

Package ‘vtpnet’

October 17, 2020

Title variant-transcription factor-phenotype networks
Version 0.28.0
Author VJ Carey <stvjc@channing.harvard.edu>
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>
License Artistic-2.0
LazyLoad yes
biocViews Network
git_url <https://git.bioconductor.org/packages/vtpnet>
git_branch RELEASE_3_11
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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)

VJ Carey <stvjc@channing.harvard.edu>

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

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