

- ☒ Euclidean
- ☐ Correlation

Method of hierachical clustering

- ☐ ward
- ☐ single
- ☒ complete
- ☐ average
- ☐ mcquitty
- ☐ median
- ☐ centroid

Re-order Rows?

- ☒ Yes
- ☐ No

Scaling?

- ☒ row
- ☐ column
- ☐ none

Cut the dendrogram into k groups, or h(eight)

- ☒ k
- ☐ h

Select k groups from the dendrogram



Constructing a heatmap from the gene list you uploaded in the Analysis Parameters tab. If you haven't uploaded a gene list, an example gene list of three genes will be used

