

Parameter Space Exploration for VisRSeq

Joseph Pober

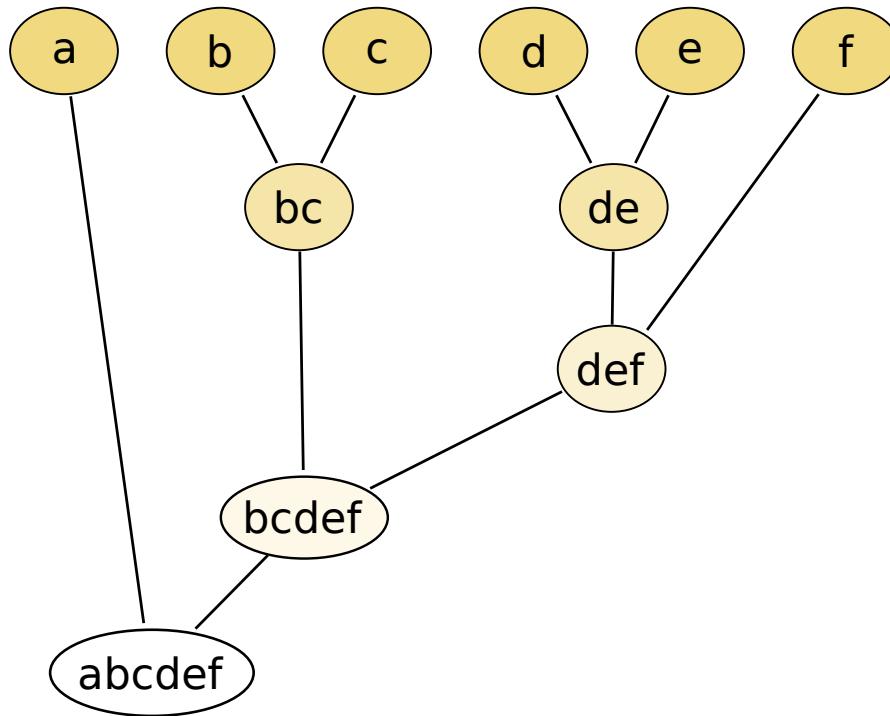
