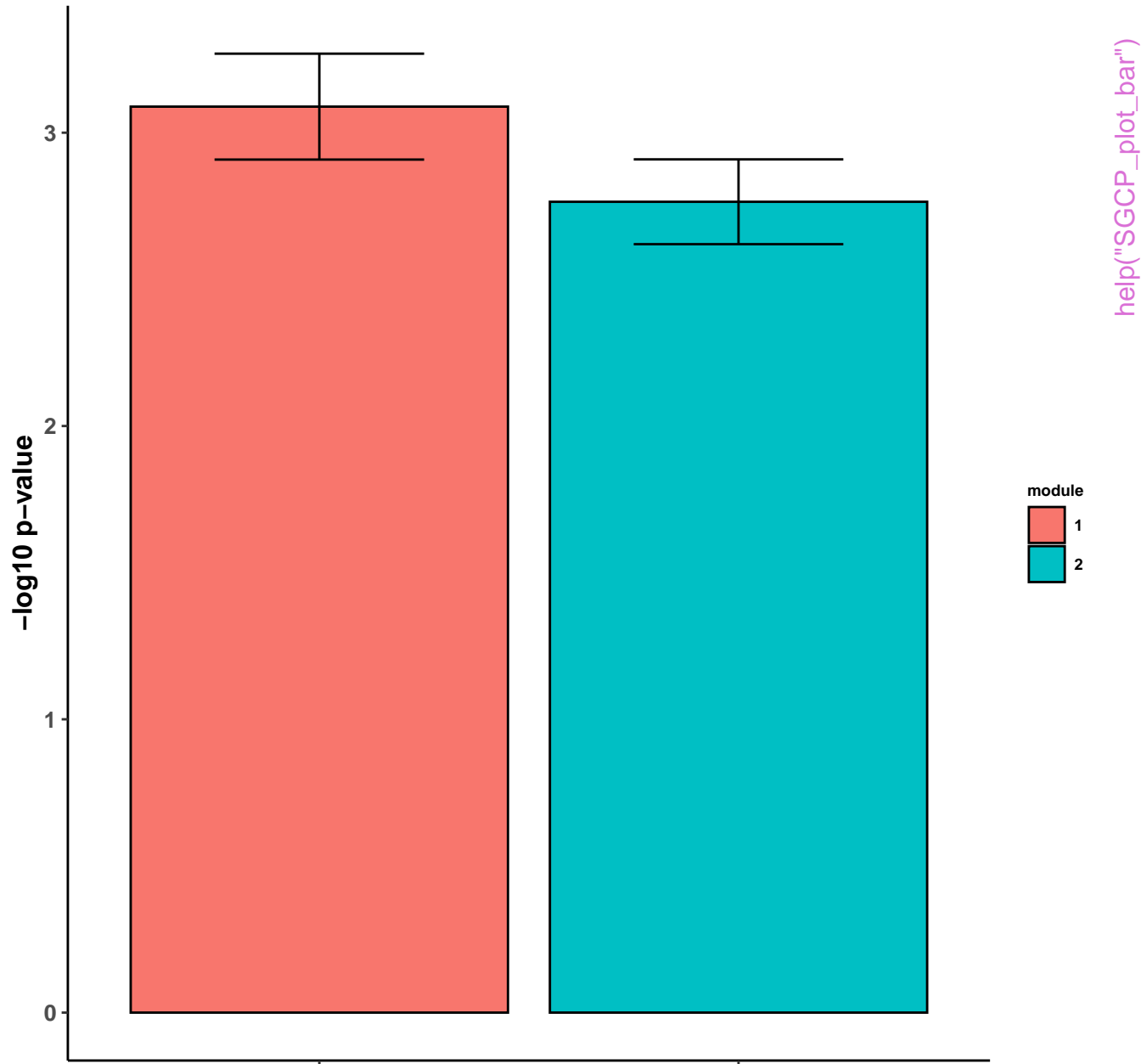
