

# Bimodality Index

Kevin R. Coombes

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## 1 Simulated Data

We simulate a dataset.

```
> set.seed(564684)
> nSamples <- 30
> nGenes <- 3000
> dataset <- matrix(rnorm(nSamples*nGenes), ncol=nSamples, nrow=nGenes)
> dimnames(dataset) <- list(paste("G", 1:nGenes, sep=''),
+                             paste("S", 1:nSamples, sep=''))
> library(BimodalIndex)
```

## 2 Appendix

This analysis was performed in the following directory:

```
> getwd()
[1] "/tmp/Rtmp5TNBOI/Rbuild6eda7c862207/BimodalIndex/vignettes"
```

This analysis was performed in the following software environment:

```
> sessionInfo()

R version 3.1.2 Patched (2015-02-05 r67741)
Platform: x86_64-unknown-linux-gnu (64-bit)
Running under: Debian GNU/Linux 8 (jessie)

locale:
 [1] LC_CTYPE=en_US.UTF-8      LC_NUMERIC=C              LC_TIME=en_US.UTF-8
 [4] LC_COLLATE=C             LC_MONETARY=en_US.UTF-8  LC_MESSAGES=en_US.UTF-8
```

```
[7] LC_PAPER=en_US.UTF-8      LC_NAME=C                LC_ADDRESS=C
[10] LC_TELEPHONE=C             LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C
```

attached base packages:

```
[1] stats      graphics  grDevices  utils      datasets  methods    base
```

other attached packages:

```
[1] BimodalIndex_1.1.1
```

loaded via a namespace (and not attached):

```
[1] mclust_4.4      oompaBase_3.1.0 tools_3.1.2
```