

narrowpeaksDiff: Analysis of differential variation across ChIP-seq experiments

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1 Rationale

Under development.

2 Details

This document was written using:

```
R> sessionInfo()
```

```
R version 3.0.2 (2013-09-25)
```

```
Platform: i386-w64-mingw32/i386 (32-bit)
```

```
locale:
```

```
[1] LC_COLLATE=C
```

```
[2] LC_CTYPE=English_United States.1252
```

```
[3] LC_MONETARY=English_United States.1252
```

```
[4] LC_NUMERIC=C
```

```
[5] LC_TIME=English_United States.1252
```

```
attached base packages:
```

```
[1] parallel splines stats graphics grDevices
```

```
[6] utils datasets methods base
```

```
other attached packages:
```

```
[1] rtracklayer_1.22.0 CSAR_1.14.0
```

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```
[3] GenomicRanges_1.14.0 XVector_0.2.0
[5] IRanges_1.20.0      BiocGenerics_0.8.0
[7] NarrowPeaks_1.6.0
```

loaded via a namespace (and not attached):

```
[1] BSgenome_1.30.0   Biostrings_2.30.0 Matrix_1.0-14
[4] RCurl_1.95-4.1    Rsamtools_1.14.0  XML_3.98-1.1
[7] bitops_1.0-6      fda_2.3.8         grid_3.0.2
[10] lattice_0.20-24   stats4_3.0.2      tools_3.0.2
[13] zlibbioc_1.8.0
```