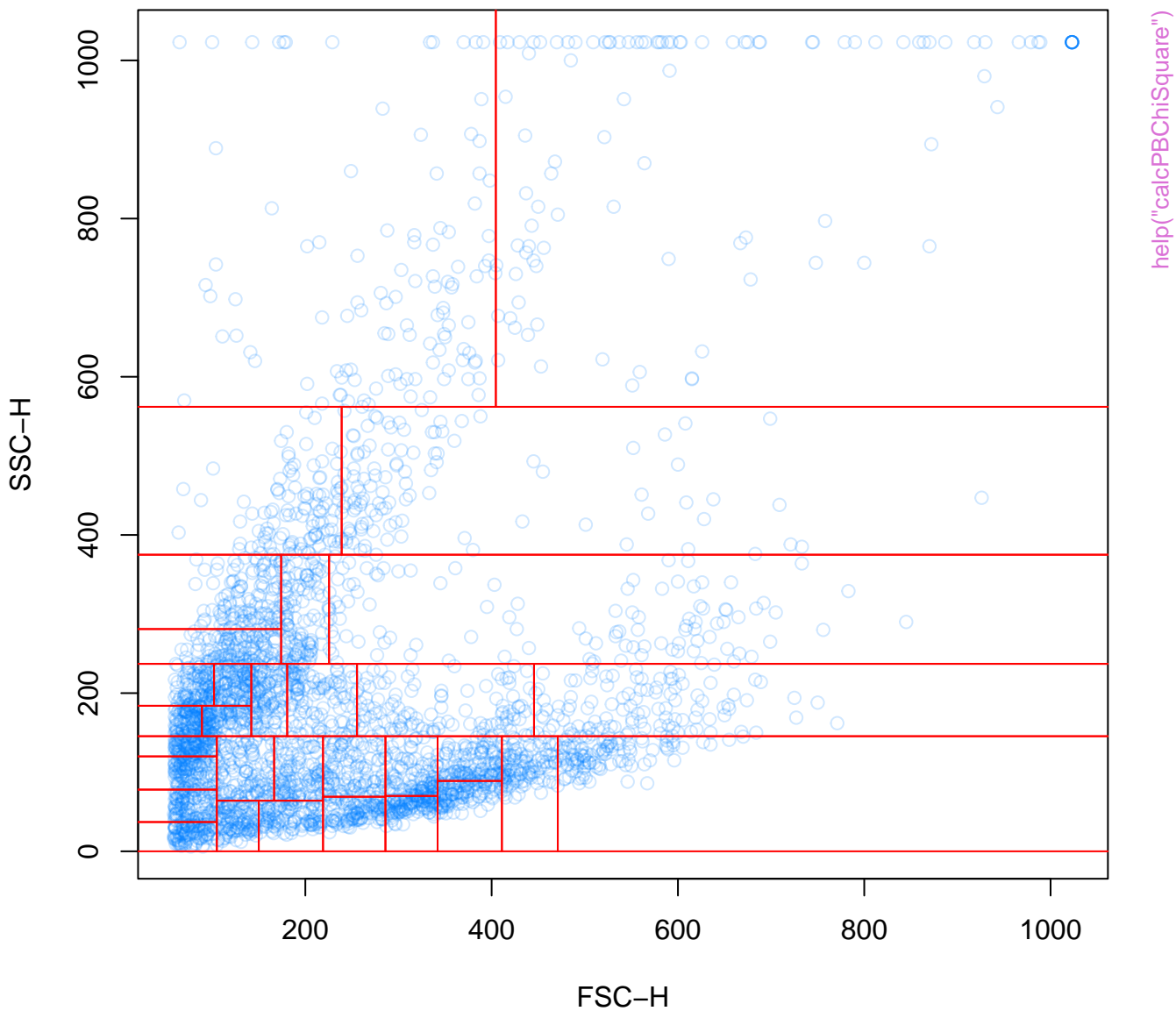
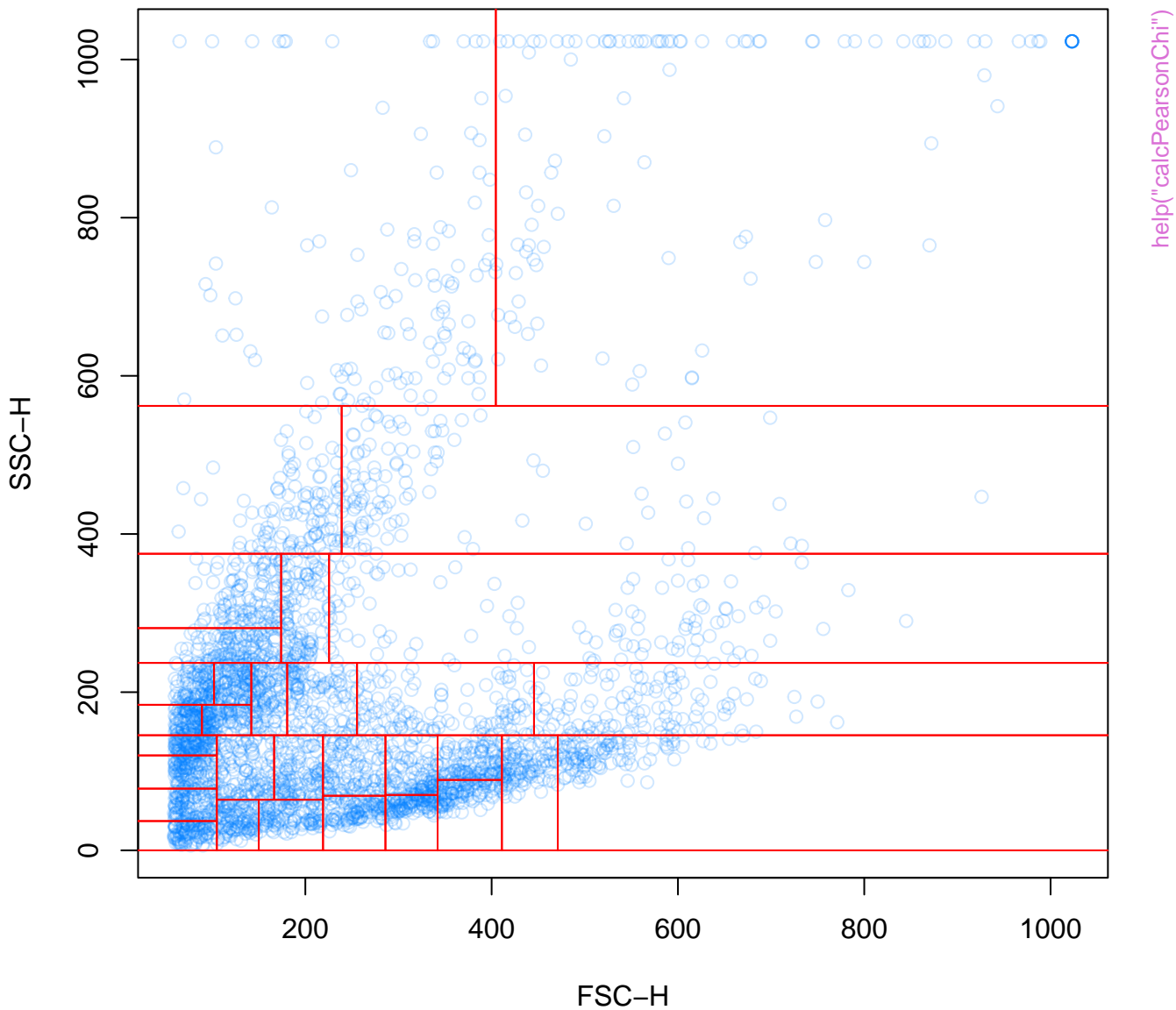


