

Package: vtpnet (via r-universe)

July 3, 2024

Title variant-transcription factor-phenotype networks
Version 0.45.0
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, gwascats, doParallel, foreach
Maintainer VJ Carey <stvjc@channing.harvard.edu>
License Artistic-2.0
LazyLoad yes
biocViews Network
Repository <https://bioc.r-universe.dev>
RemoteUrl <https://github.com/bioc/vtpnet>
RemoteRef HEAD
RemoteSha 55ee11c3bf6c7b6b5be11444a8f88c1d4cc75e03

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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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