NDMG Comparison

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```
library(ggplot2)
library(grid)
library(gridExtra)
library(reshape2)
library(data.table)

##
## Attaching package: 'data.table'
## The following objects are masked from 'package:reshape2':
##
## dcast, melt
```

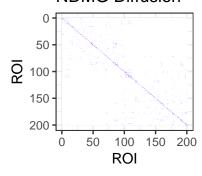
Loading

```
ndmg.dwi.adj = as.matrix(read.csv('./ndmg-dwi.csv', sep=" ", header=FALSE))
derek.dwi.adj = as.matrix(read.csv('./derek-dwi.csv', sep=" ", header=FALSE))
ndmg.func.adj = as.matrix(read.csv('./ndmg-func.csv', sep=" ", header=FALSE))
colnames(ndmg.dwi.adj) <- as.character(seq(1, dim(ndmg.dwi.adj)[1]))
colnames(derek.dwi.adj) <- as.character(seq(1, dim(derek.dwi.adj)[1]))
colnames(ndmg.func.adj) <- as.character(seq(1, dim(ndmg.func.adj)[1]))
ndmg.dwi.dat <- melt(ndmg.dwi.adj)
derek.dwi.dat <- melt(derek.dwi.adj)
ndmg.func.dat <- melt(ndmg.func.adj)</pre>
```

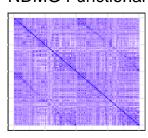
Plotting

```
titles=list("NDMG Diffusion", "Probabilistic Diffusion", "NDMG Functional")
mtxs = list(ndmg.dwi.dat, derek.dwi.dat, ndmg.func.dat)
plots = lapply(1:length(mtxs), function(i) {
    mtx = mtxs[[i]]
    ggplot(mtx, aes(x=Var1, y=Var2, fill=value)) +
        geom_tile() +
        scale_y_reverse() +
        xlab("ROI") +
        ylab("ROI") +
        ggtitle(titles[[i]]) +
        scale_fill_gradient(name="Connectivity", low="white", high="blue") +
        theme_bw()
})
sim_leg <- g_legend(plots[[1]])
plots <- lapply(1:length(plots), function(i) {</pre>
```

NDMG Diffusion



NDMG Functional



Probabilistic Diffusion