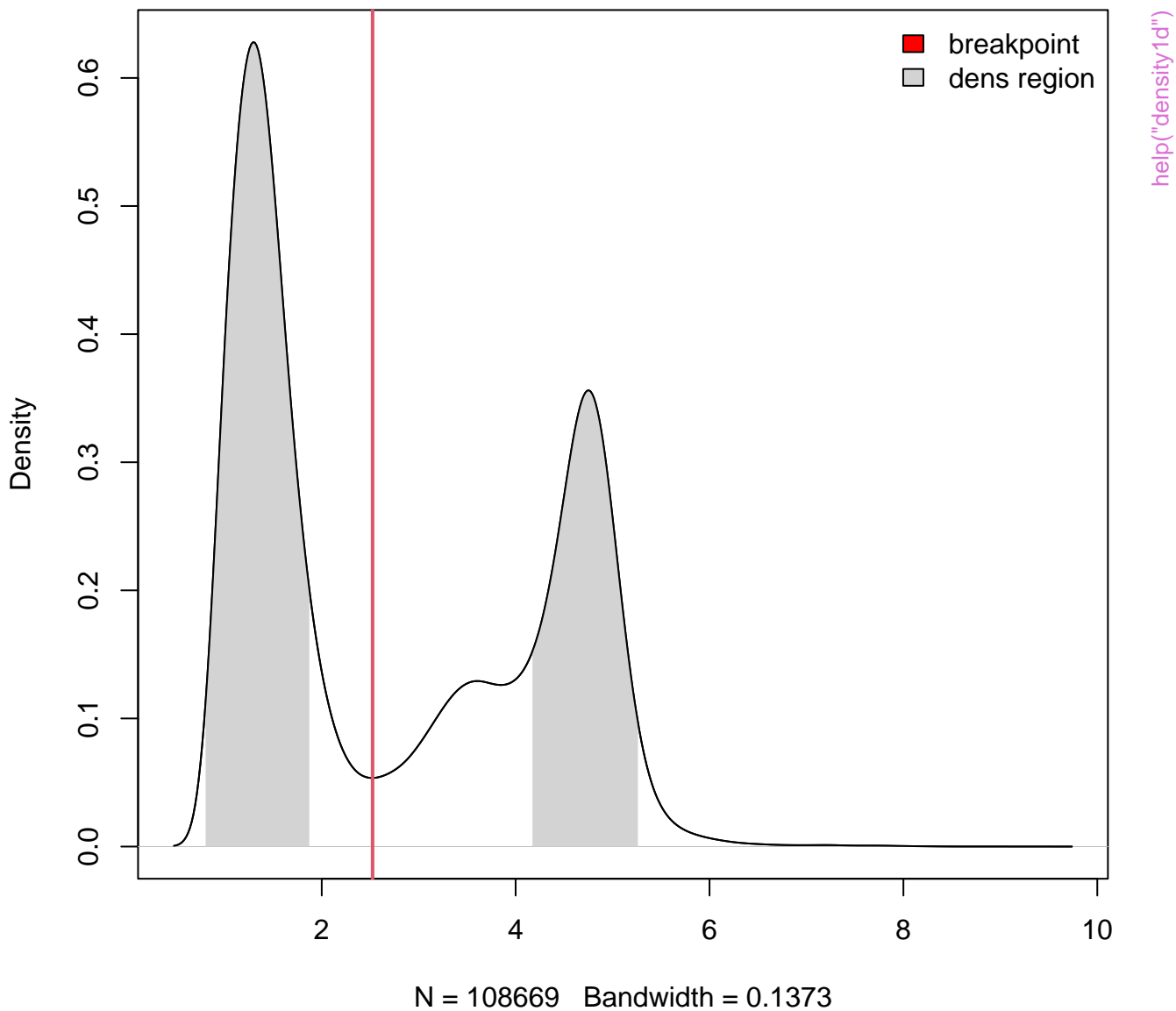
Binned control data



Binned control data



breakpoint for parameter FL4-H



FL4-H

s10a07

s10a06

s10a05

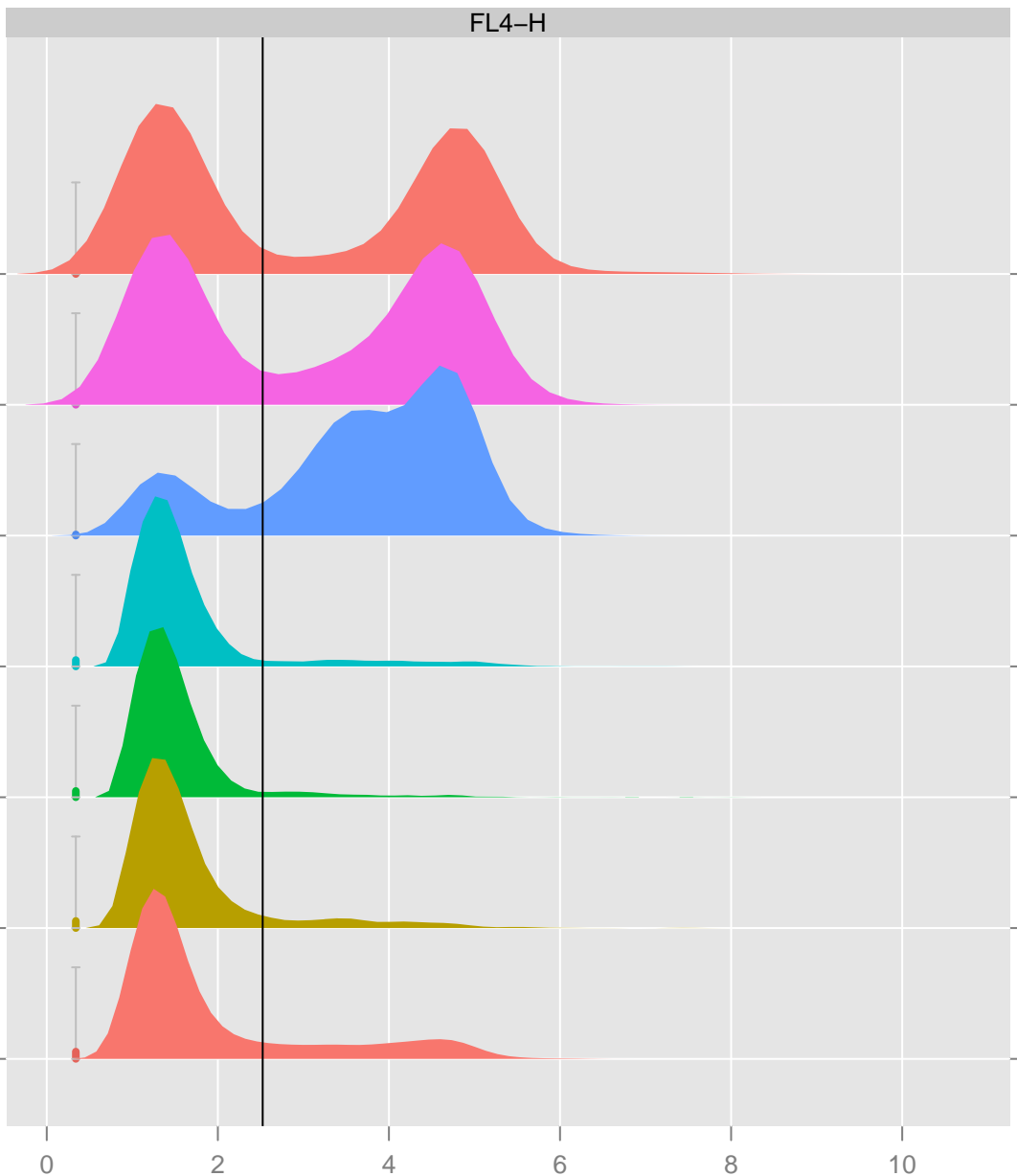
s10a04

s10a03

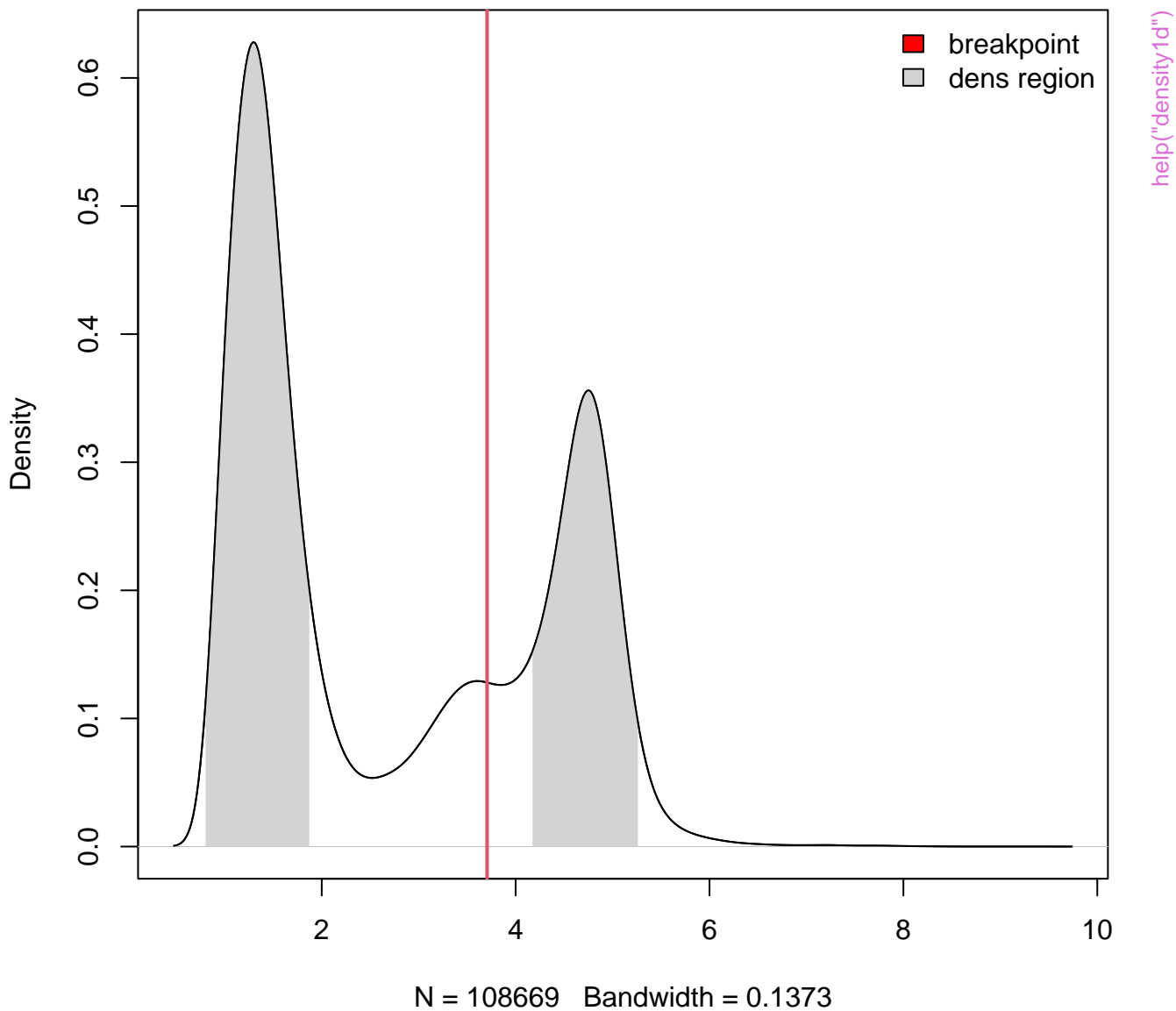
s10a02

s10a01

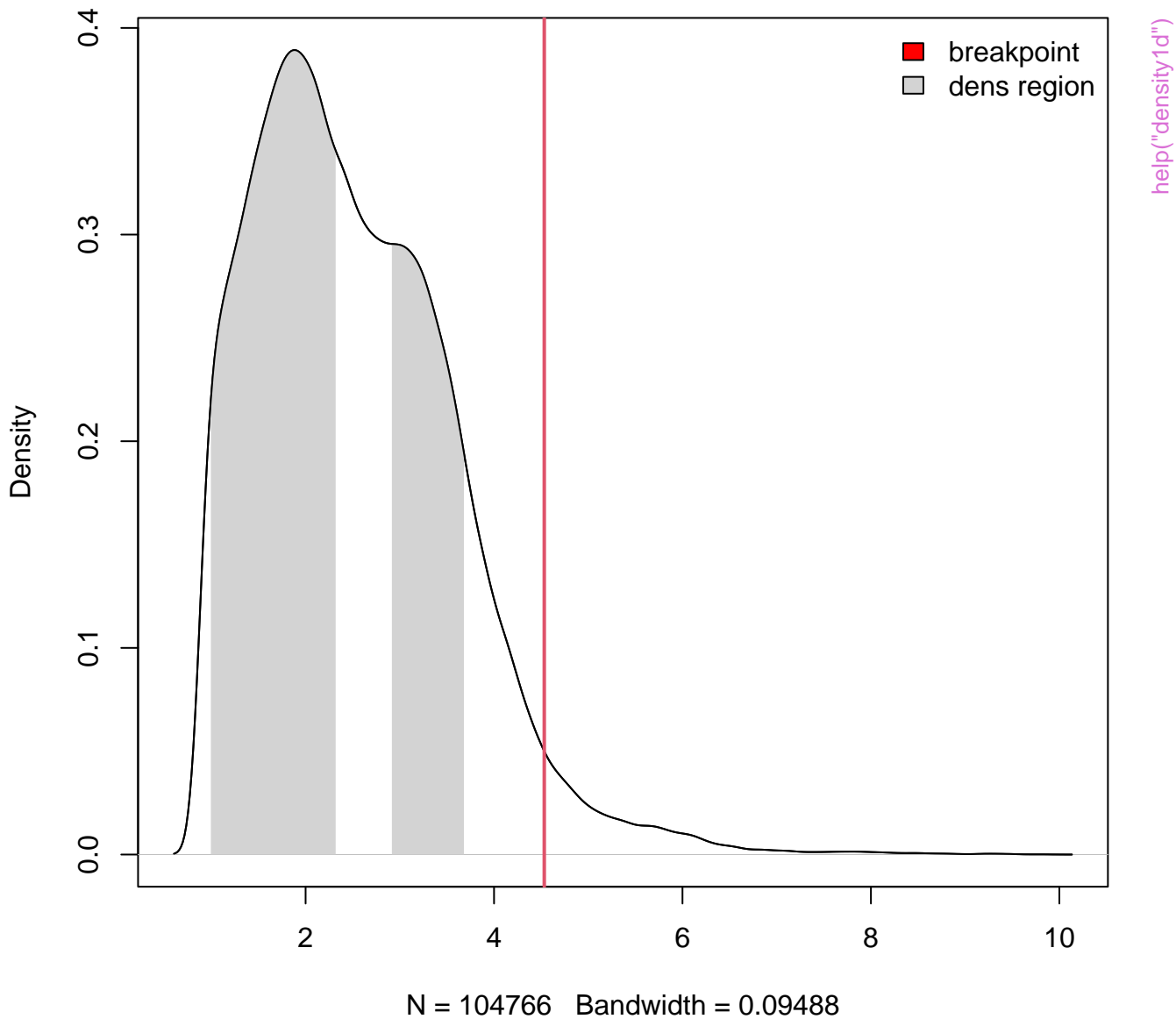
