Subchallenge 1 Report

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load libraries

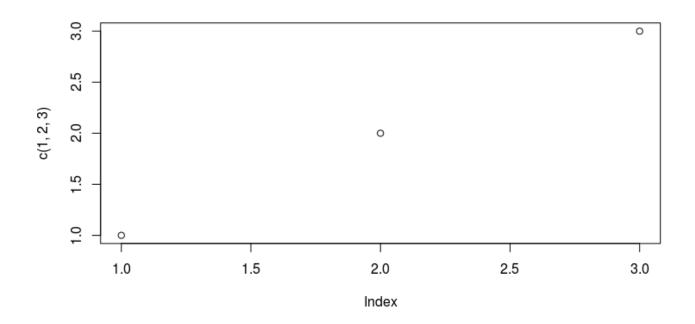
```
library(knitr)
```

set global options

download data

calculate genesets

```
[1] 2374 1
```



generate heatmaps
perform pathway analysis