# narrowpeaksDiff: Analysis of differential variation across ChIP-seq experiments

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January, 2013

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## 1 Rationale

Under development.

## 2 Details

This document was written using:

```
R> sessionInfo()

R version 3.0.0 (2013-04-03)
Platform: x86_64-apple-darwin10.8.0 (64-bit)

locale:
[1] C

attached base packages:
[1] parallel  splines   stats     graphics  grDevices
[6] utils     datasets  methods   base

other attached packages:
[1] rtracklayer_1.20.0   CSAR_1.12.0
[3] GenomicRanges_1.12.1 IRanges_1.18.0
[5] BiocGenerics_0.6.0   NarrowPeaks_1.4.0

loaded via a namespace (and not attached):
```

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```
 [1] BSgenome_1.28.0   Biostrings_2.28.0 Matrix_1.0-12
 [4] RCurl_1.95-4.1    Rsamtools_1.12.0  XML_3.96-1.1
 [7] bitops_1.0-5      fda_2.3.2         grid_3.0.0
[10] lattice_0.20-15   stats4_3.0.0      tools_3.0.0
[13] zlibbioc_1.6.0
```