help("density1d")



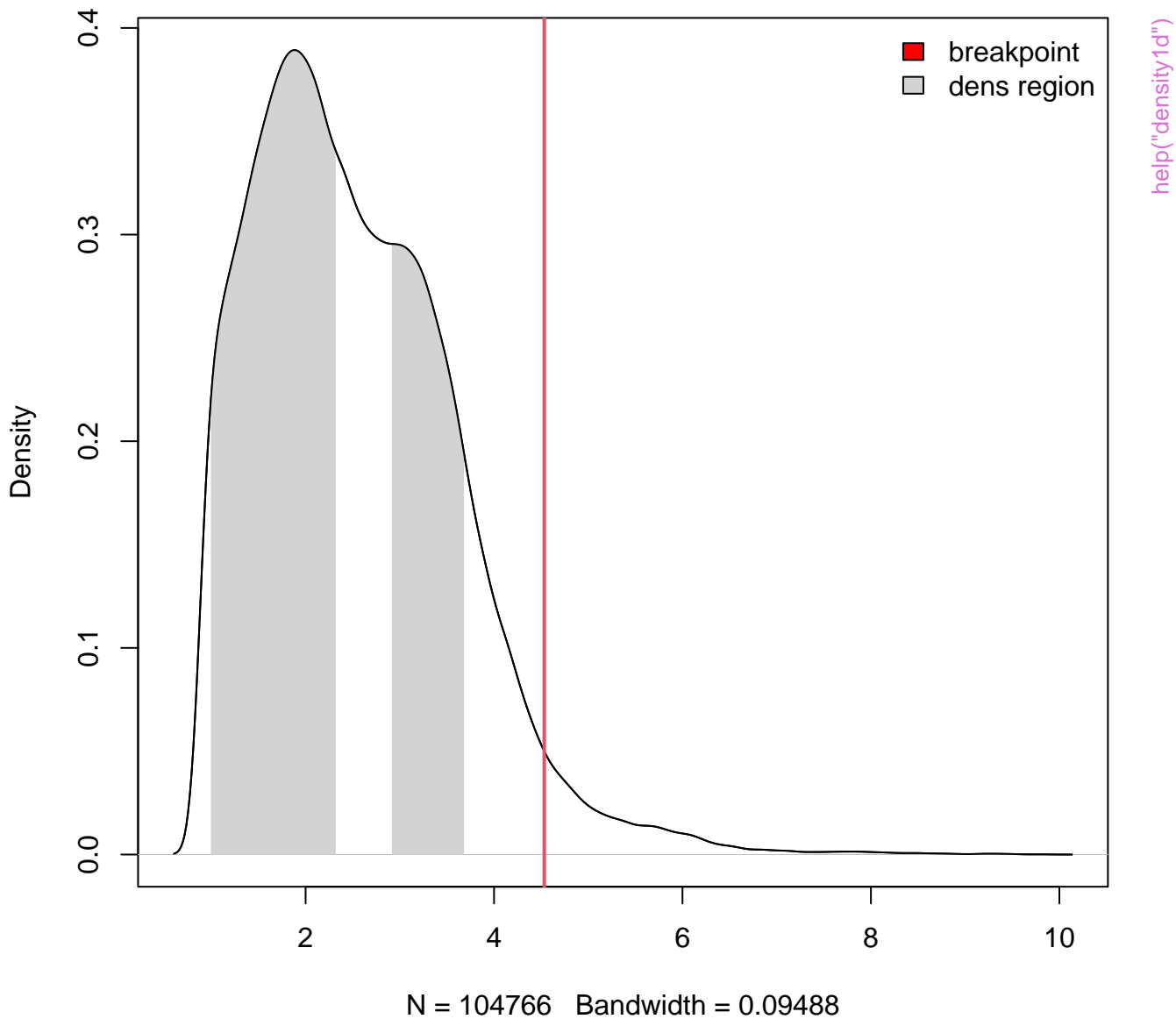
breakpoint for parameter FL4-H



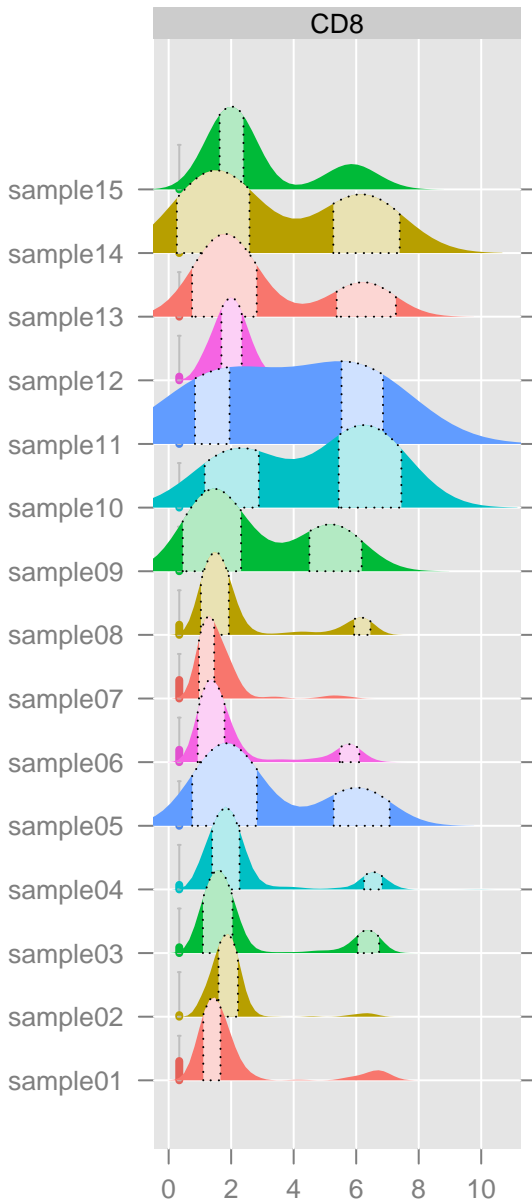
breakpoint for parameter FL3-H



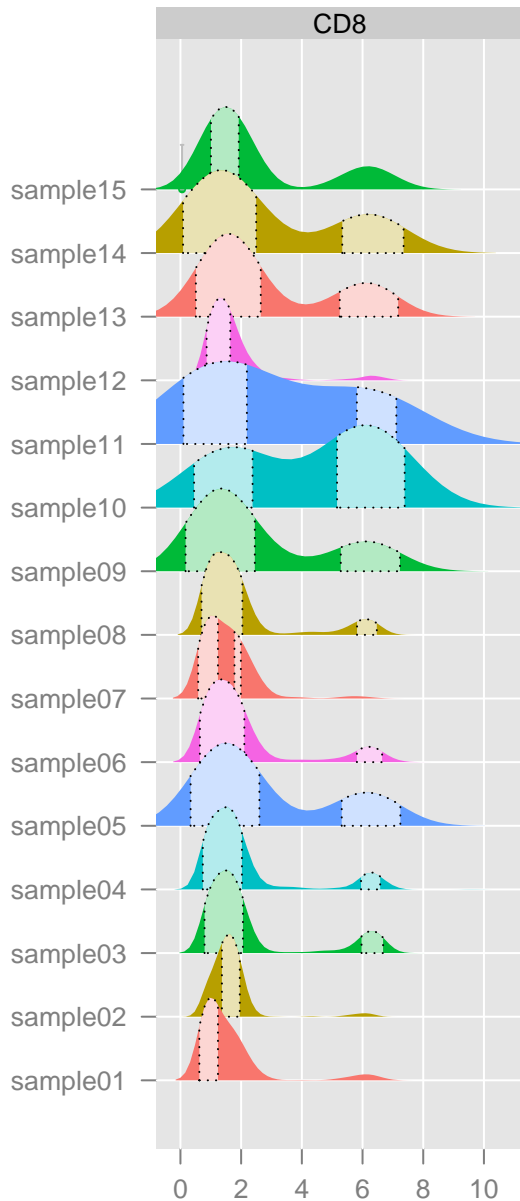
breakpoint for parameter FL3-H



original



normalized



help("gaussNorm")

