

Package ‘vtpnet’

February 7, 2023

Title variant-transcription factor-phenotype networks

Version 0.38.0

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Description variant-transcription factor-phenotype networks, inspired by Maurano et al., Science (2012), PMID 22955828

Suggests MotifDb, VariantAnnotation, Rgraphviz

Depends R (>= 3.0.0), graph, GenomicRanges, gwascats, doParallel, foreach

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LazyLoad yes

biocViews Network

git_url <https://git.bioconductor.org/packages/vtpnet>

git_branch RELEASE_3_16

git_last_commit a0b6d4b

git_last_commit_date 2022-11-01

Date/Publication 2023-02-06

R topics documented:

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vtpnet-package *variant-transcription factor-phenotype networks*

Description

variant-transcription factor-phenotype networks, inspired by Maurano et al., Science (2012), PMID 22955828

Details

Package: vtpnet
Version: 0.5.3
Suggests: MotifDb, VariantAnnotation, Rgraphviz
Depends: R (>= 3.0.0), graph, GenomicRanges, gwascats, foreach
License: Artistic-2.0
LazyLoad: yes
biocViews: Network
Built: R 3.1.1; ; 2014-10-08 14:42:02 UTC; unix

Not for release at this time (Oct 2014).

Author(s)

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