Content

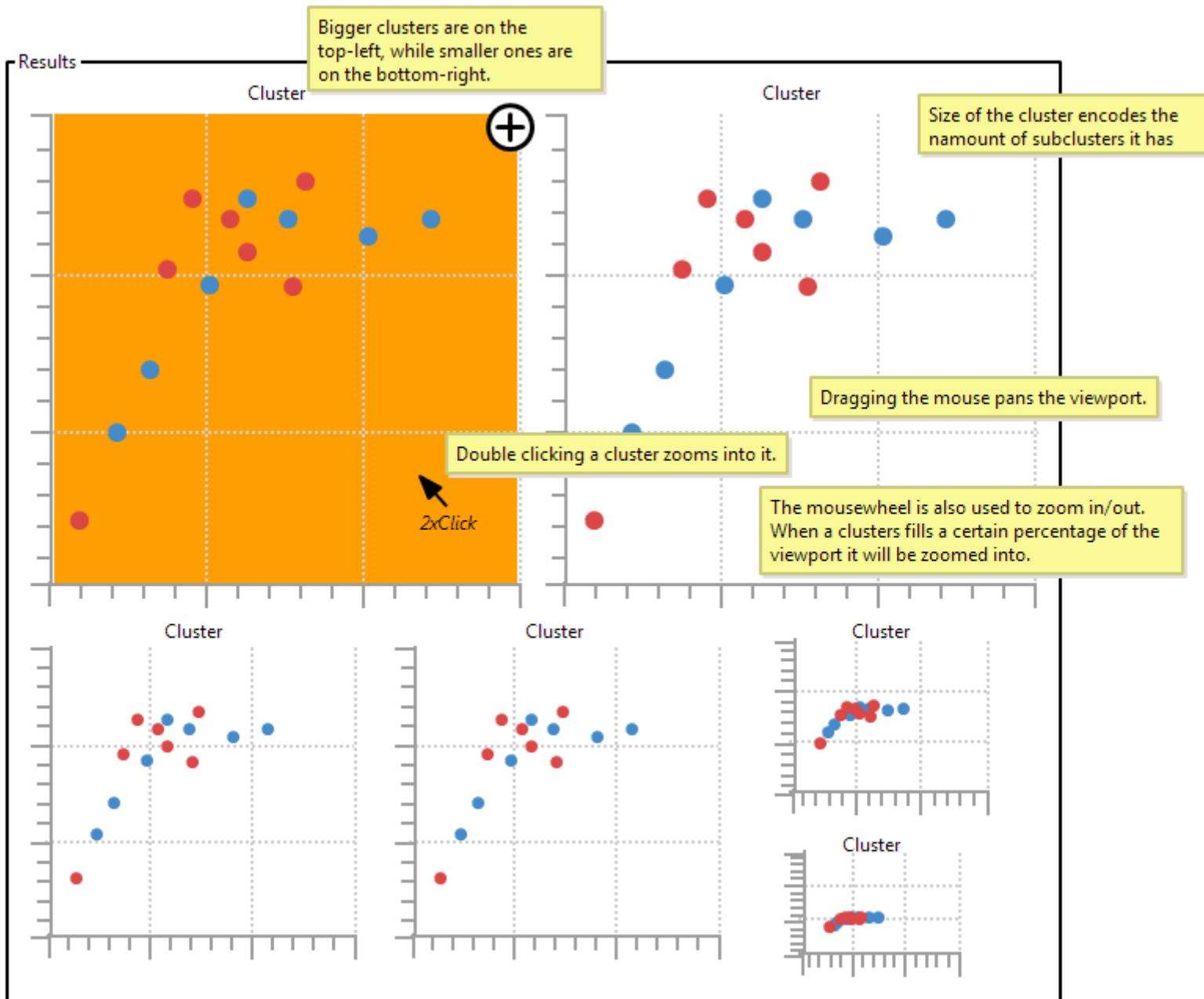
- Recap of project
- Visualization design
- Prototype