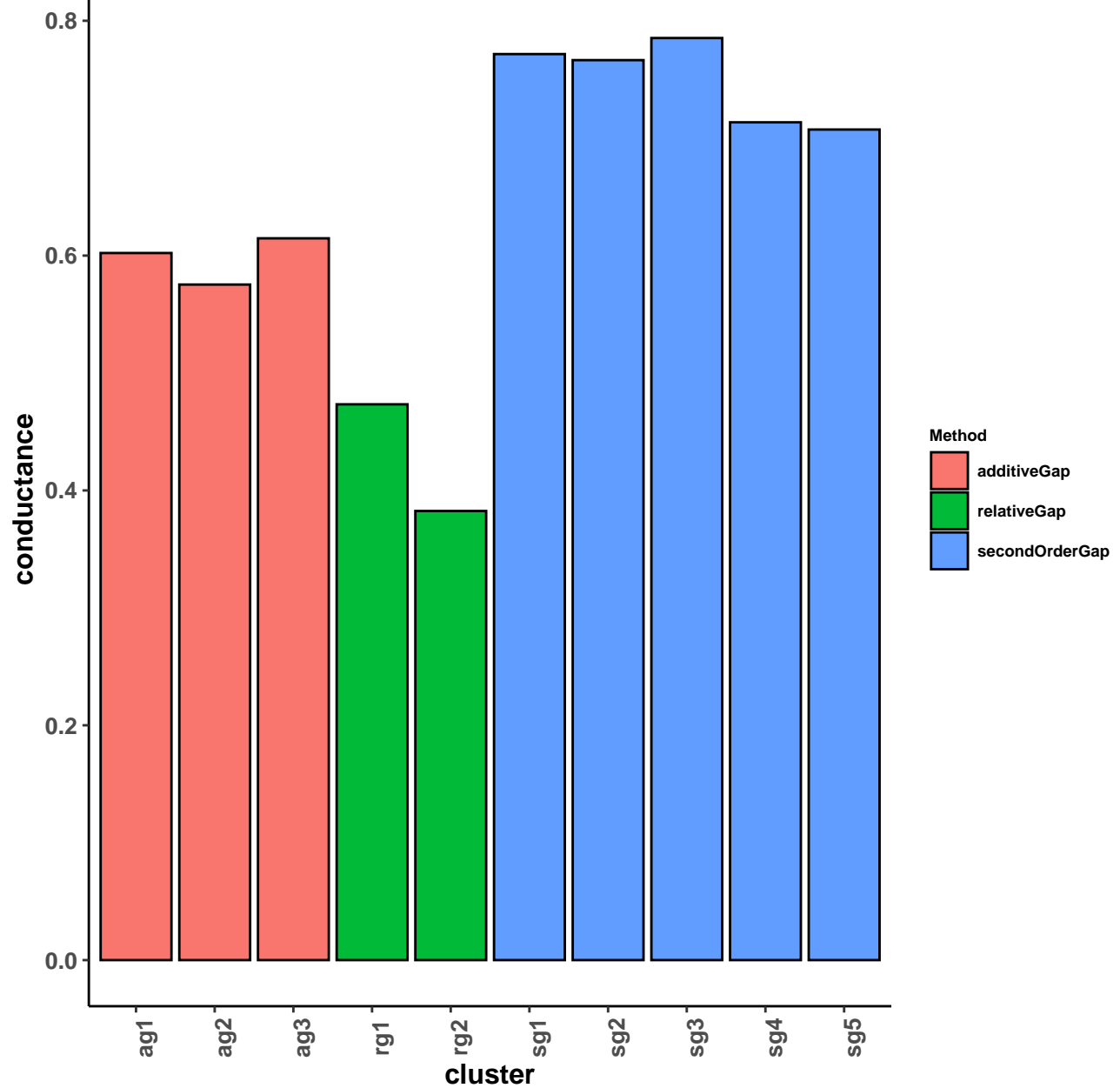


