

Package ‘vtpnet’

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Title variant-transcription factor-phenotype networks

Version 0.14.0

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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

NeedsCompilation no

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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, 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).

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