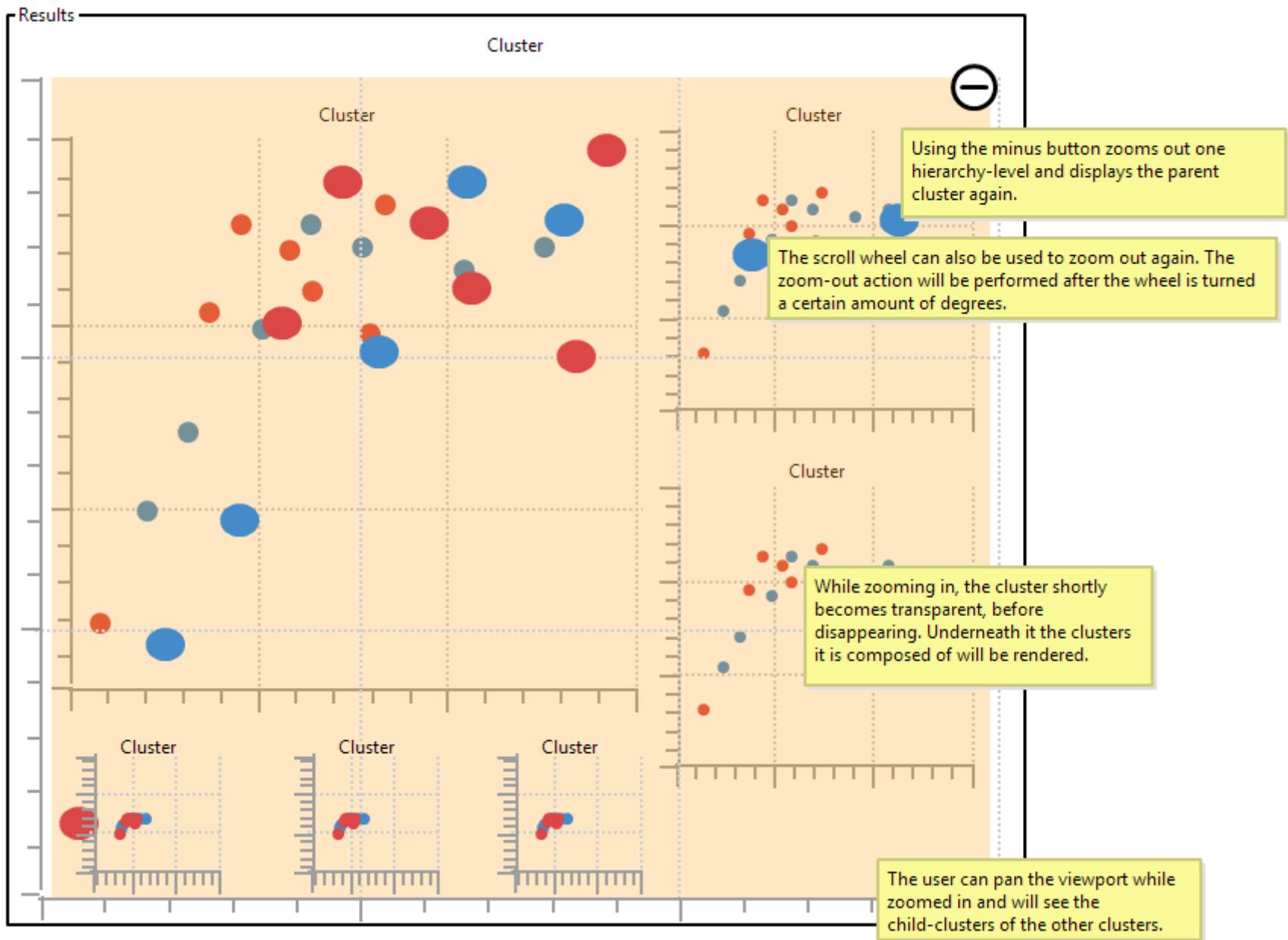
Project recap

- VisRSeq lets non-programmers use R
- VisRSeq allows only one plot at a time
- Parameter space exploration
- Clustering

Hierarchical clustering







Normal View

VisRSeq (v0.72.9)

File

Data workspace k-means [x]

Apps

Search [Clustering]

- Hierarchical Clustering (hclust)
- K-Means Clustering

Partitioning Around Medoids

Parameters

Input Code

k-means

variables:
 standardize variables

centers:
3

maximum number of c...
15

Plot Options

choice of plot:
K-means clustering

iterations:
10

start:
1

algorithm:
Hartigan-Wong

Output Column Name

cluster ID: clusterID

Console

Normal Run

Ranges

VisRSeq (v0.72.9)

File

Data workspace k-means [x]

Apps

Search [Clustering]

- Hierarchical Clustering (hclust)
- K-Means Clustering
- Partitioning Around Medoids

Parameters

Input Code

k-means

variables: Yes No

standardize variables: Yes No

centers: 3 3

maximum number of c... 15 15

Plot Options

Plot the within groups sums of square...

