

Package ‘pcaGoPromoter.Hs.hg19’

October 17, 2019

Version 1.20.0

Date 2011-12-07

Title pcaGoPromoter.Hs.hg19 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 Homo Sapiens with package pcaGoPromoter

biocViews Homo_sapiens_Data

LazyLoad yes

License GPL (>= 2)

git_url <https://git.bioconductor.org/packages/pcaGoPromoter.Hs.hg19>

git_branch RELEASE_3_9

git_last_commit ec1cadf

git_last_commit_date 2019-05-02

Date/Publication 2019-10-17

R topics documented:

pcaGoPromoter.Hs.hg19	1
Index	2

pcaGoPromoter.Hs.hg19 *The pcaGoPromoter.Hs.hg19 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.Hs.hg19, [1](#)

pcaGoPromoter.Hs.hg19, [1](#)

promotersRefseqs

(pcaGoPromoter.Hs.hg19), [1](#)

thresholds (pcaGoPromoter.Hs.hg19), [1](#)