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# Bioconductor installs

source("https://bioconductor.org/biocLite.R")

biocLite ()

biocLite("flowCore")
biocLite('flowMerge')
biocLite("flowQ")
biocLite('flowType')

biocLite('RchyOptimyx')

biocLite('rrcov')
biocLite('codetools')
biocLite('foreach')


install.packages("ROCR")
install.packages("sfsmisc")
install.packages("GEOmap")

install.packages("car"); install.packages("RFOC")

library(flowCore)
library(flowDensity)
library(flowType)
library(RchyOptimyx);
library(ROCR)
library(sfsmisc)
library(GEOmap)
```