choice of plot: Recommended number of clusters u... K-means clustering

iterations: 10 10

start: 1 1

Hartigan-Wong
 Lloyd
 Forgy
 MacQueen

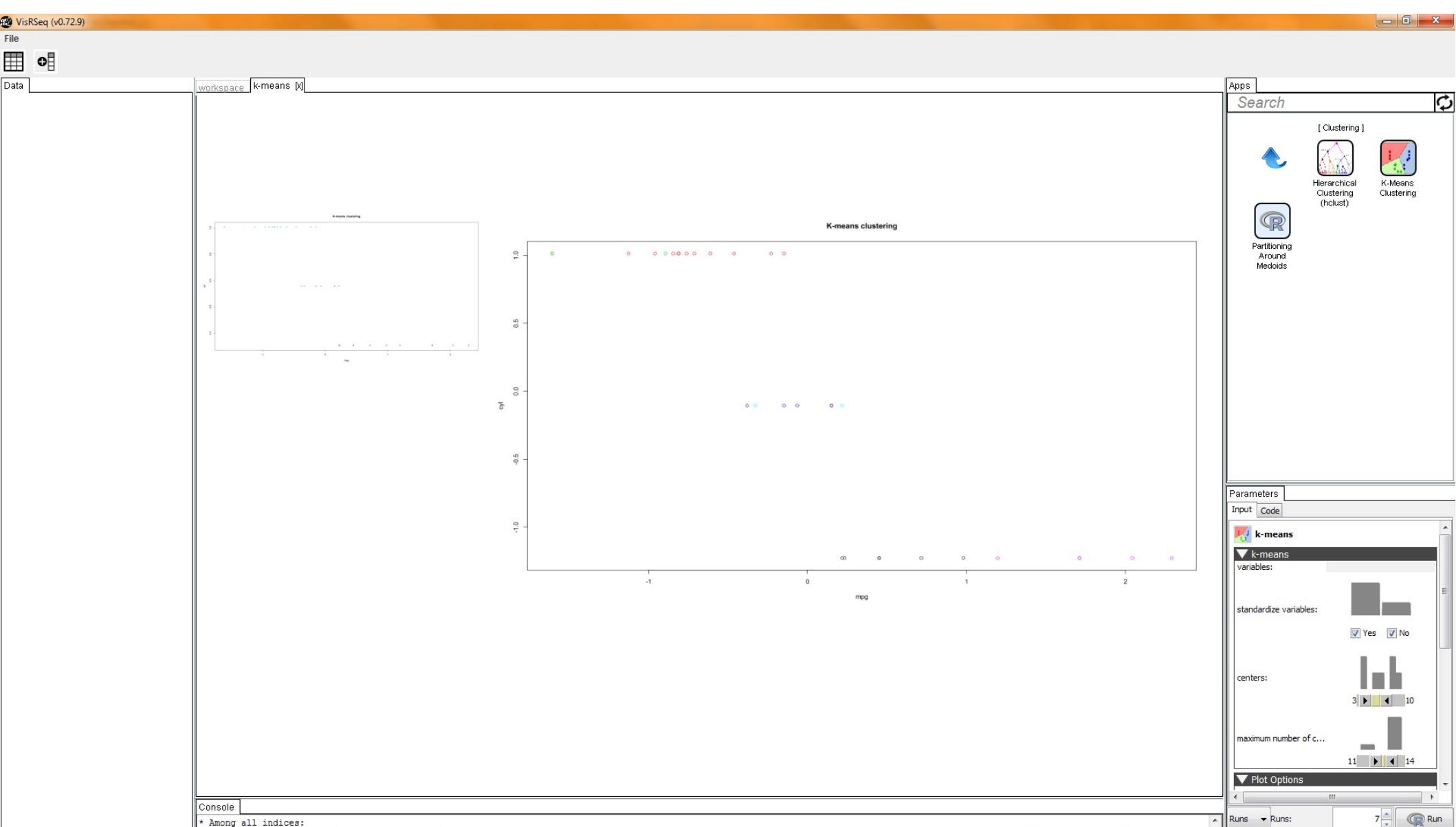
Output Column Name

