

# Package ‘pcaGoPromoter.Mm.mm9’

March 25, 2013

**Version** 0.99.3

**Date** 2011-12-07

**Title** pcaGoPromoter.Mm.mm9 is a data package used by pcaGoPromoter

**Author** Morten Hansen, <mhansen@sund.ku.dk>

**Maintainer** Morten Hansen, <mhansen@sund.ku.dk>

**Description** Contains the data for transcription factors analysis of organism Mus musculus with package pcaGoPromoter

**biocViews** AnnotationData, Mus\_musculus

**LazyLoad** yes

**License** GPL (>= 2)

## R topics documented:

pcaGoPromoter.Mm.mm9 . . . . . 1

**Index** 2

---

pcaGoPromoter.Mm.mm9

*The pcaGoPromoter.Mm.mm9 data set is for use with package pcaGoPromoters*

---

### Description

This data package contains two lists, thresholds and promotersRefseqs, which is used by the function primo in the software package ‘pcaGoPromoter’. See manual pages for function primoData in packages ‘pcaGoPromoter’ for description of the two variables.

# Index

## \*Topic datasets

pcaGoPromoter.Mm.mm9, [1](#)

pcaGoPromoter.Mm.mm9, [1](#)

promotersRefseqs

(pcaGoPromoter.Mm.mm9), [1](#)

thresholds (pcaGoPromoter.Mm.mm9), [1](#)