After GPA using bg

FSC

SSC

sample15

sample14

sample13

sample12

sample11

sample10

sample09

sample08

sample07

sample06

sample05

sample04

sample03

sample02

sample01

0

200

400

600

800

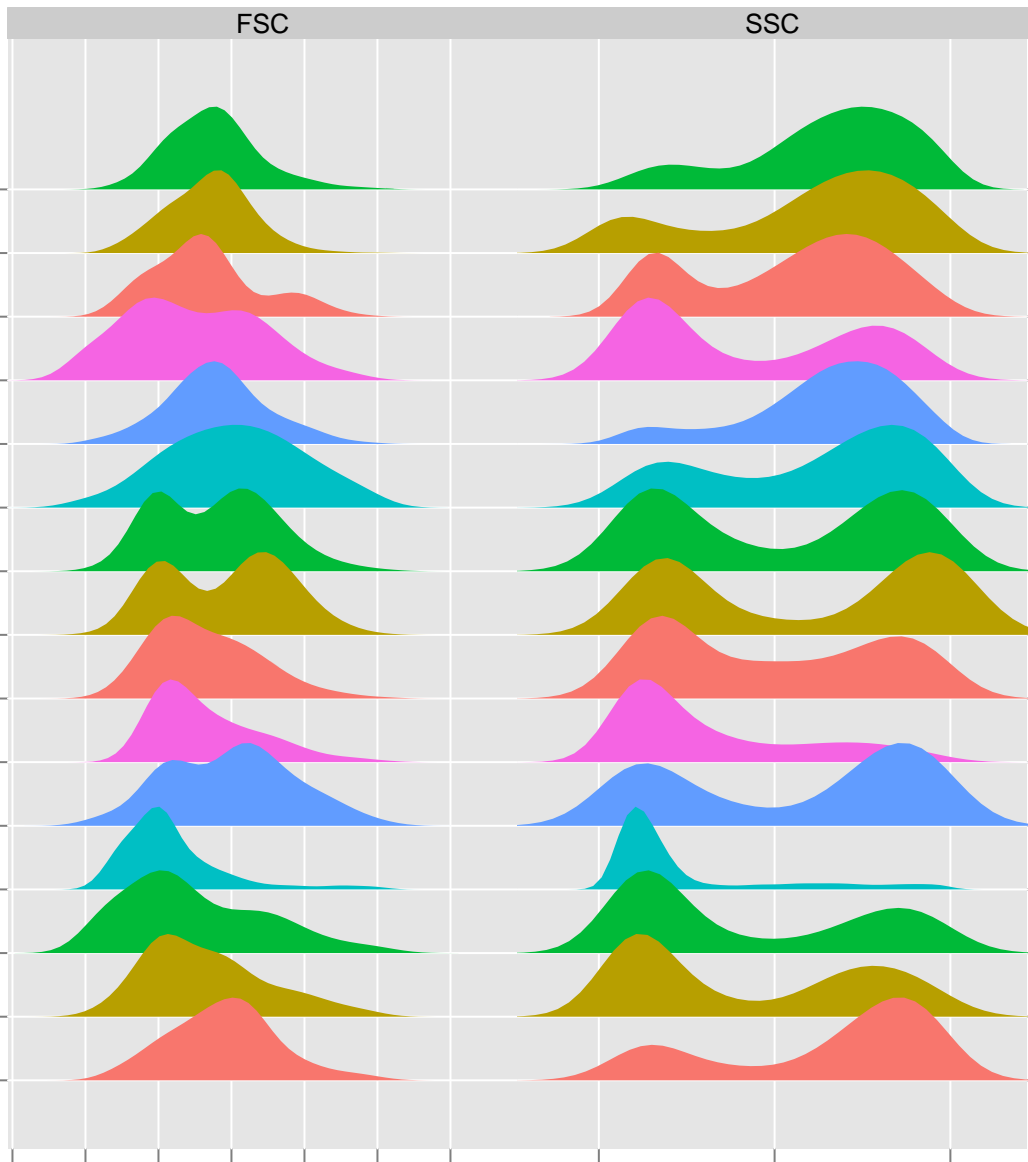
1000

0

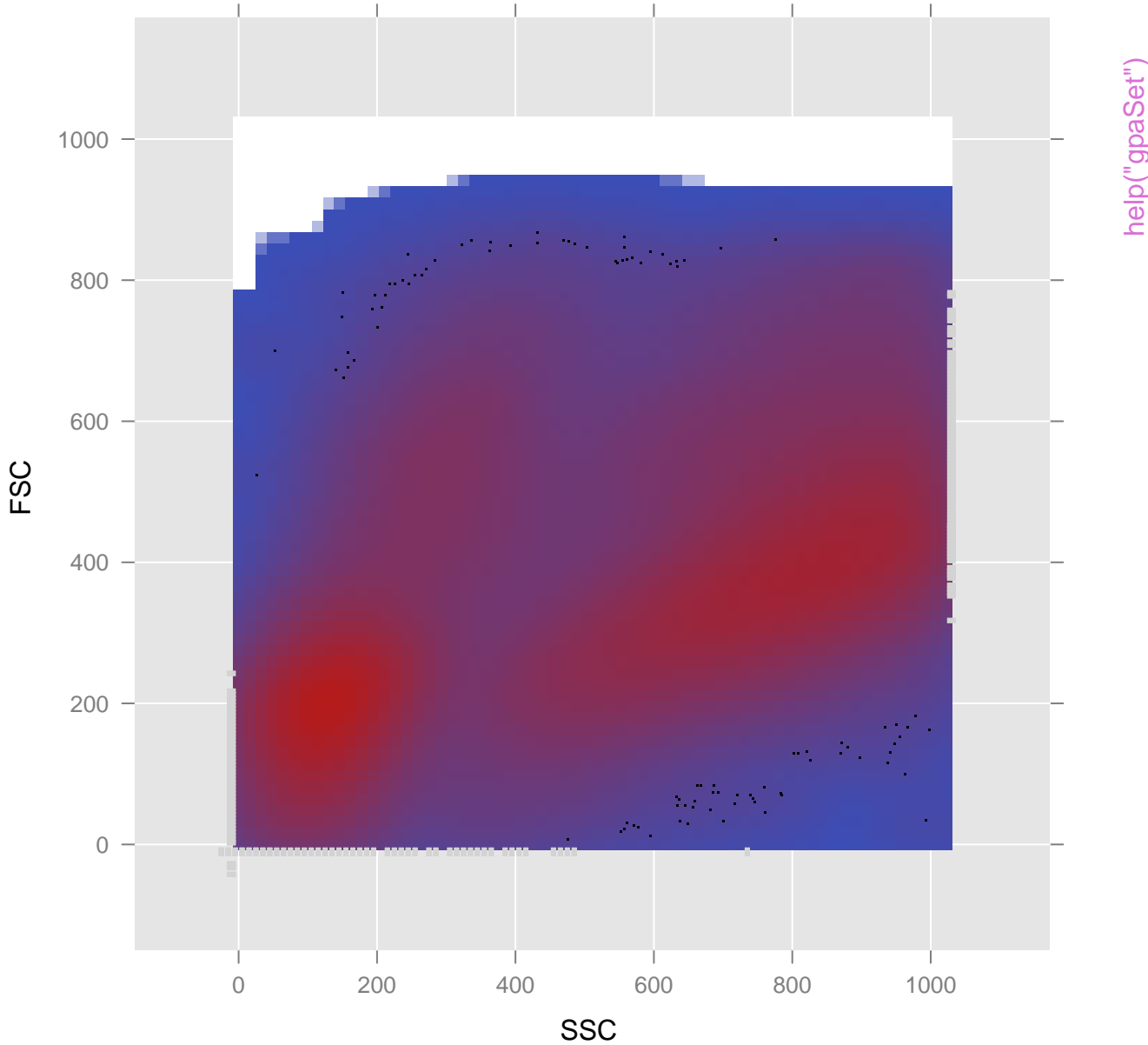
500

1000

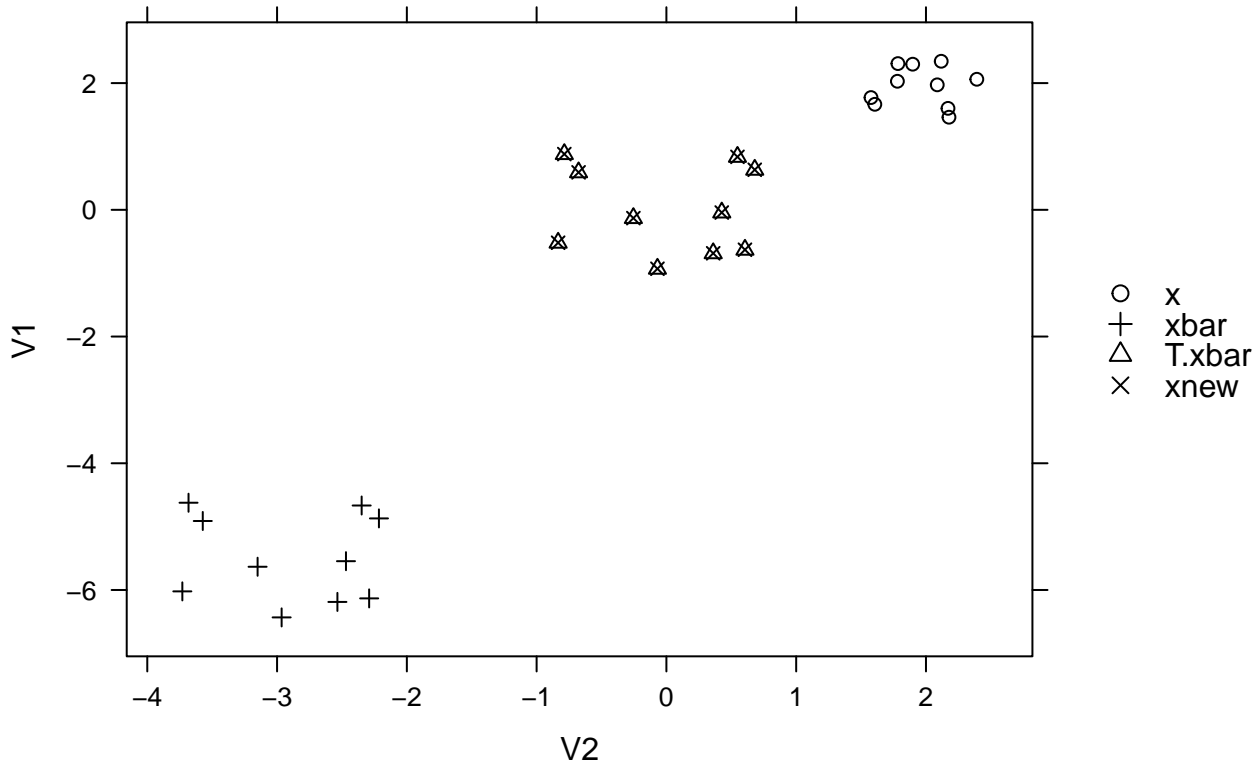
help("gpaSet")

