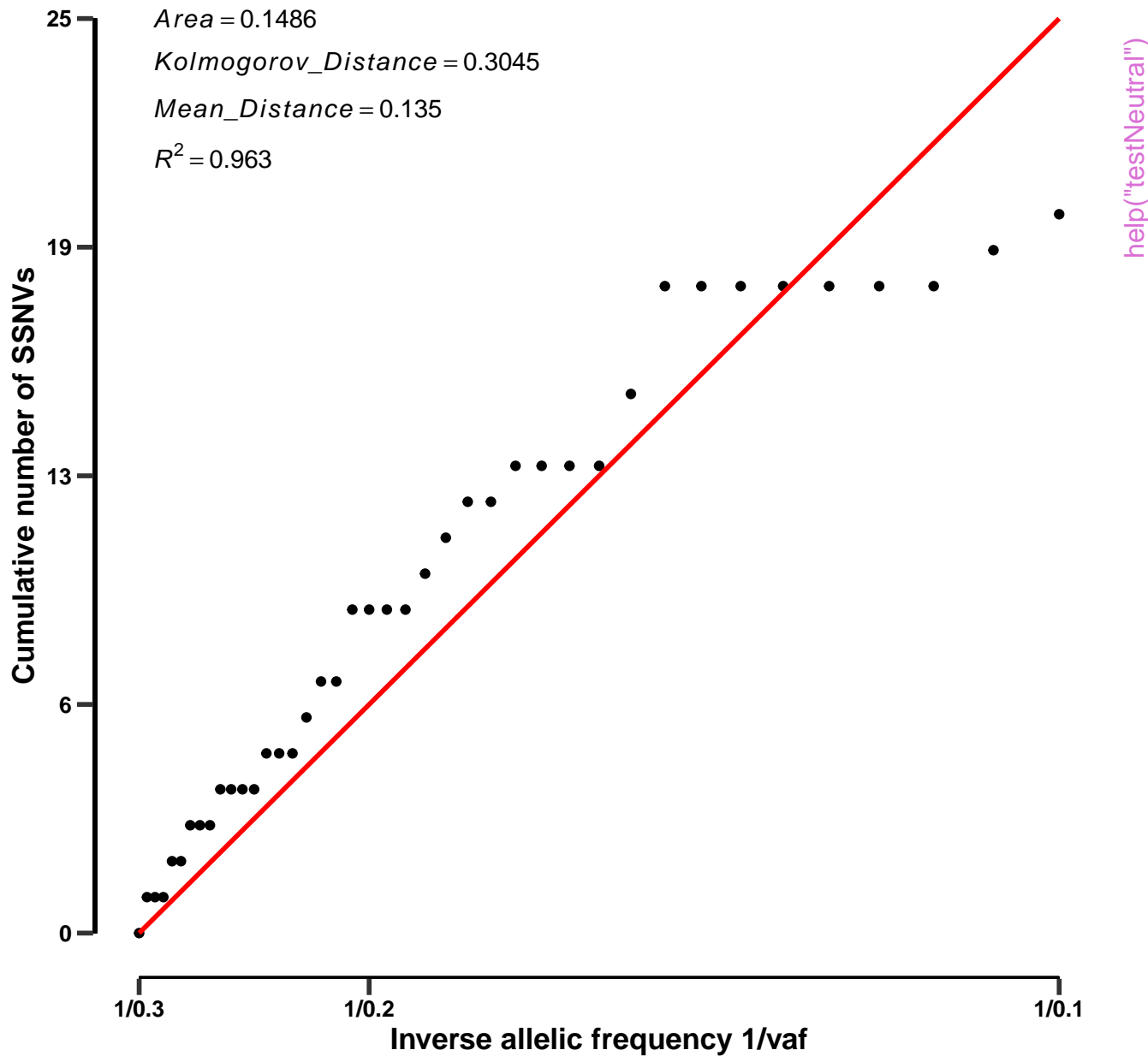


help("plotPhyloTree")

