

Package ‘vtpnet’

August 18, 2022

Title variant-transcription factor-phenotype networks

Version 0.37.0

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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 master

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R topics documented:

vtpnet-package 2

Index 3

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, doParallel, 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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Index

* **package**

vtpnet-package, [2](#)

cancerMap (vtpnet-package), [2](#)

filterGWASbyMap (vtpnet-package), [2](#)

getMtags (vtpnet-package), [2](#)

getOneHits (vtpnet-package), [2](#)

gwMatchPWM (vtpnet-package), [2](#)

matrc (vtpnet-package), [2](#)

maurGWAS (vtpnet-package), [2](#)

MdbSearchToDisk (vtpnet-package), [2](#)

pax4 (vtpnet-package), [2](#)

pax4_75 (vtpnet-package), [2](#)

pax4_85 (vtpnet-package), [2](#)

rng2facHits (vtpnet-package), [2](#)

runfimo (vtpnet-package), [2](#)

setupGWPM (vtpnet-package), [2](#)

vtpnet (vtpnet-package), [2](#)

vtpnet-package, [2](#)