mean $-\log_{10}$ p-values



Clustering Conductance Index



help("SGCP_plot_conductance")

p-values Density

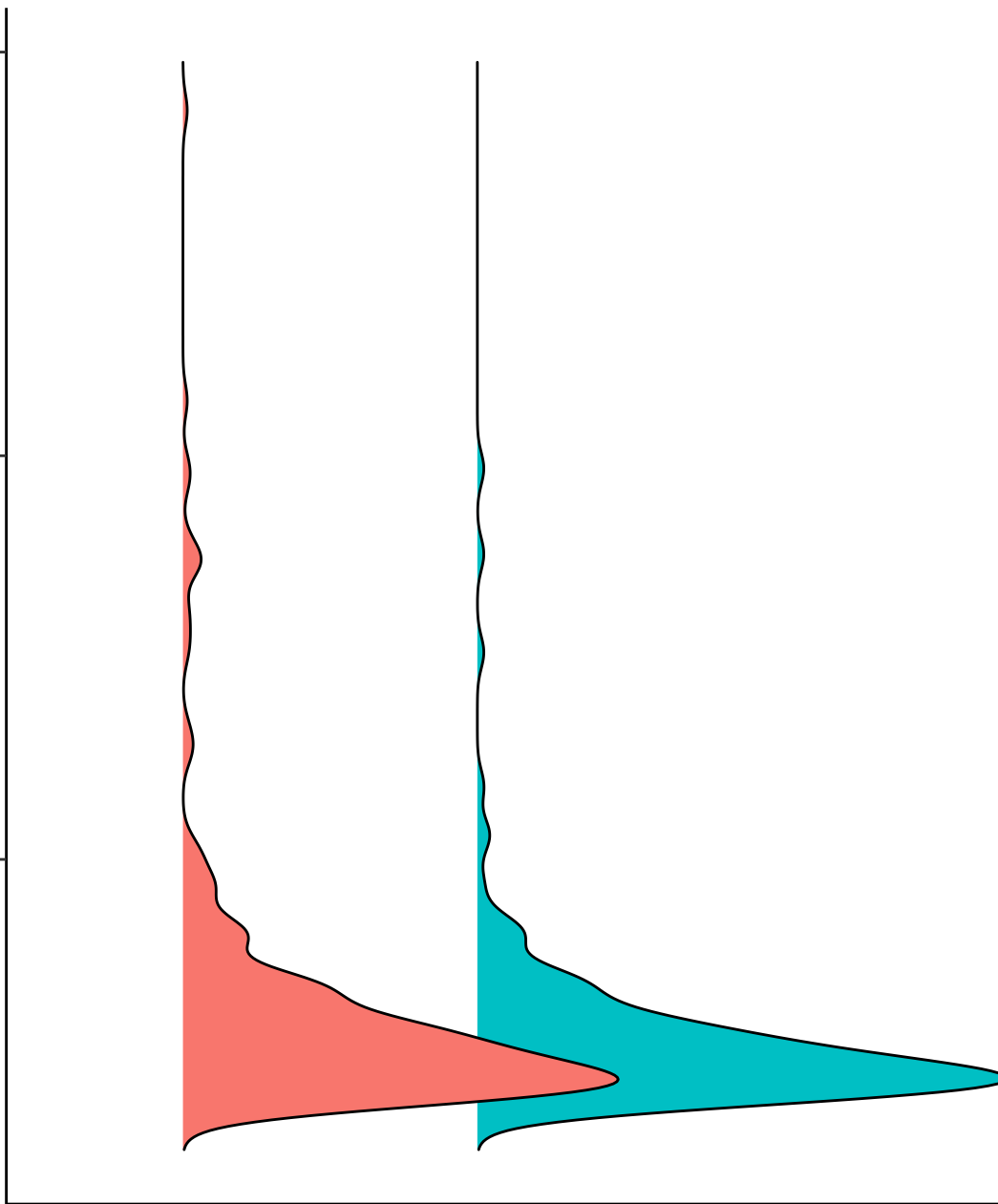