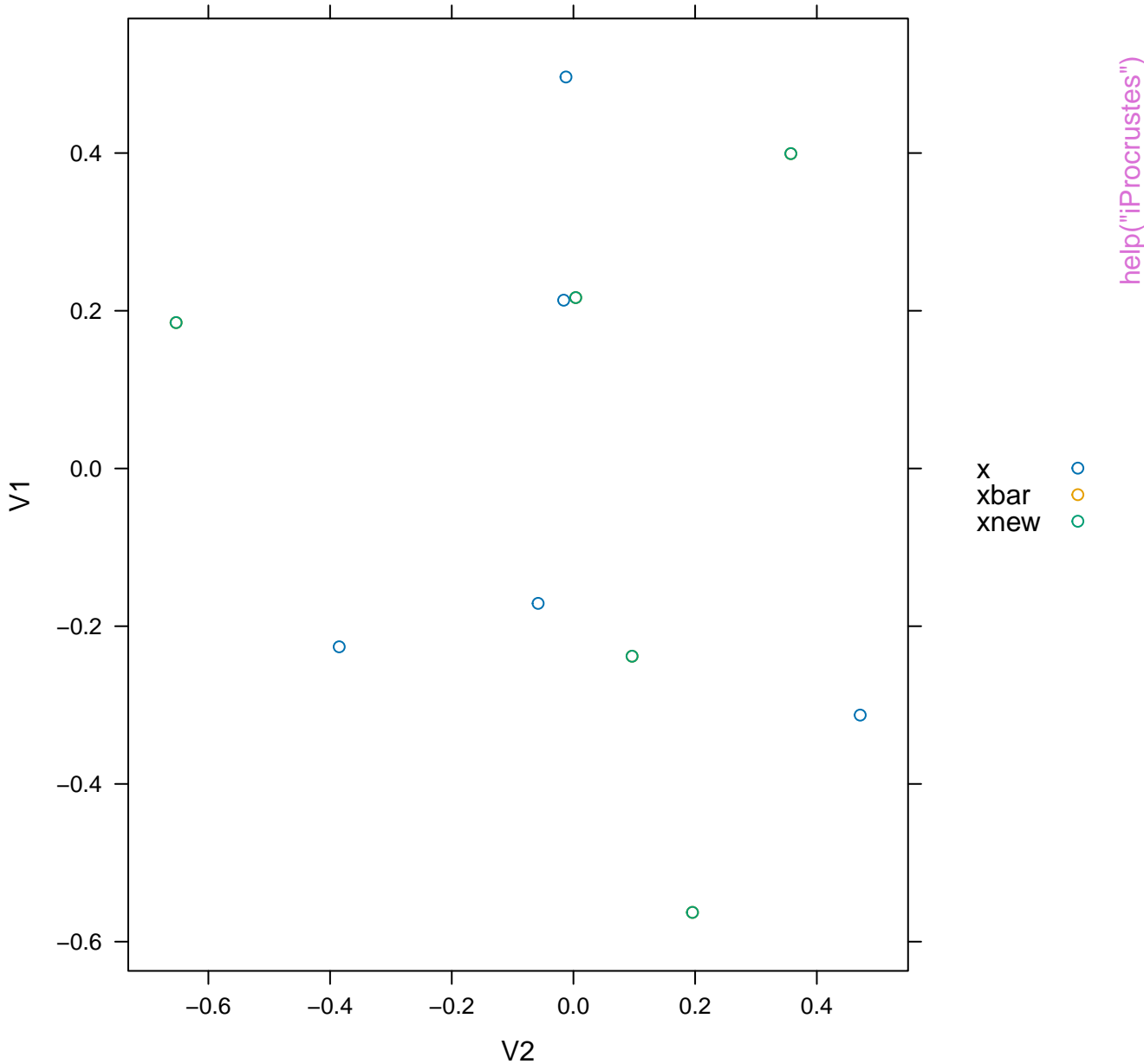


All flowFrames

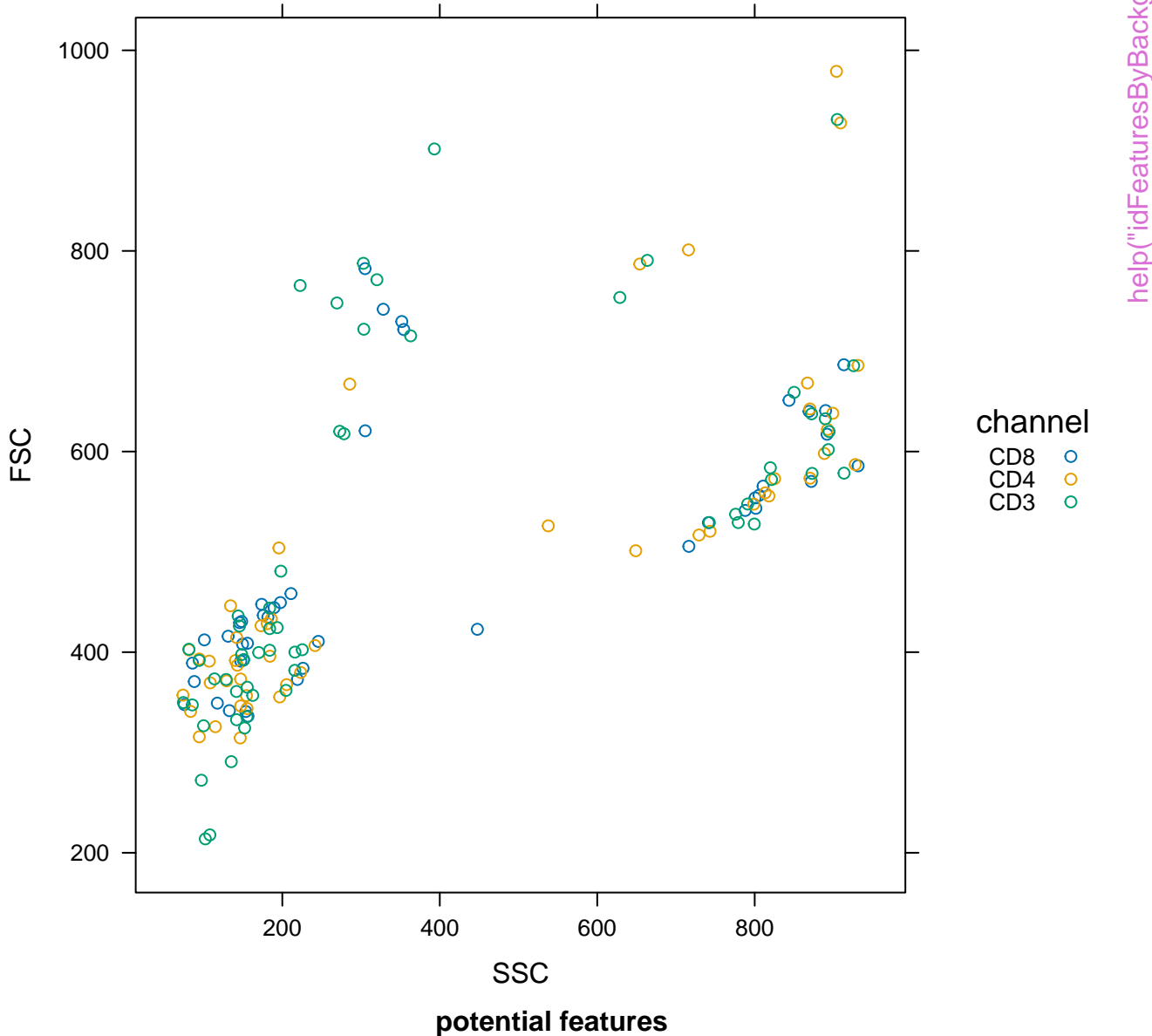


Align the points in x to xbar

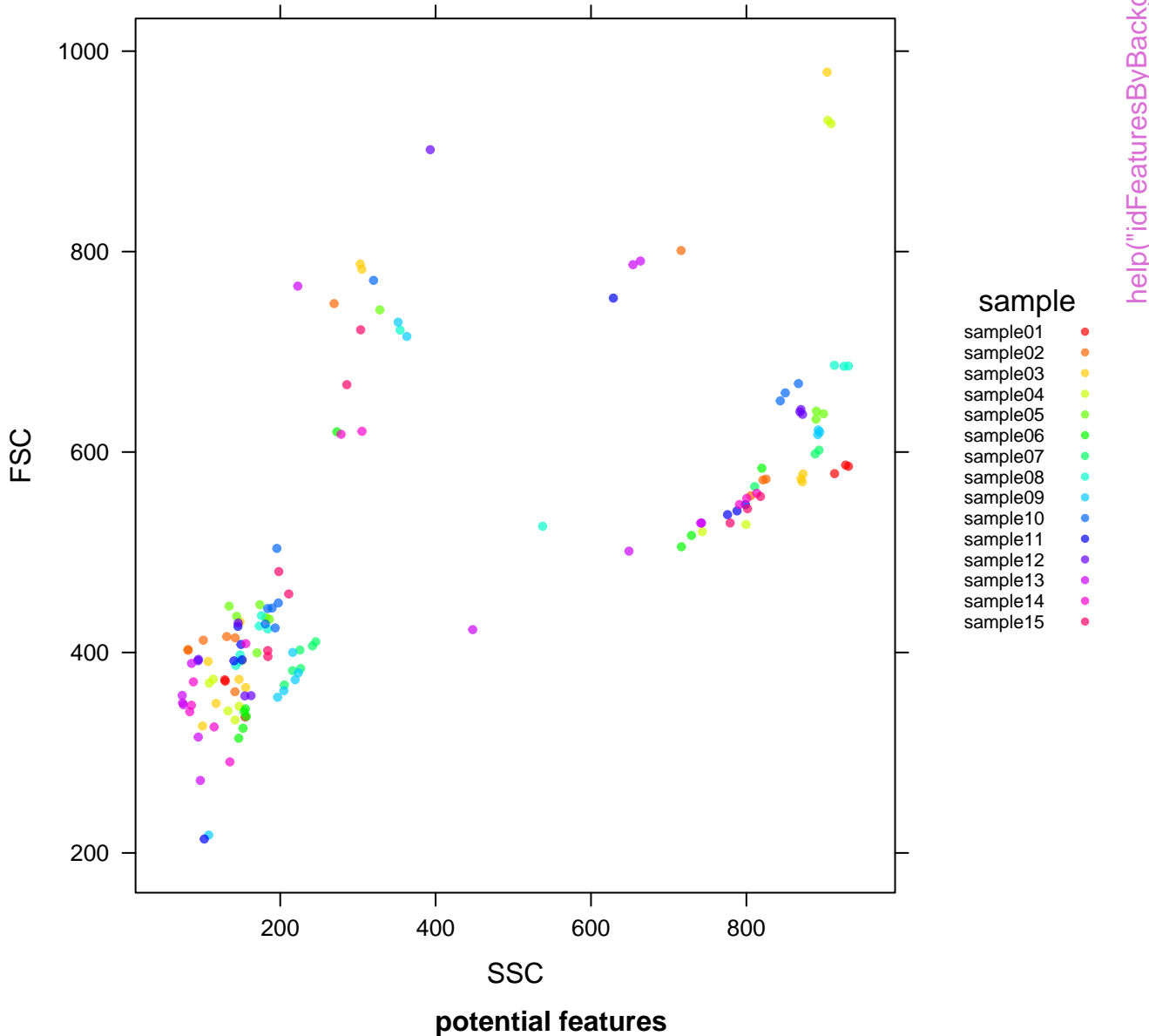




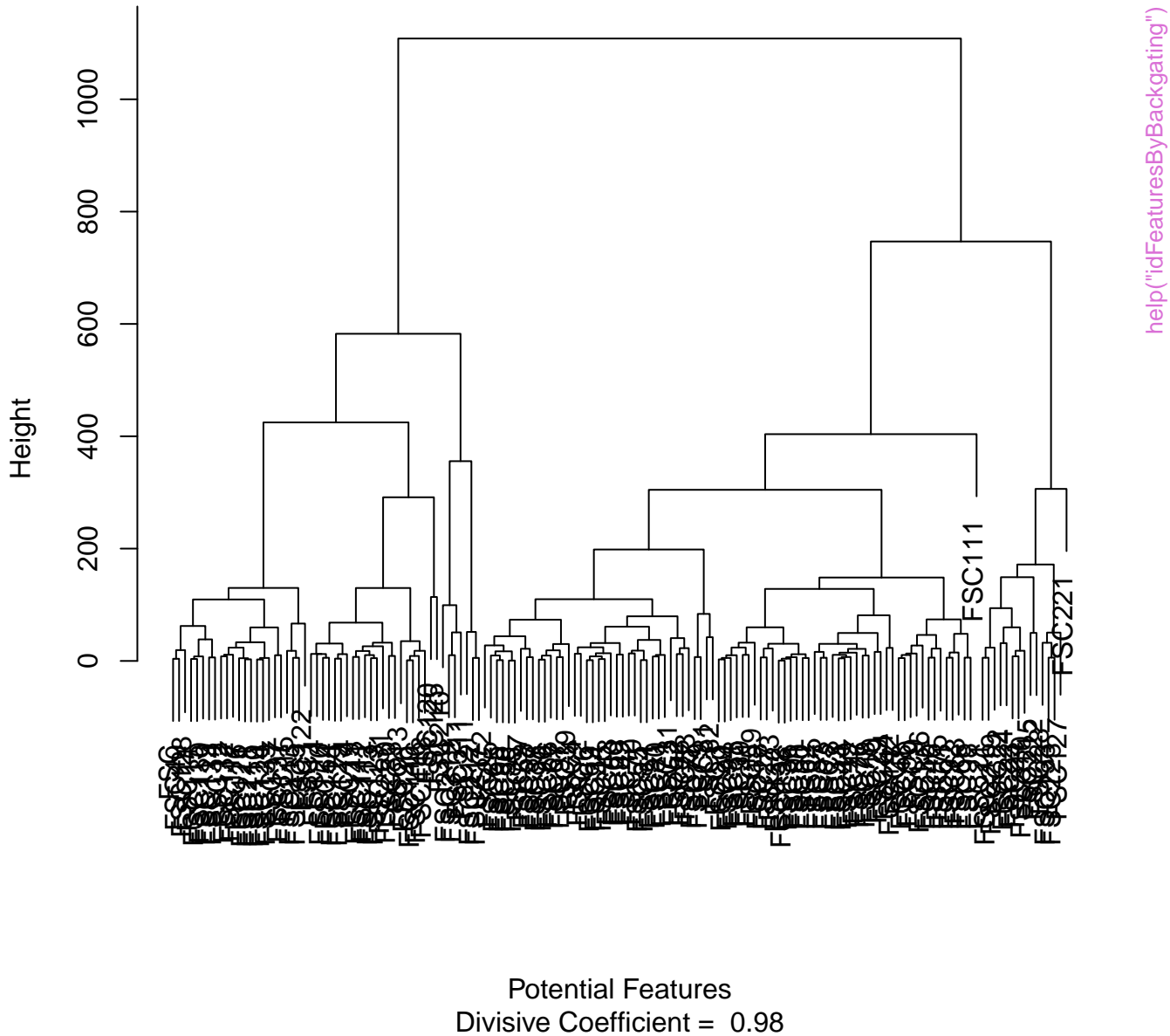
