BIOS 7659 Homework 8

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# 1. Cell Type Composition

Read in the data:

baseDir1 <- "blood/plate1"  
targets1 <- read.metharray.sheet(baseDir1)

## [1] "blood/plate1/selected\_plate1.csv"

baseDir2 <- "blood/plate2"  
targets2 <- read.metharray.sheet(baseDir2)

## [1] "blood/plate2/selected\_plate2.csv"

targets <- rbind(targets1, targets2)  
rgSet <- read.metharray.exp(targets=targets,extended=T)