

Package ‘vtpnet’

October 17, 2024

Title variant-transcription factor-phenotype networks

Version 0.44.0

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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, gwascat, doParallel,
foreach

Maintainer VJ Carey <stvjc@channing.harvard.edu>

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LazyLoad yes

biocViews Network

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

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Contents

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, gwascat, 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](#)