Centroids of high-density areas yielded by backgating on channel CD8 CD4 CD3



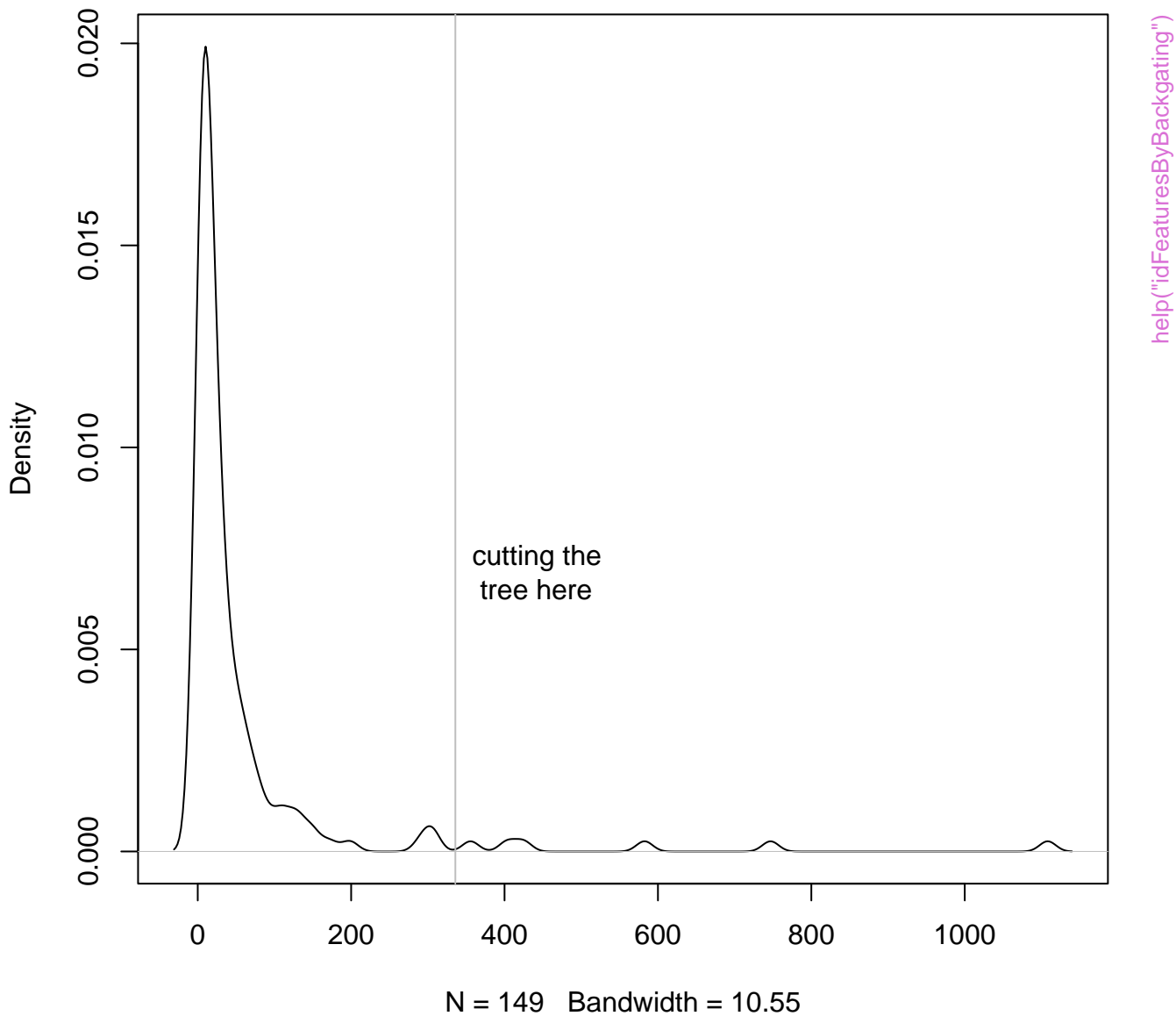
Centroids of high-density areas yielded by backgating on channel CD8 CD4 CD3



