

Package ‘vtpnet’

April 10, 2022

Title variant-transcription factor-phenotype networks

Version 0.34.0

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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

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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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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