cluster ID: clusterID

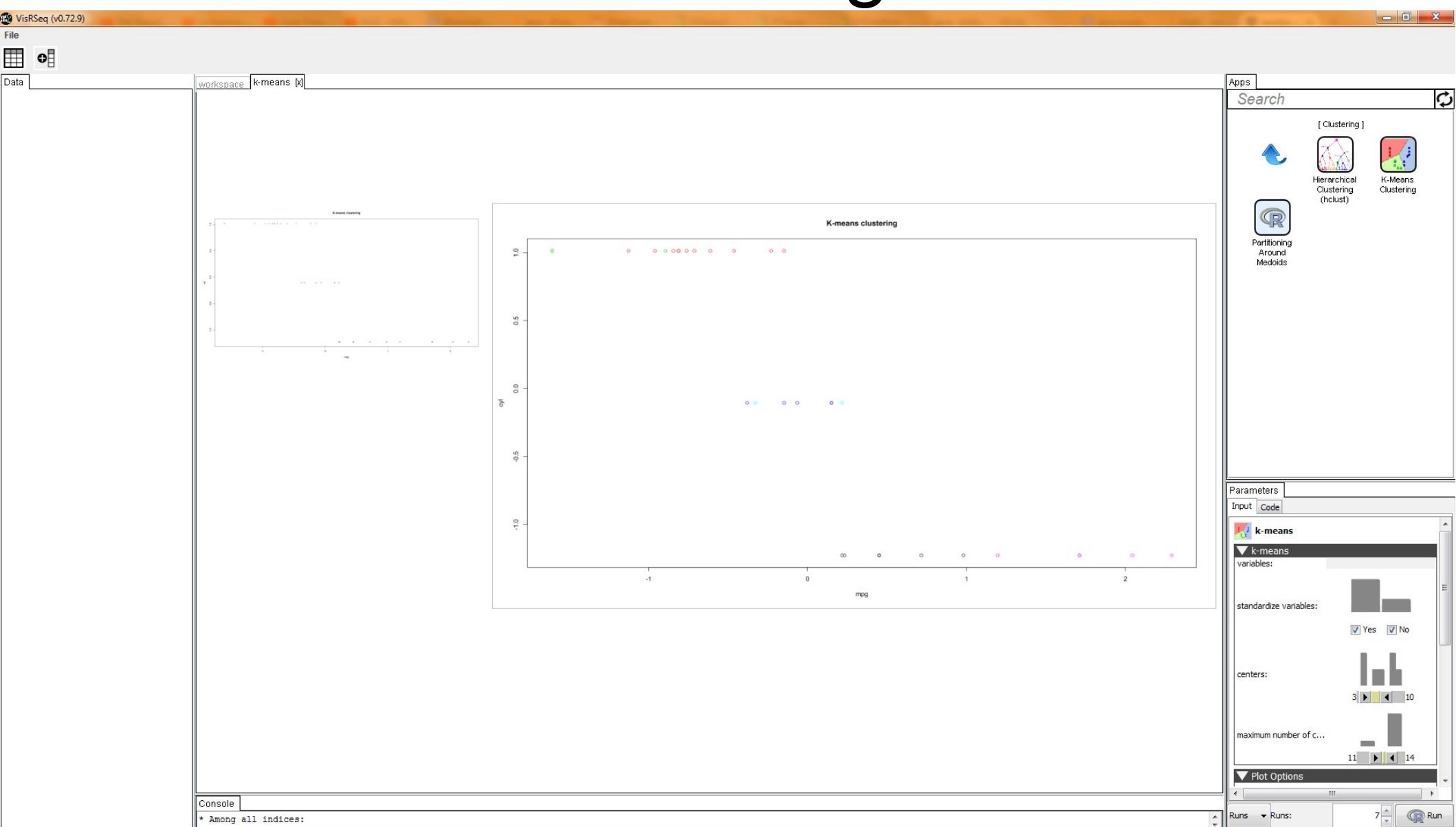
Runs: Runs: 2 Run

Console

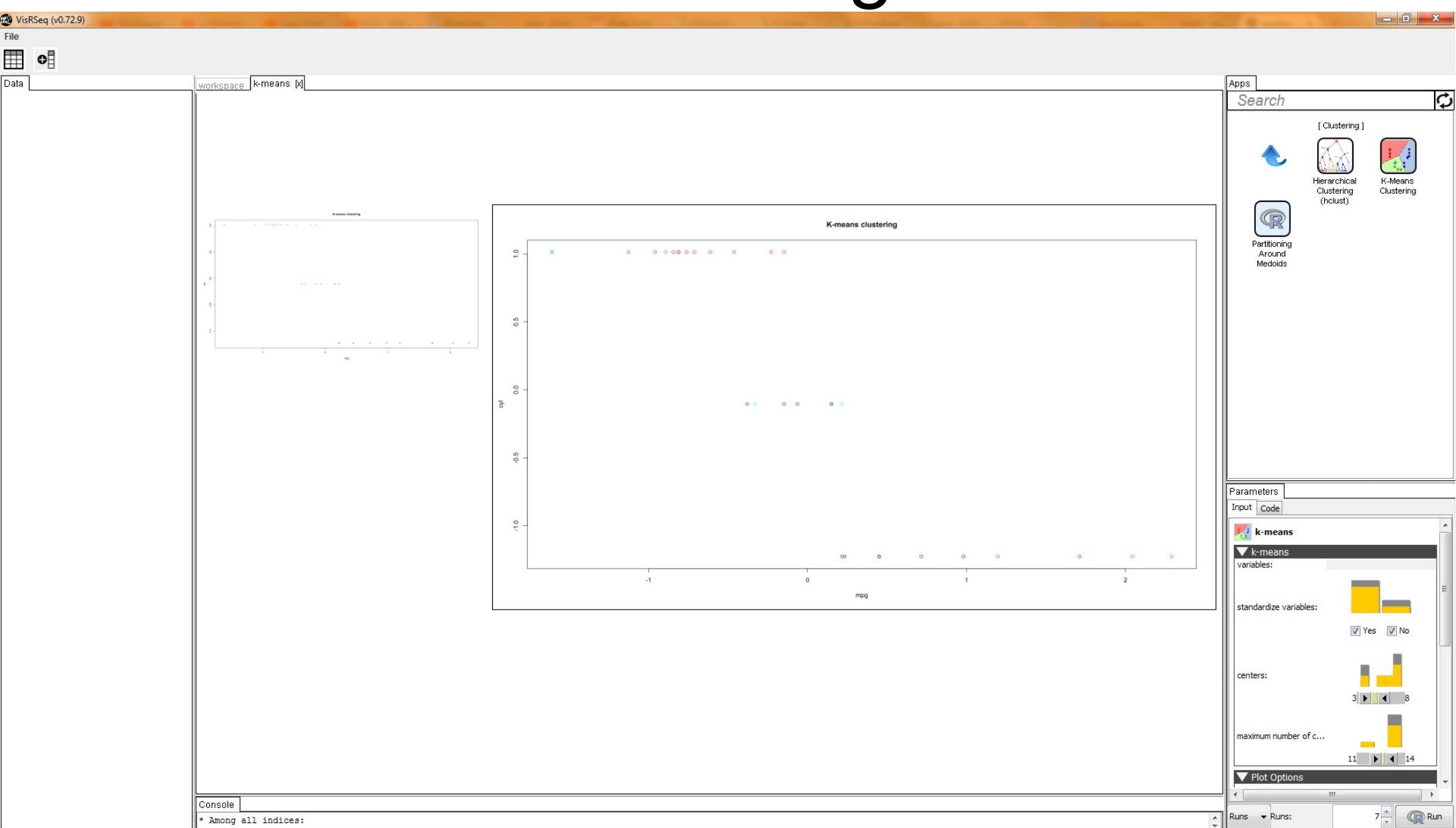
Distribution



Hovering



Selecting



Inspecting

VisRSeq (v0.72.9)

File

Data workspace k-means [x]

Back

Search [Clustering]

- Hierarchical Clustering (hclust)
- K-Means Clustering

Partitioning Around Medoids

Parameters Input Code

k-means

variables:

standardize variables: Yes No

centers: 3 ▶ 8

maximum number of c... 11 ▶ 14

Plot Options

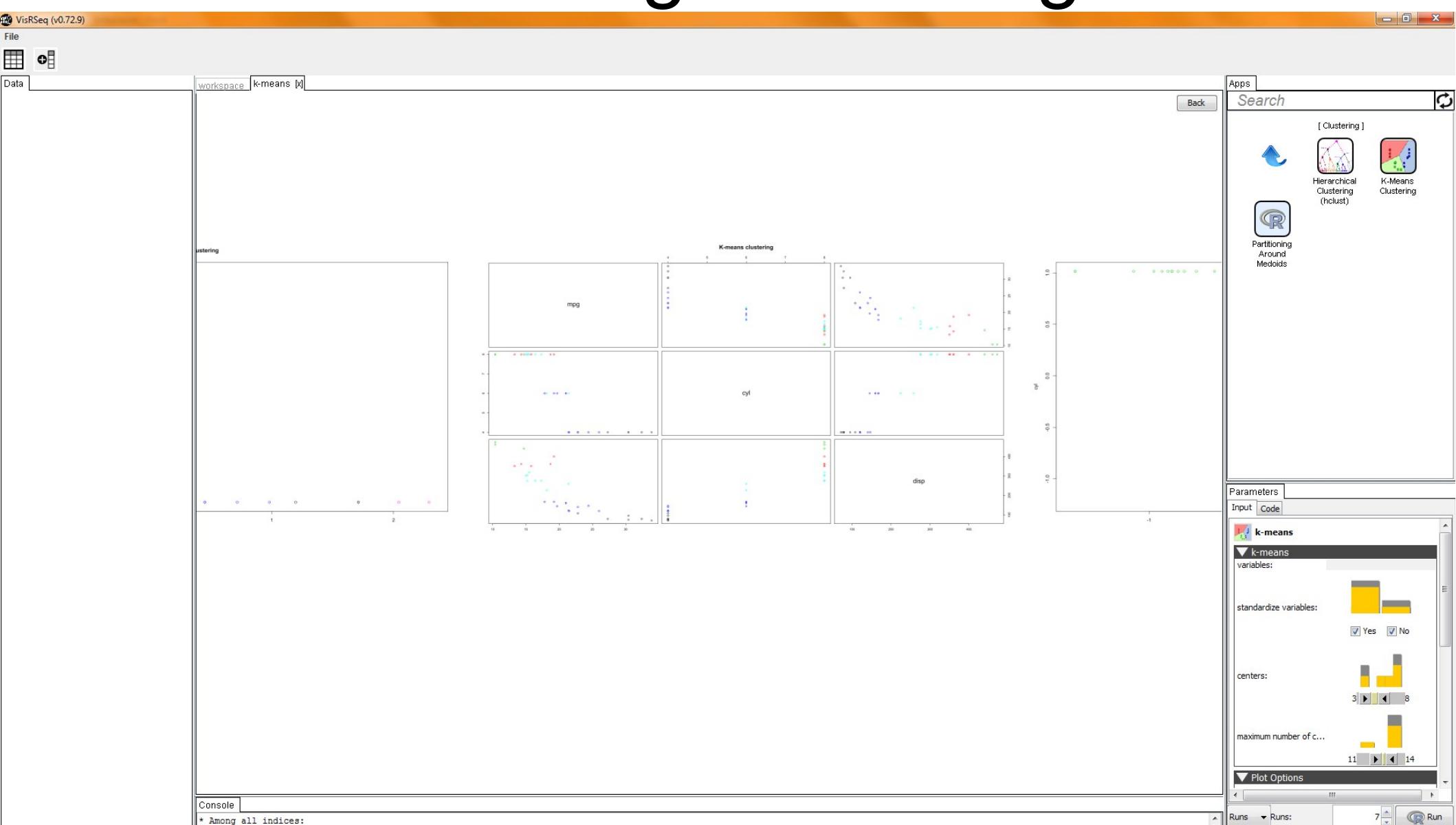
Console

* Among all indices:

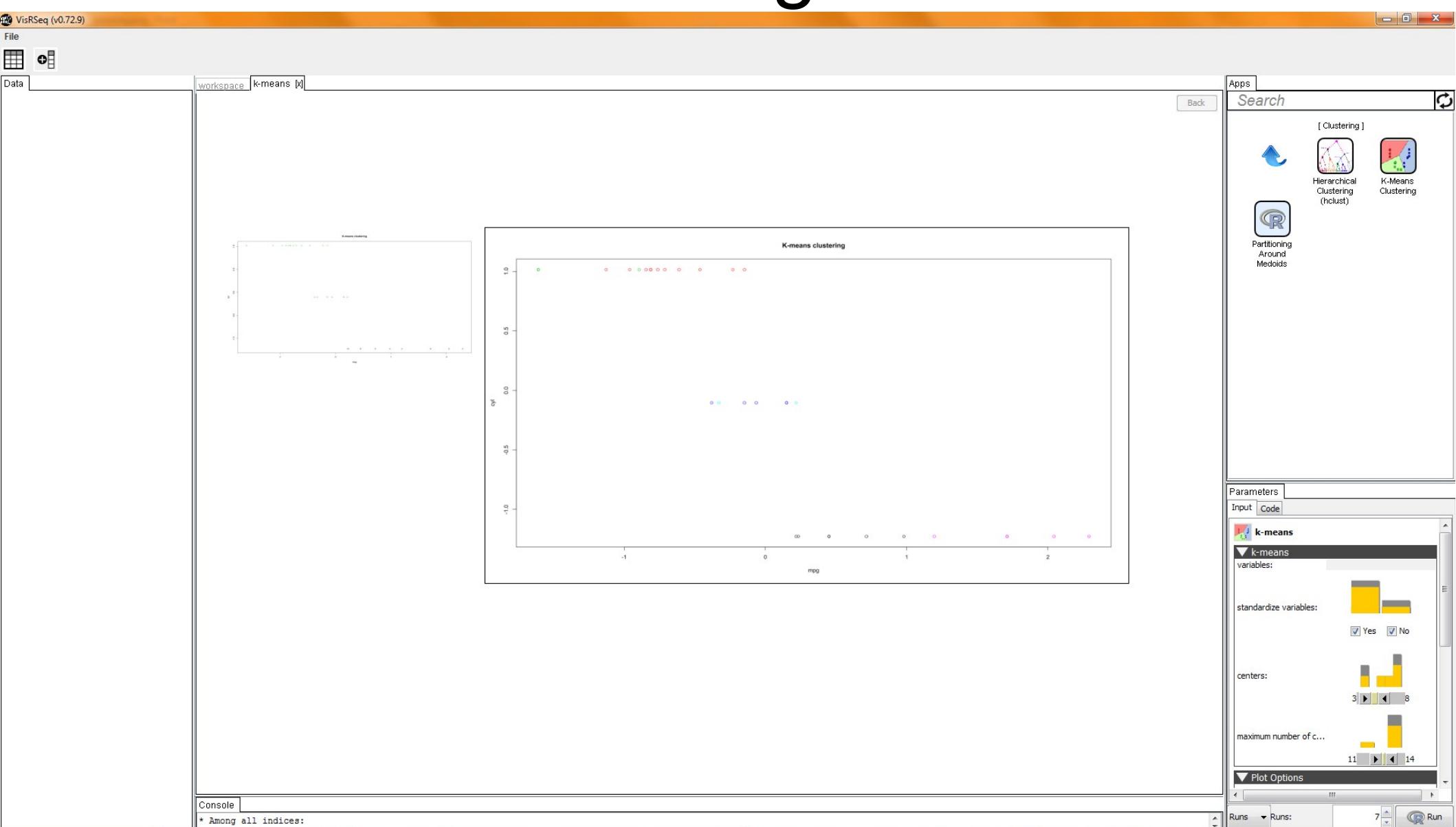
Runs Runs: 7 Run

The screenshot shows the VisRSeq software interface. At the top, there's a toolbar with icons for file operations like Open, Save, and Print. Below the toolbar, the main window has tabs for 'Data' and 'workspace' (which is currently selected), and a 'k-means [x]' tab. On the right side, there's a sidebar titled 'Search' with sections for 'Clustering' (Hierarchical Clustering (hclust) and K-Means Clustering) and 'Partitioning Around Medoids'. The central area displays four scatter plots labeled 'k-means clustering' with axes from 0 to 10. Below these plots is a 'k-means' parameter panel. It includes a 'variables:' section with a checkbox for 'standardize variables' (checked), a 'centers:' section with a slider set between 3 and 8, and a 'maximum number of c...' section with a slider set between 11 and 14. At the bottom, there's a 'Plot Options' section and a 'Console' tab showing the message '* Among all indices:'. A 'Runs' section at the very bottom shows 'Runs Runs: 7' and a 'Run' button.

Zooming & Panning



Zooming out



Challenges

- Visual comparison of plots
- Parallelization

Thank you for your attention!