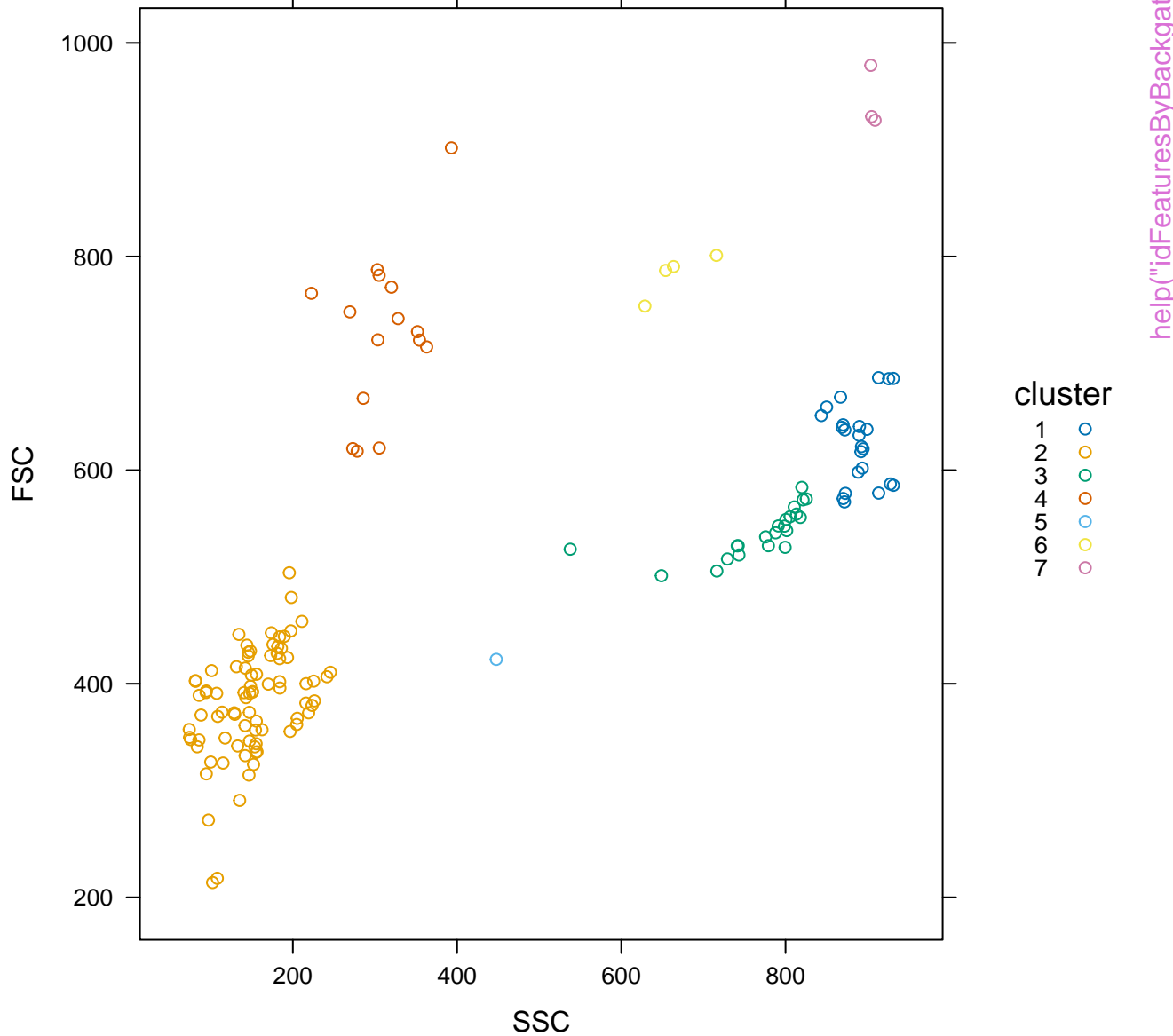
Cut the tree at height 336



Distribution of the cluster (diana) height

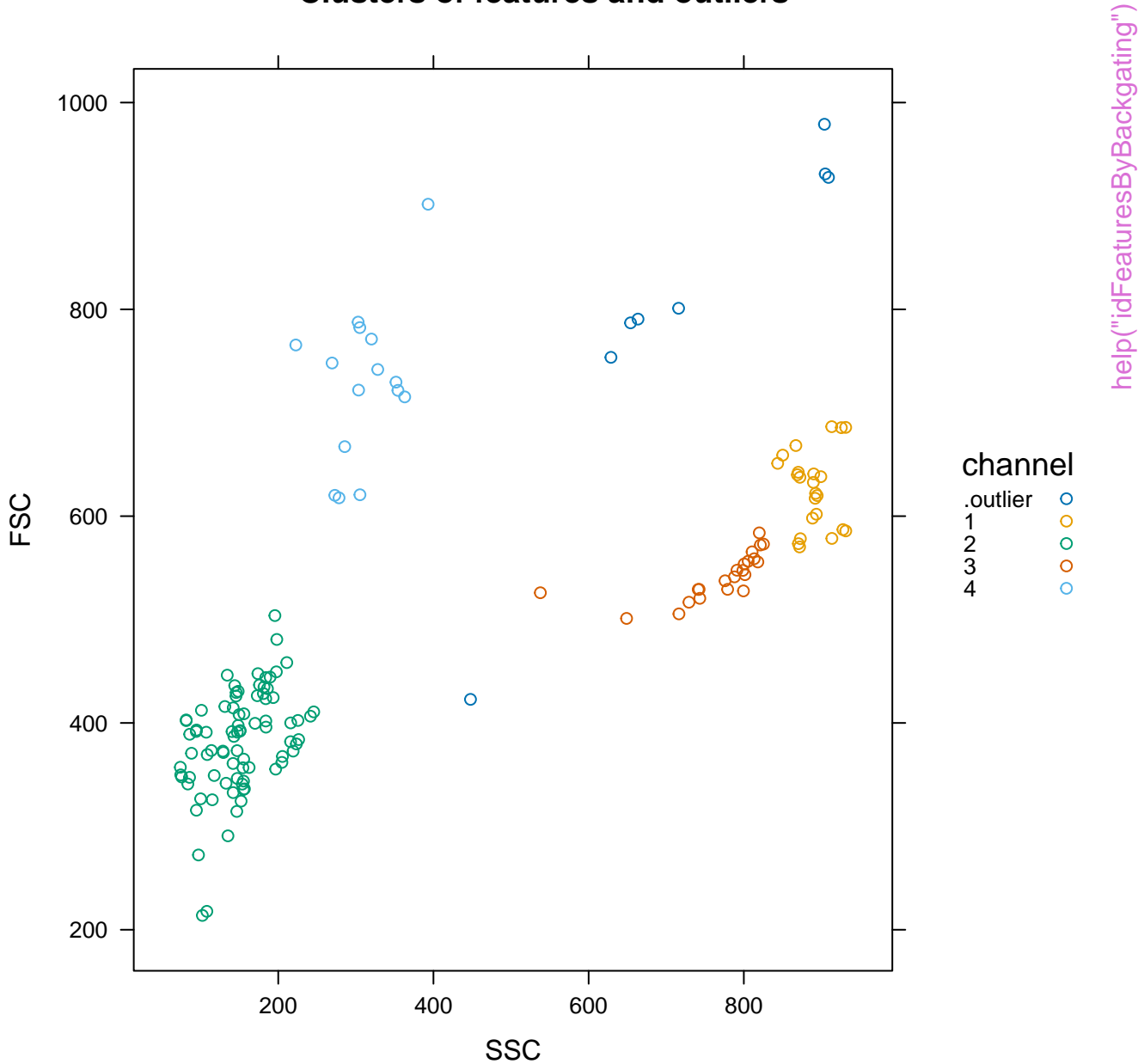


Clusters of potential features



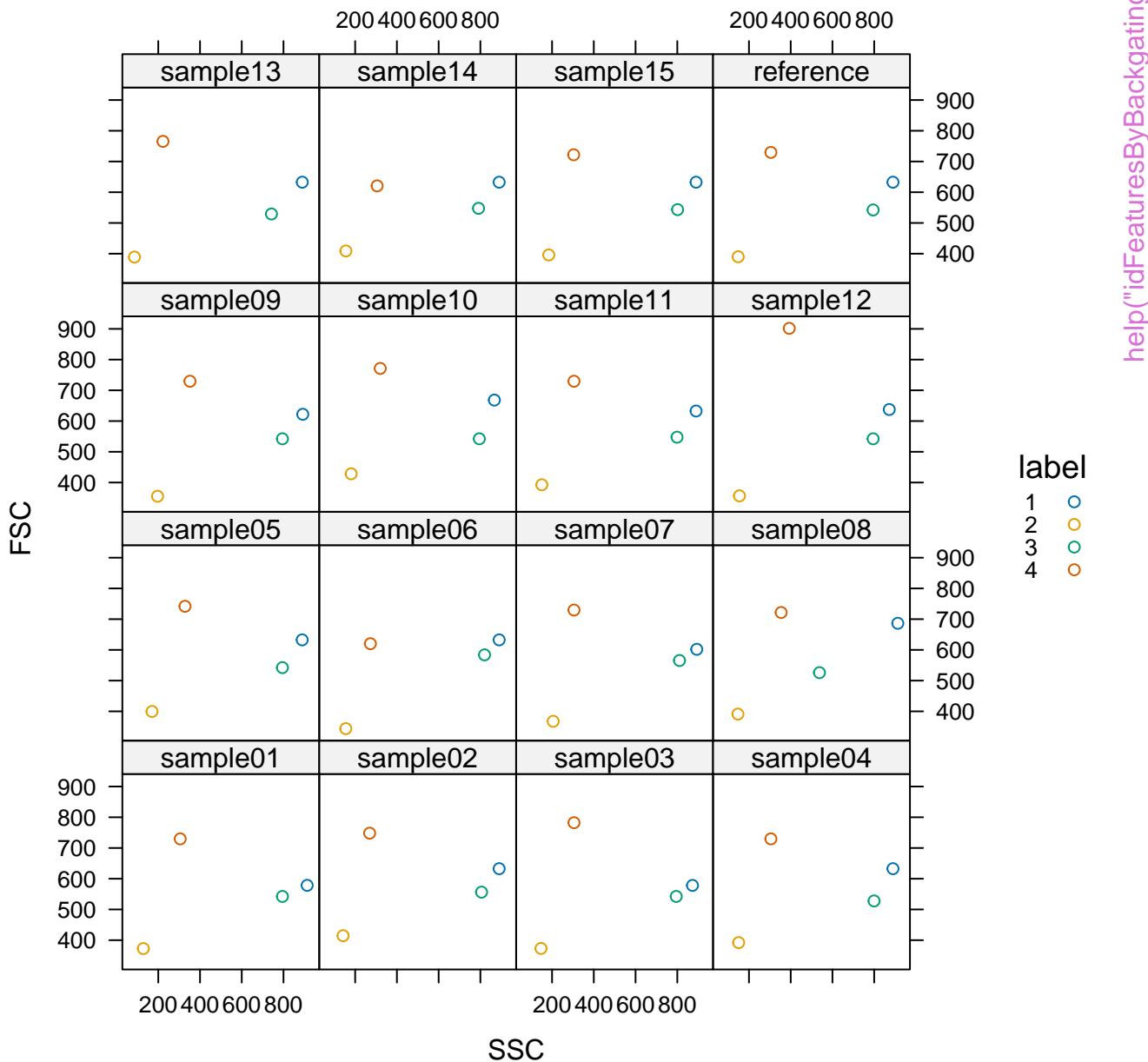
Clusters containing < 336 members are considered outliers.

Clusters of features and outliers