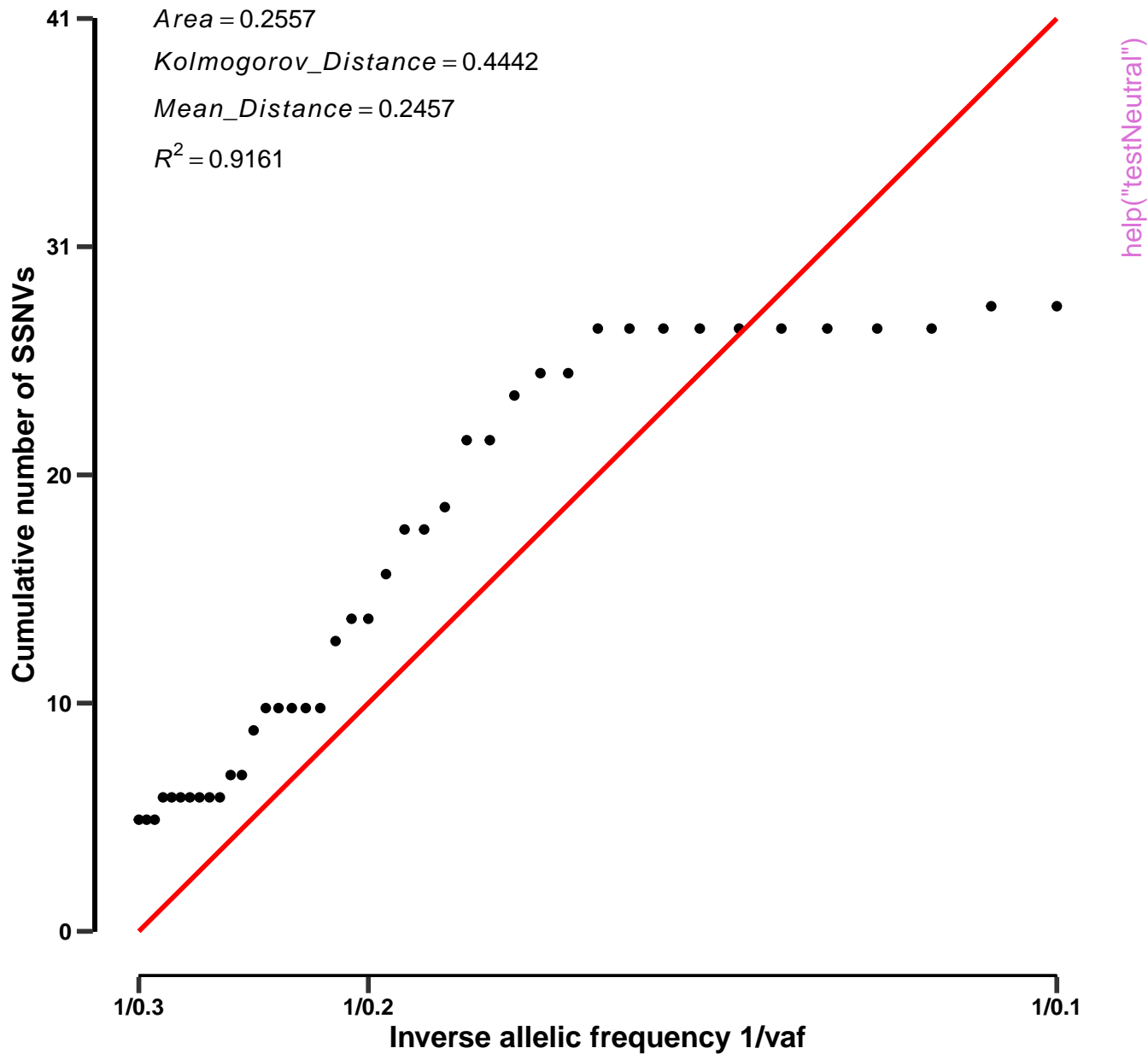
V402:V402_P_2



