# Package 'RnBeads.mm9'

### April 17, 2025

Title RnBeads.mm9

Description Automatically generated RnBeads annotation package for the assembly mm9. Author RnBeadsAnnotationCreator Maintainer RnBeadsAnnotationCreator <rnbeads@mpi-inf.mpg.de> Date 2021-11-21 License GPL-3 Version 1.41.0 Depends R (>= 3.0.0), GenomicRanges Suggests RnBeads NeedsCompilation no git\_url https://git.bioconductor.org/packages/RnBeads.mm9 git\_branch devel git\_last\_commit\_f7a3f5d git\_last\_commit\_date 2025-04-15 Repository Bioconductor 3.22 Date/Publication 2025-04-17

## Contents

mm9
-----

#### Index

mm9

Annotation tables for mm9

#### Description

Scaffold of annotation tables for the mm9 assembly. This structure is automatically loaded upon initialization of the annotation, that is, by the first valid call to any of the following functions: rnb.get.assemblies, rnb.get.chromosomes, rnb.get.annotation, rnb.set.annotation, rnb.get.mapping, rnb.annotation.size. Adding an annotation amounts to attaching its table(s) and mapping structures to this scaffold.

3

#### Format

- list of up to six elements "GENOME", "CHROMOSOMES", "regions", "sites", "controls" and "mappings". These elements are described below.
- "GENOME" Name of the Bioconductor package that contains the genomic sequence for this genome assembly.
- "CHROMOSOMES" Supported chromosomes for this genome assembly. The elements of this character vector follow the Ensembl convention ("1", "2", ...), and the names of this vector the convention of the UCSC Genome Browser ("chr1", "chr2", ...).
- "regions" list of NULLs; the names of the elements correspond to the built-in region annotation tables. Once the default annotations are loaded, the attribute "builtin" is a logical vector storing, for each region annotation, whether it is the default (built-in) or custom.
- "sites" list of NULLs; the names of the elements correspond to the site and probe annotation tables.
- "controls" list of NULLs; the names of the elements correspond to the control probe annotation tables. The attribute "sites" is a character vector pointing to the site annotation that encompasses the respective control probes.
- "mappings" list of NULLs; the names of the elements correspond to the built-in region annotation tables.

#### Author(s)

**RnBeads Annotation Creator** 

# Index

\* datasets mm9, 1

mm9, 1

regions (mm9), 1
rnb.annotation.size, 1
rnb.get.annotation, 1
rnb.get.assemblies, 1
rnb.get.chromosomes, 1
rnb.get.mapping, 1
rnb.set.annotation, 1

sites(mm9), 1