

# Package ‘pcaGoPromoter.Mm.mm9’

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

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

**biocViews** Mus\_musculus\_Data

**LazyLoad** yes

**License** GPL (>= 2)

**git\_url** <https://git.bioconductor.org/packages/pcaGoPromoter.Mm.mm9>

**git\_branch** RELEASE\_3\_10

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

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## 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.

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