$-\log_{10}$ p-value

15
10
5

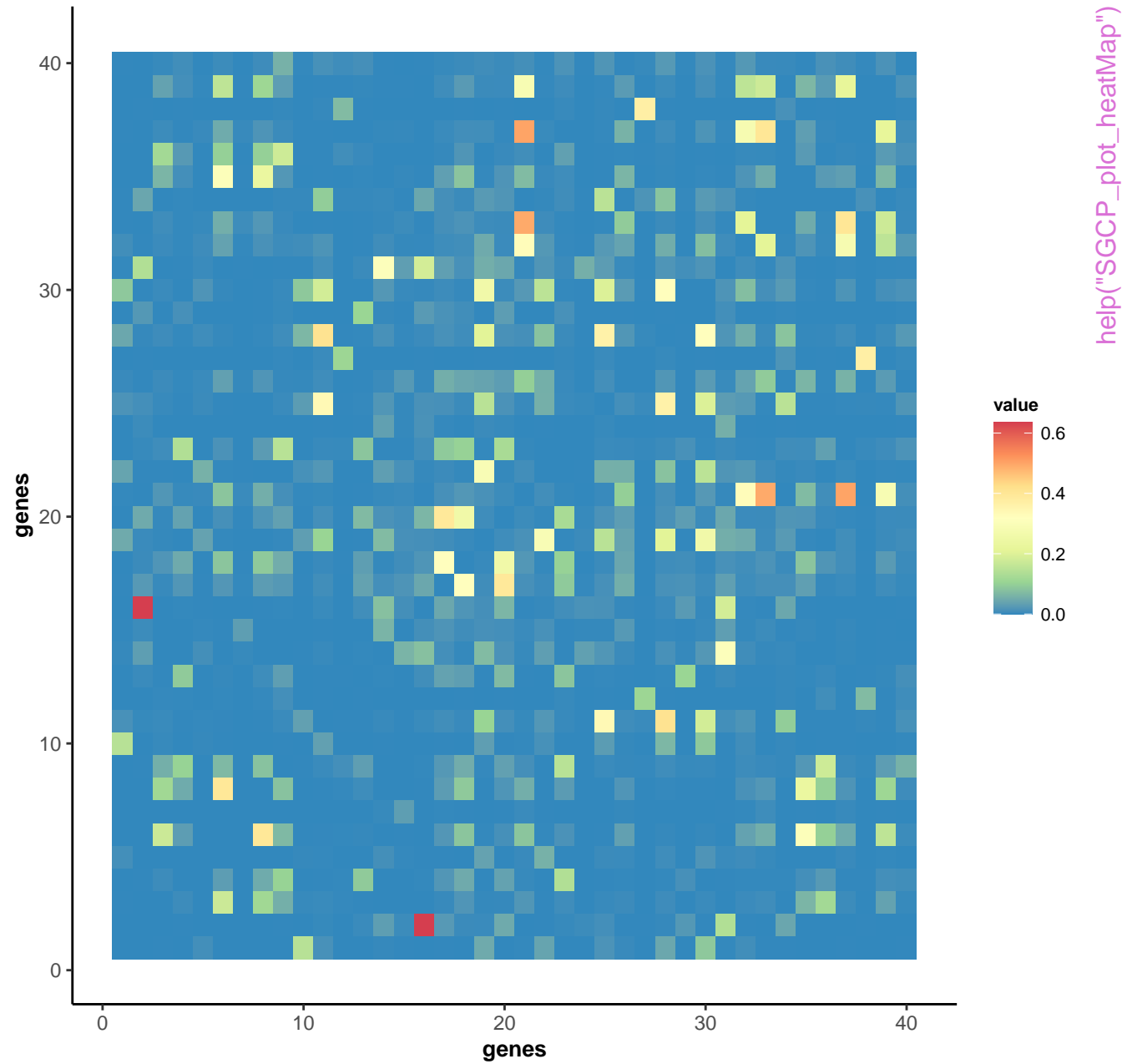
module

1
2

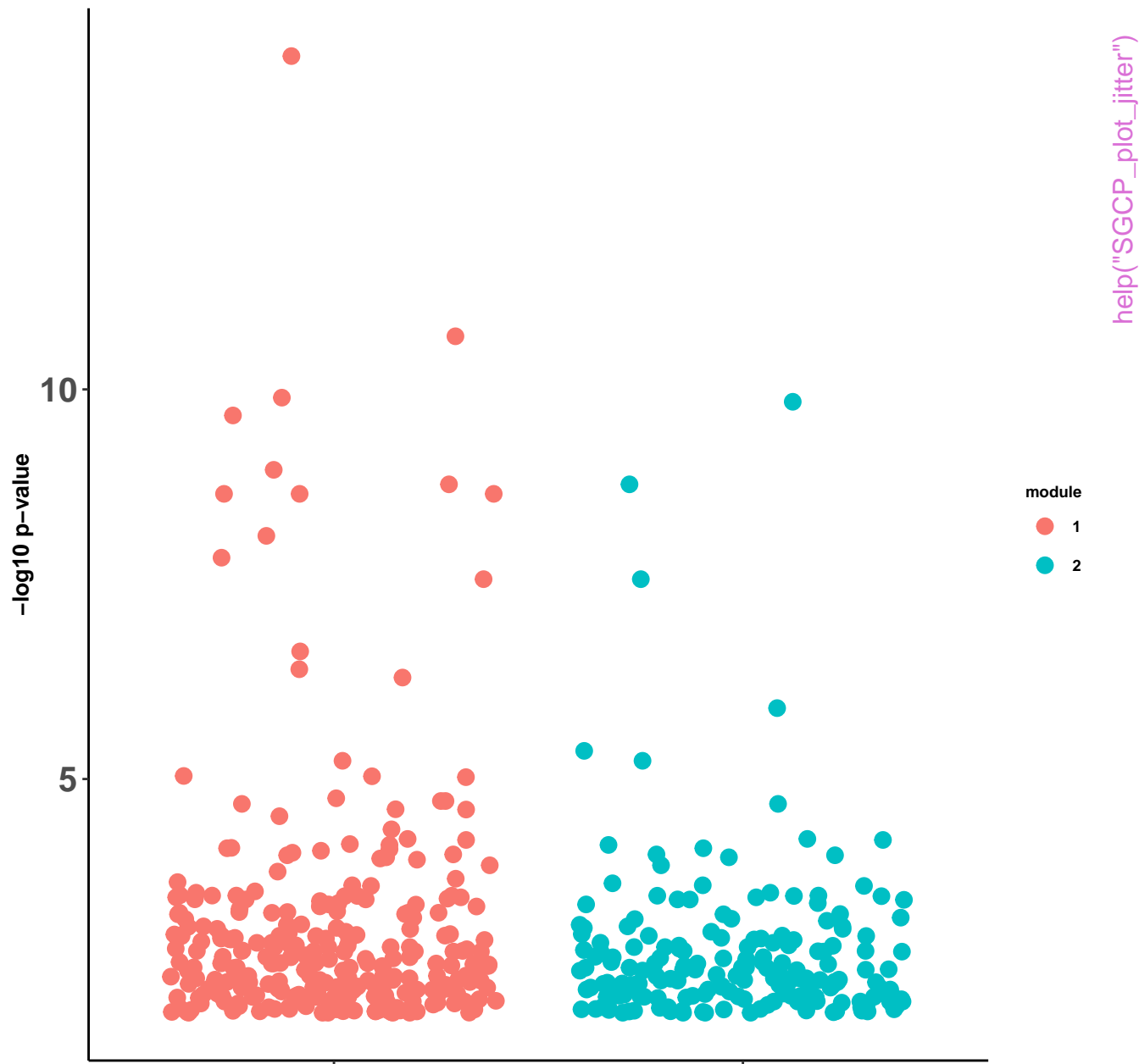
help("SGCP_plot_density")



