Analyzing ChIP-chip data using Bioconductor

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This is only a placeholder. The real vignette has been pre-created since it takes quite some time to process.

Software versions

This tutorial was generated using the following package versions:

- R version 2.9.0 (2009-04-17), x86_64-unknown-linux-gnu
- $\bullet \ \ Locale: \ LC_CTYPE=en_US; LC_NUMERIC=C; LC_TIME=en_US; LC_COLLATE=en_US; LC_MONETARY=C; LC_MESS = locale: \ LC_CTYPE=en_US; LC_NUMERIC=C; LC_TIME=en_US; LC_COLLATE=en_US; LC_MONETARY=C; LC_MESS = locale: \ lo$
- Base packages: base, datasets, graphics, grDevices, methods, stats, tools, utils