

Project.R

Divy

Sun Dec 28 20:35:54 2014

```
"Hello World This is gene mapping project"
```

```
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```

```
"Pleas"
```

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```

```
dat <- matrix( 0, 8, 4 )  
colnames( dat ) <- c("CCStatus", "XRAY", "BINARYOB", "weights")  
dat[5:8,1] <- 1  
dat[c(3,4,7,8),2] <- 1  
dat[c(2,4,6,8),3] <- 1
```