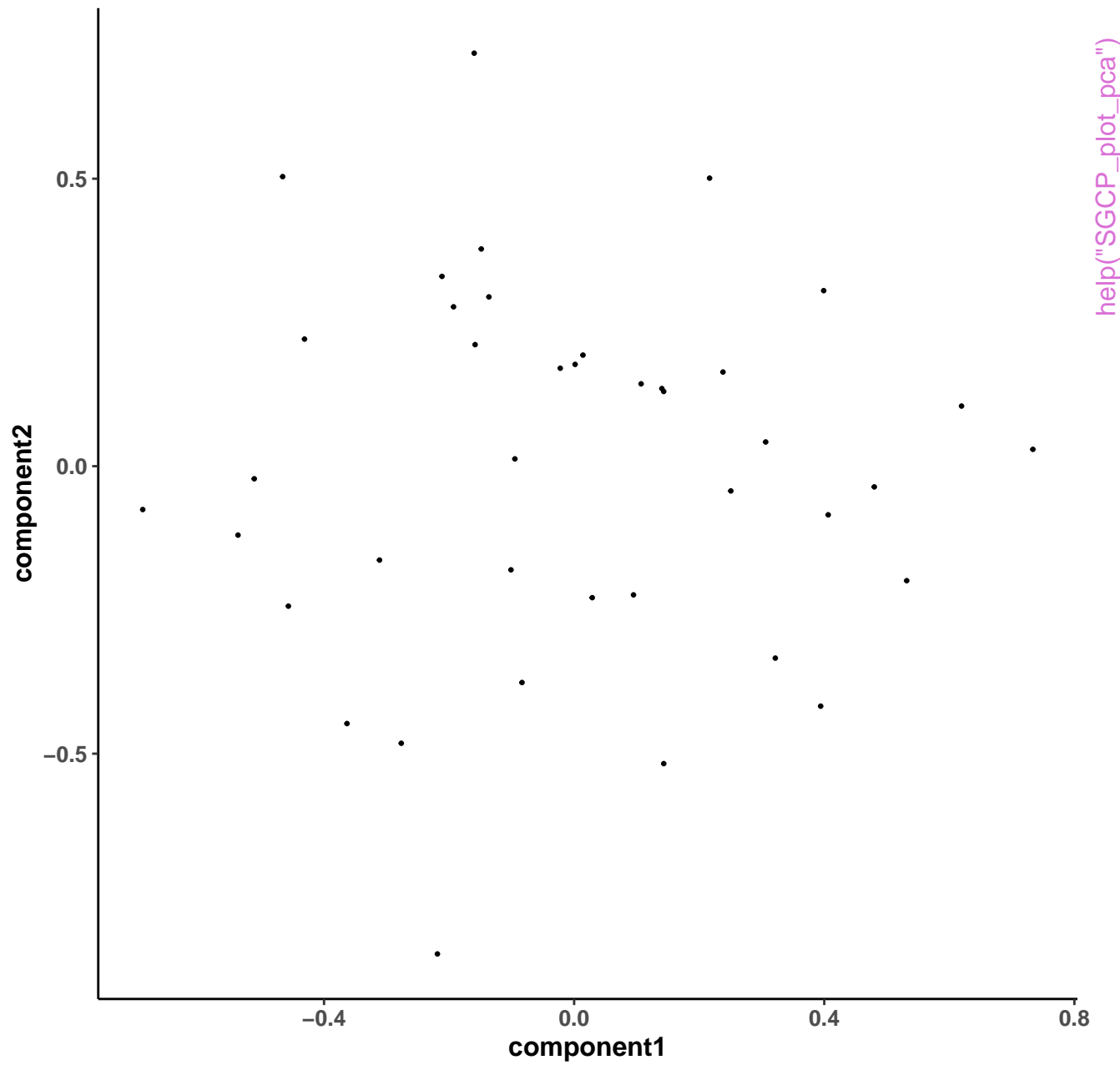
Adjacency Heatmap



p-values Distribution

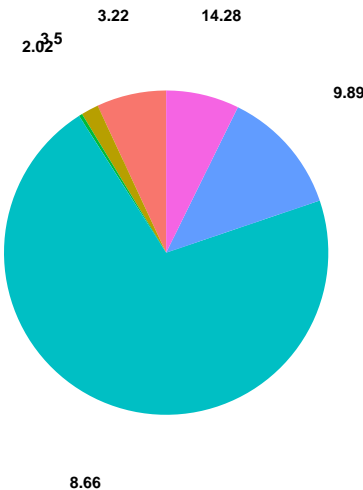


