**Results and Associated Methods**

**Enrich:**measureNames  
groupDataByRegion  
getDescription(name,raw)  
groupDataByRegion(query,descVector)

**Classify:**  
getIndices(percentTrain,dataset)  
setTrainAndTest(percentTrain,dataset)  
funKNN(p,set)  
compClassRes(res,actual)  
loadClassifyTest(sets)

**Normalize:**  
logNorm(sampleSet,nsam)  
loessNorm(dataSet)  
compileNormData(sampleSet)  
writeNormData(sampleSet,name)  
loadNormalizeWrite(PPVector)  
plotOldNorm(sampleSet,sample)  
plotOldNorms(sampleSet,samples)

[**PreProcess:**](../preProcess.R)  
[loadDatasets(query)](fp/loadDatasets.txt)  
loadDatasetsAndSub(query)  
convertDatasets(fullVector)  
fixColNames(allNames,setName)  
getDiagnoses(raw)  
applyDiagnoses(diagn,data)  
preProcess(query)  
preProcessShort(name,raw=NULL,sub=NULL)

**Main:**  
queryVector  
queryCF  
PPVector  
init\_workspace()  
runFullAnalysis(namesOfSets)  
runShortAnalysis(namesOfSets)