

QlikView-OpenCPU Demo

OpenCPU and QlikView Demo Configuration

spa

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Installation and Setup

This brief document's purpose is to enable you to integrate R to the Cluster Analysis.qvw on demand using a cloud solution. More specifically, this paper will demonstrate how you can send data to and from other systems using the web as the pipeline. In this example, the Cluster Analysis.qvw will send data to Open CPU, Open CPU will process the data and send the data back to Cluster Analysis on demand. Open CPU is an initiative from Jeroem Ooms and has been implemented by Stat/Dev consulting in the Neitherlands.

These are steps you will go through:

- 1. Add a R script to the Open CPU store
- 2. Deploy the Cluster Analysis, qvw to QlikView server
- 3. Install & Configure the openCPU extension
- 4. Demonstrate QlikView and R on demand.

Adding an R Script to the Open CPU Store

OpenCPU.org

As I mention earlier, OpenCPU is being developed by Jeroen Ooms. The basic philosophy of OpenCPU is to have scientific computing in the cloud. Let's now publish our first set of R code to OpenCPU by clicking here.

Post a function

Publish Function



R Script to be published

Below is a function written in R that leverages the kmeans (klustering mean) function to cluster the data. The function takes as inputs a few parameters (e.g. AccountCount, AvgOrderSize, CustomerCnt) and returns back a new set of data that is clustered. Therefore, copy the code below and paste to OpenCPU link shown above. After pasting the code, press the publish function.

```
function (X, AccountCount, AvgOrderSize, CustomerCnt, Row, clusterGroup)
{

data = data.frame(X = X, AccountCount = AccountCount, AvgOrderSize = AvgOrderSize, CustomerCnt = CustomerCnt, Row = Row)

cleanData <- subset(data, select = -c(X))

fit <- kmeans(cleanData, clusterGroup)

aggregate(cleanData, by = list(fit$cluster), FUN = mean)

cleanData <- data.frame(cleanData, fit$cluster)

cleanData <- subset(cleanData, select = c(Row, fit.cluster))

cleanData <- merge(x = data, y = cleanData, by = "Row", all.x = TRUE)

cleanData <- subset(cleanData, select = -c(Row))

return(cleanData)
```

Post a function

```
function (X, AccountCount, AvgOrderSize, CustomerCnt, Row, clusterGroup)
{
    data = data.frame(X = X, AccountCount = AccountCount, AvgOrderSize = AvgOrderSize, CustomerCnt = CustomerCnt, Row = Row)
    cleanData <- subset(data, select = -c(X))
    fit <- kmeans(cleanData, clusterGroup)
    aggregate(cleanData, by = list(fitsColuster), FUN = mean)
    cleanData <- data.frame(cleanData, fitsColuster)
    cleanData <- subset(cleanData, select = -c(Row, fit.cluster))
    cleanData <- merge(x = data, y = cleanData, by = "Row", all.x = TRUE)
    cleanData <- subset(cleanData, select = -c(Row))
}
return(cleanData)
}</pre>
```

Publish Function

After pressing the publish button, you will be redirected to anew page as shown below:



```
{
    "object" : "x8ed46d775e",
    "graphs" : [],
    "files" : {}
}
```

Please copy down your object key (e.g. x8ed46d775e). This hashkey tells your extension where to find your function. You can think of the object as an unique street address for your function. Once you have your object key, you have finished setting up Open CPU.

Deploy the Cluster Analysis.qvw to QlikView Server

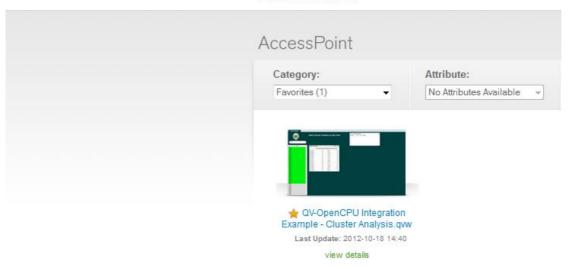
QV-OpenCPU Integration Example - Cluster Analysis.qvw

The zip file includes a QlikView document that needs to be deployed to the QlikView server. Therefore, copy and paste the application to the 'User Document'. For Windows 7, the default directory is C:\ProgramData\QlikTech\Documents. Once the file has been copied over, open up QlikView Access point using *Firefox**.

*Please note that version 1 of the extension only supports Firefox and Google Chrome.

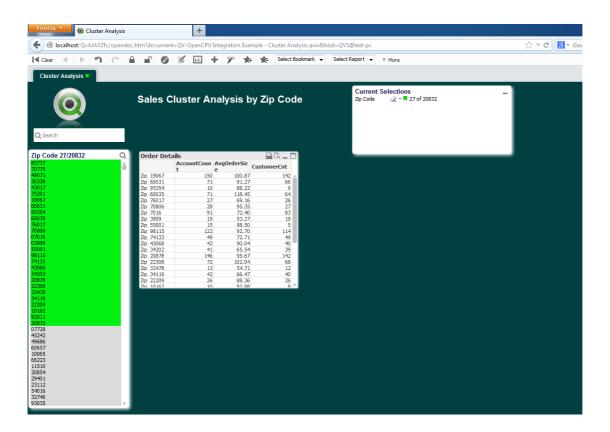


QlikView



When you open the QlikView file your application should look like the following:

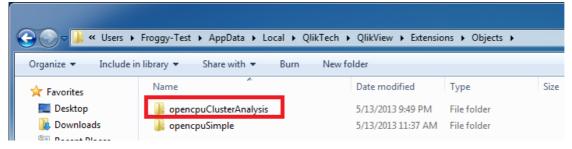




Install & Configure the openCPU Extension

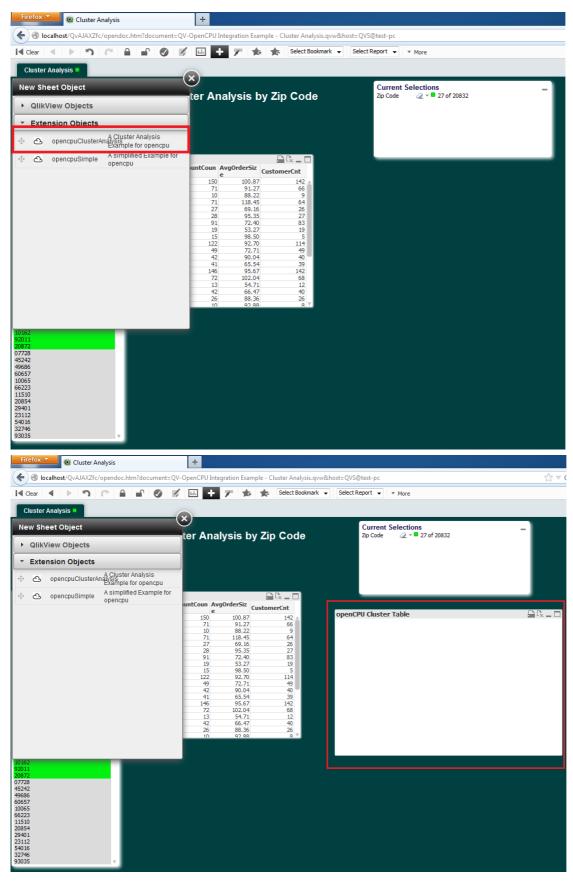
opencpuClusterAnalysis.qar

The opencpuClusterAnalysis extension will need to be installed on the server. If you double click the opencpuClusterAnalysis.qar to install the extension, it will only install it on the client's machine. Please ensure that the extension is located at:



Once the extension is correctly install, go to the QlikView Access point and launch the Cluster Analysis application which was describe earlier in the document. Inside the application, add a new sheet object and click on the extension objects. You will see the opencpuClusterAnalysis extensions. Drag the extension into the application.





Next right click on the openCPU Cluster Table object and click on properties. The properties

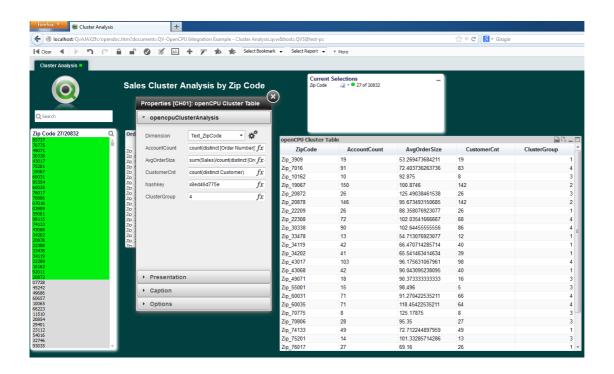


dialog will open with the following options:

Dimension Properties [CH01]: openCPU Cluster Table This field holds the dimension of which you will cluster against opencpuClusterAnalysis Input: Text ZipCode Dimension Text_ZipCode **AccountCount** AccountCount count(distinct [Order Number]) fxThis field represents the # of distinct order number against the dimension. AvgOrderSize sum(Sales)/count(distinct [Orc fx-Input: count(distinct [Order Number]) CustomerCnt count(distinct Customer) fxhashkey x8ed46d775e AvgOrderSize: fxClusterGroup This field represents the average order size per dimension. Input: sum(Sales)/count(distinct [Order Number1) CustomerCnt This field represents the # of distinct customer per dimension Input: count(distinct Customer) Presentation Hashkey Caption This field is required to link the extension Options with the function hosted in openCPU. Input: x8ed46d775e ClusterGroup This field informs the k mean cluster how to cluster the various dimensions. Input: 4

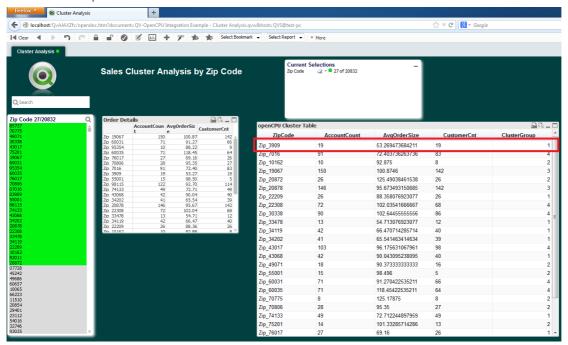
Once it is all set up, the extension will display a straight table.





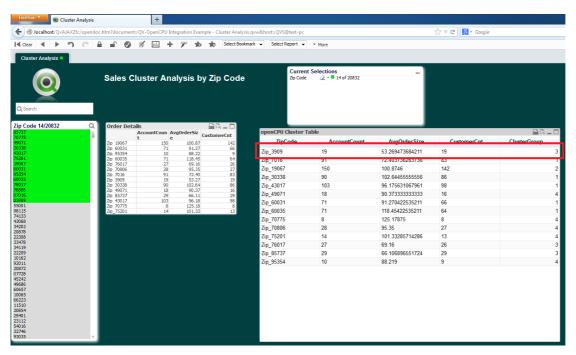
Demonstrate QlikView and R on demand

After the extension has been set up, click on a few zip codes and watch the straight table recluster each zip code on demand.



With 27 zip codes selected





With 14 Zip codes selected



Appendix

Below is the process workflow on the exchange between R, OpenCPU and the extension.