PCA plot

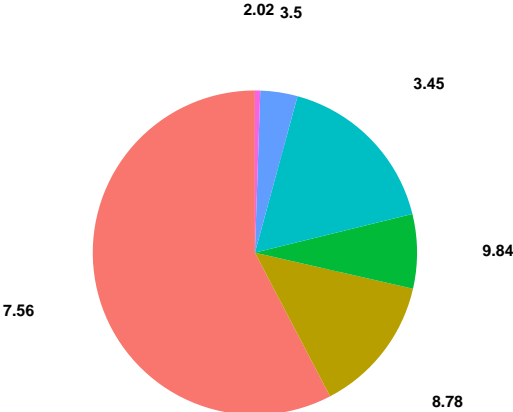


GO Analysis

module1



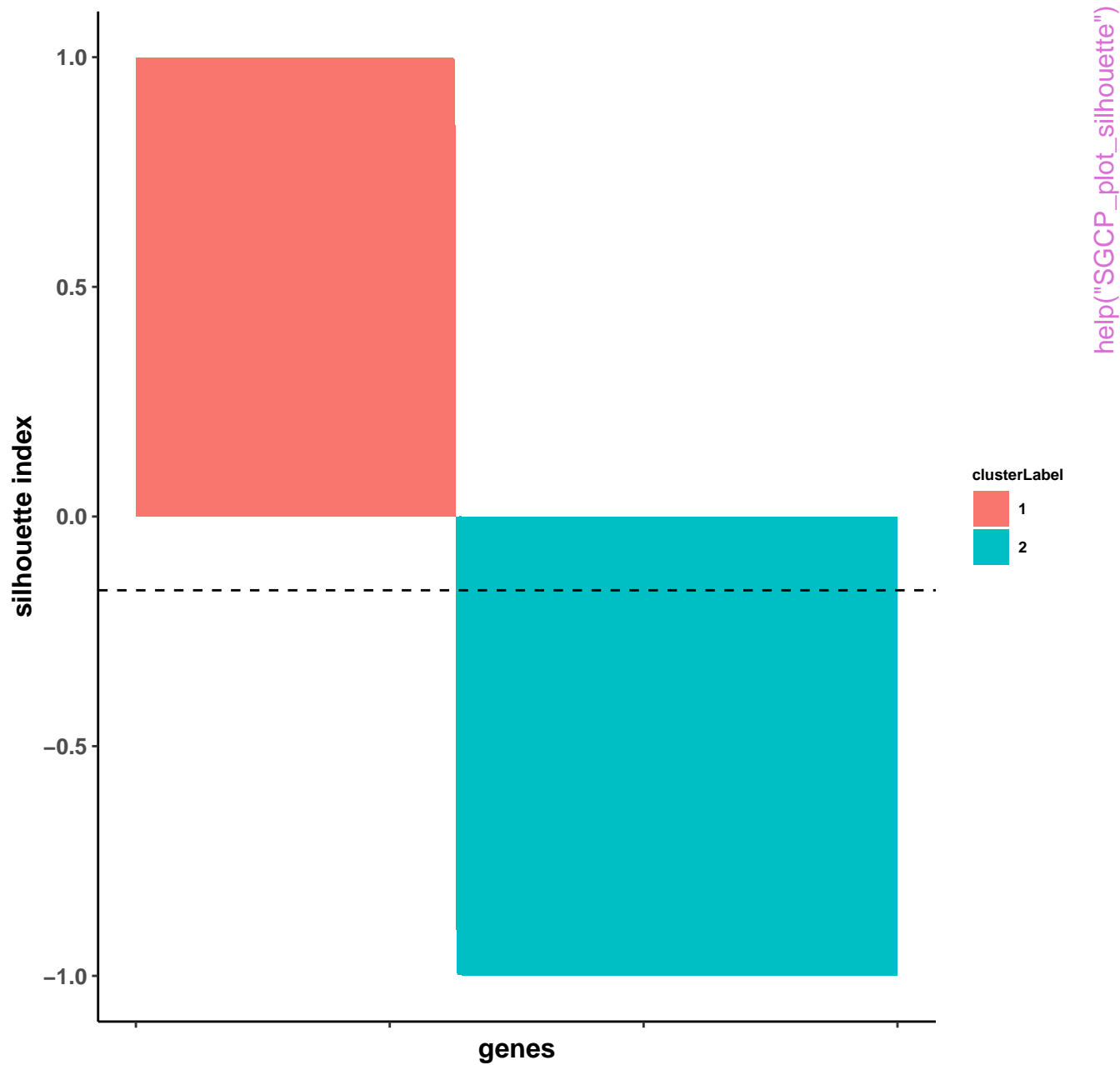
module2