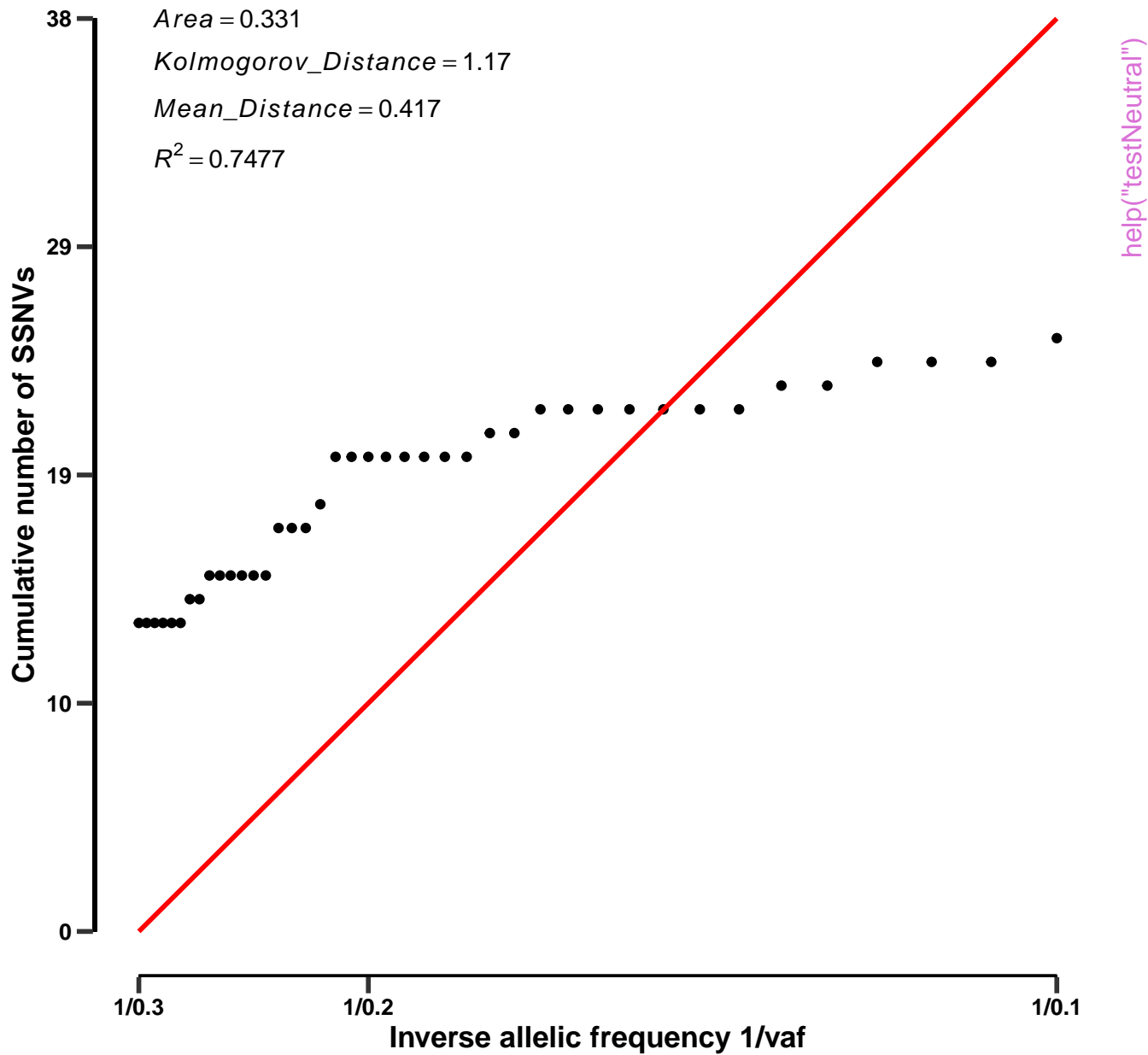
V402:V402_P_3



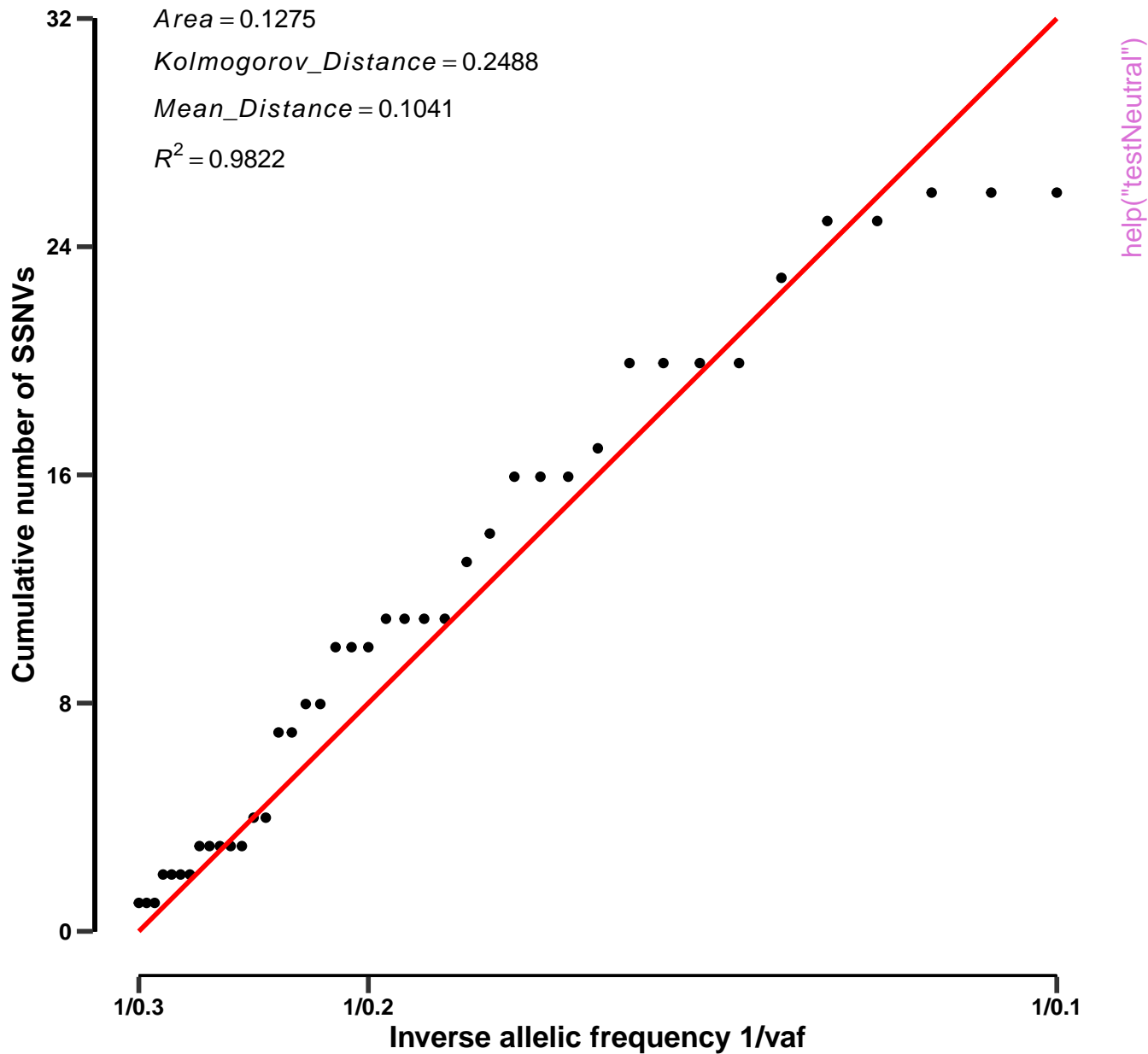