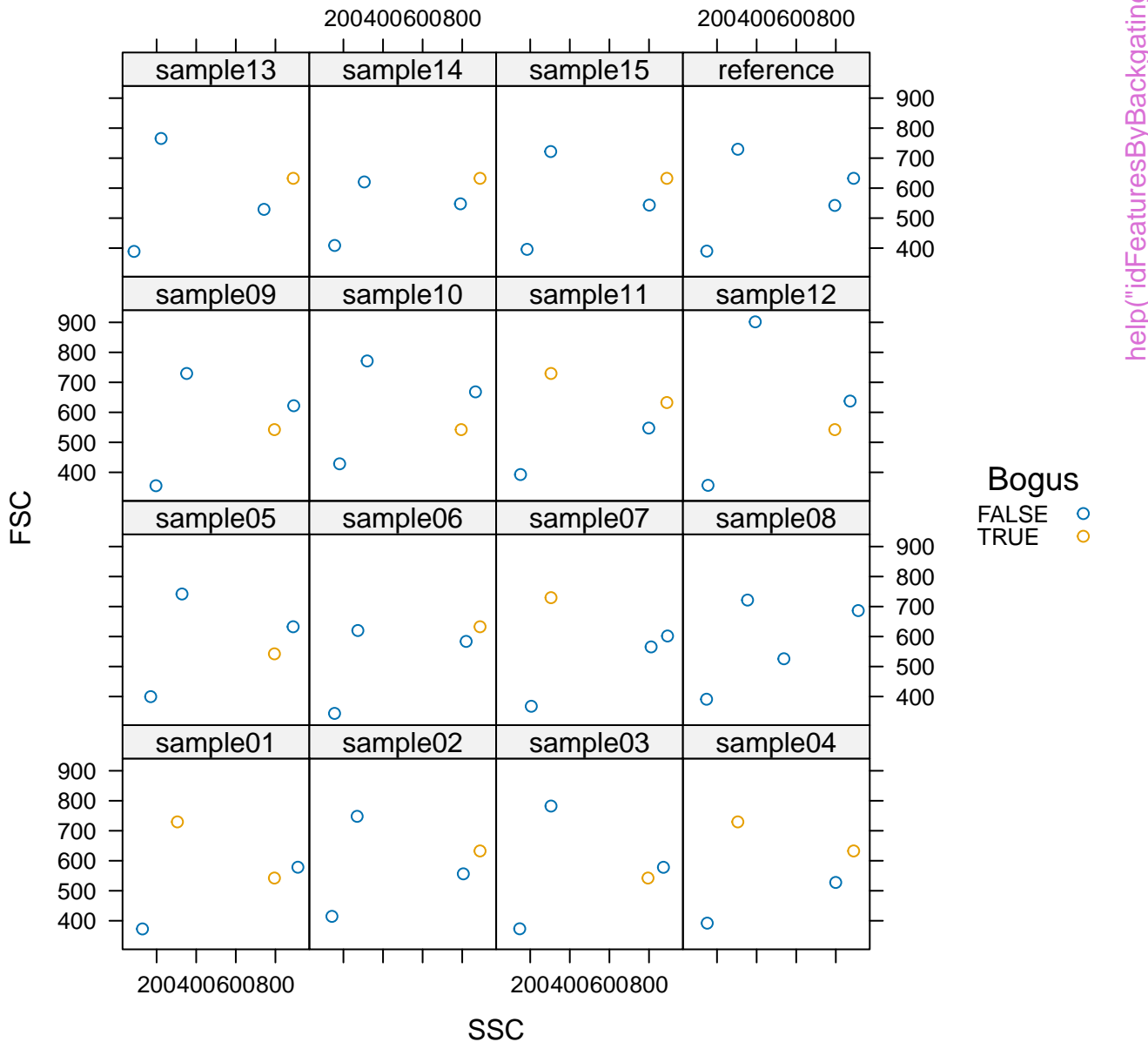


Clusters containing < 336 members are considered outliers.

Labeled features

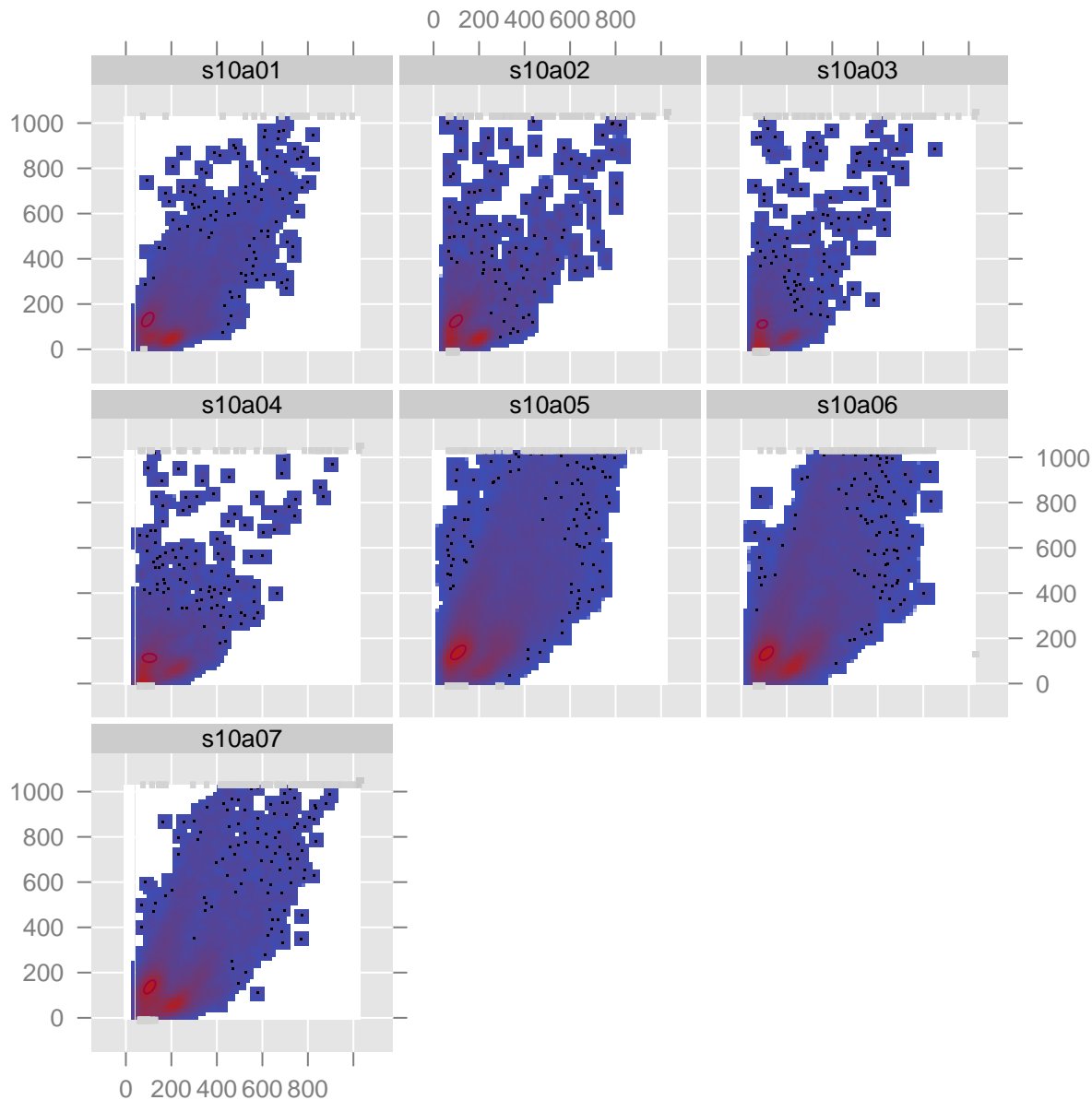


Labeled features



Missing features are those with bogus=TURE.

SSC-H SSC-Height



FSC-H FSC-Height

help("lymphGate")

Binned control data