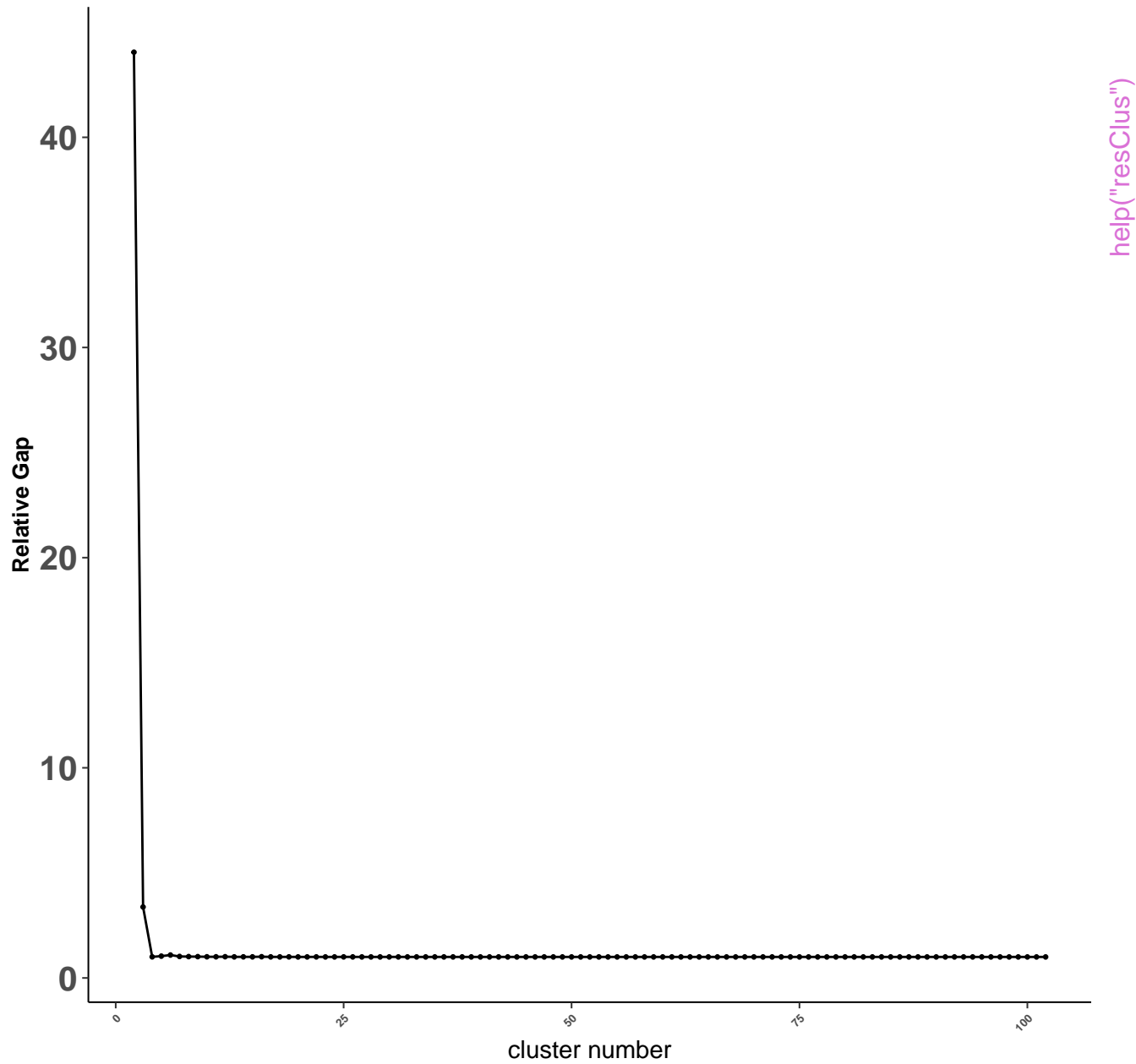
GeneOntology

	overBP		overMF		underCC
	overCC		underBP		underMF

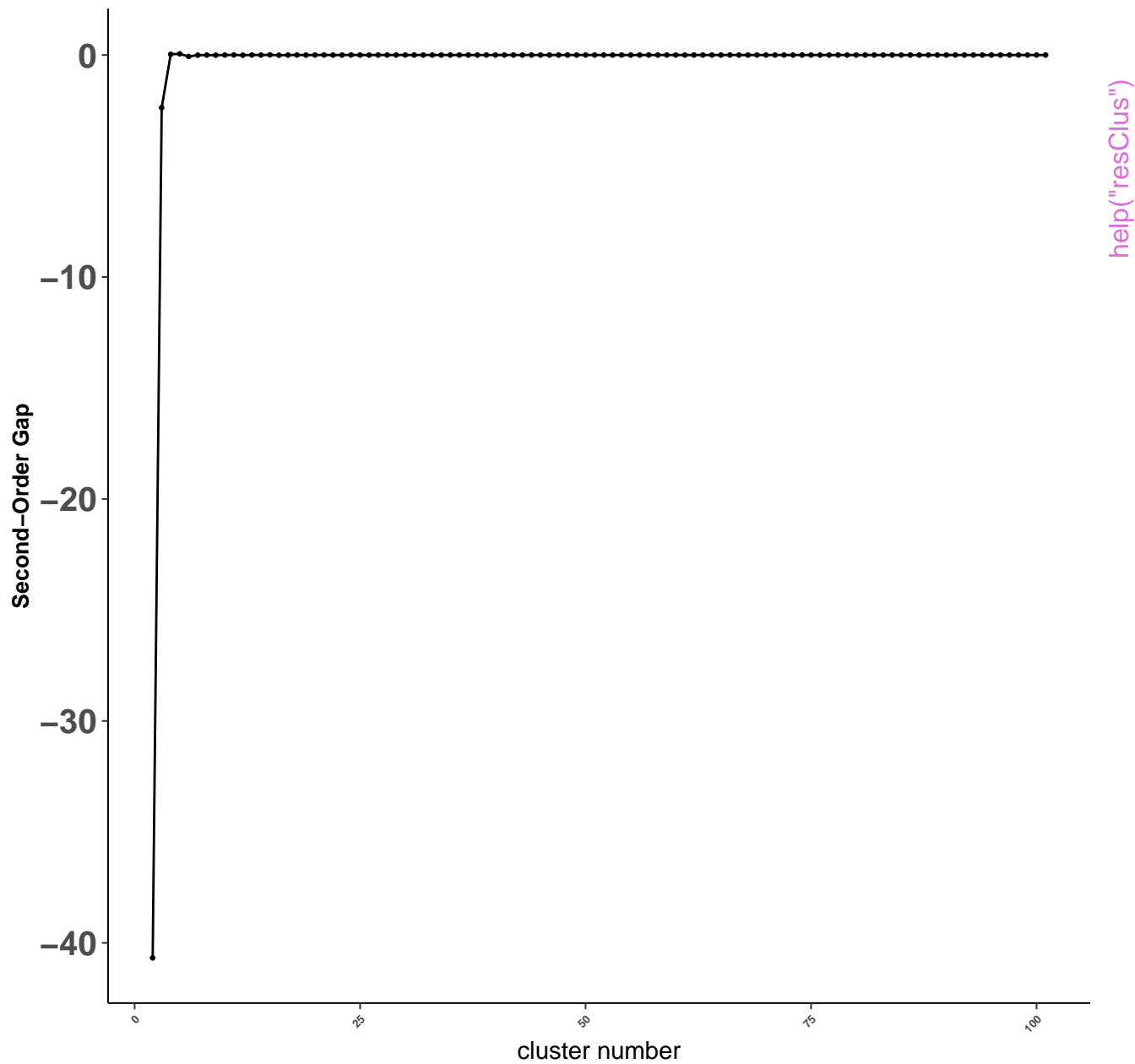
Gene Silhouette Index



Relative Gap Method



Second-Order Gap Method



help("resClus")

Additive Gap Method