V402:V402_P_4



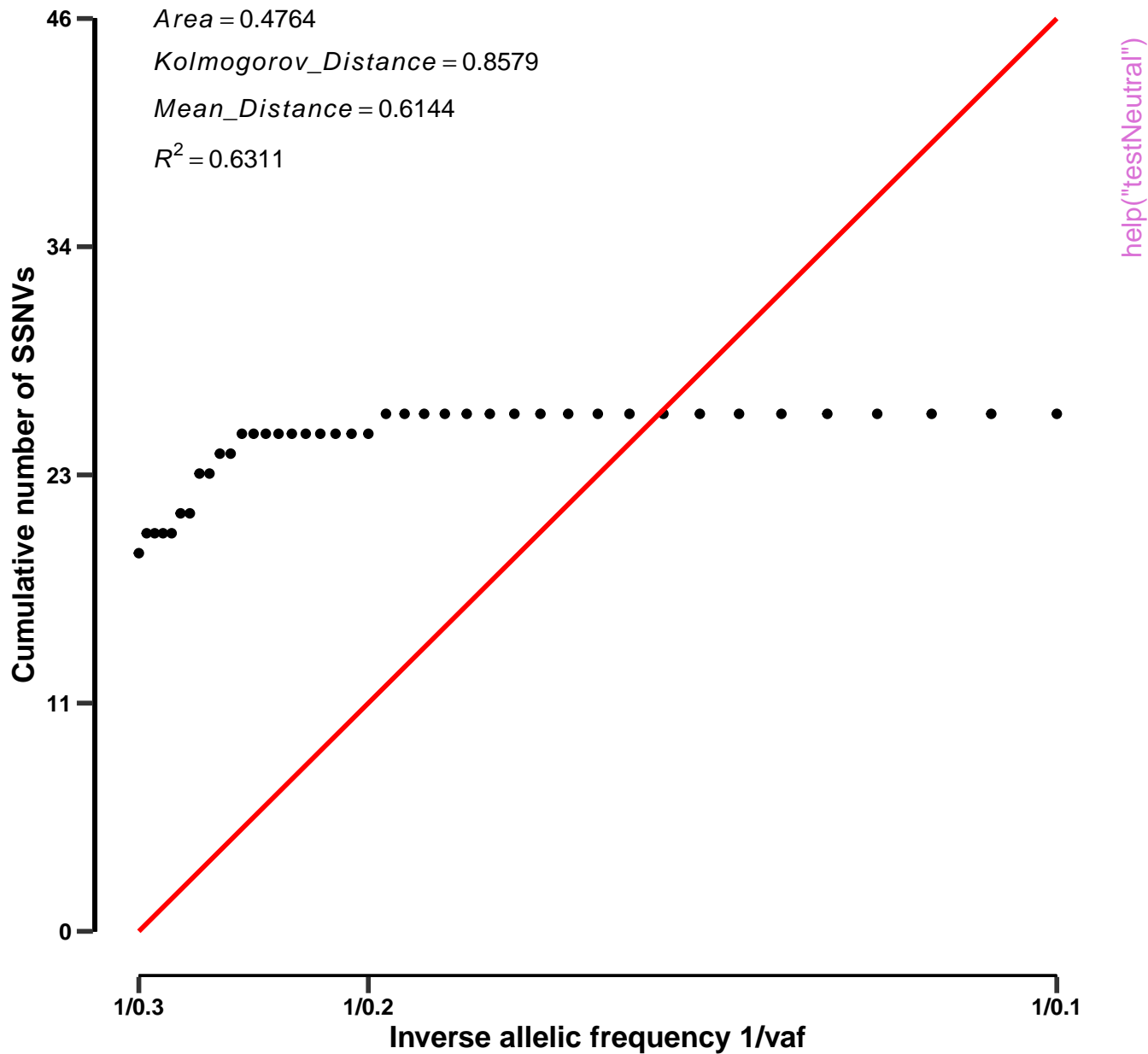
V750:V750_BM_2



V930:V930_P_4



V953:V953_P_1



V953:V953_BM_4

