Package: vtpnet (via r-universe)

October 18, 2026

Title variant-transcription factor-phenotype networks

Version 0.53.0

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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

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License Artistic-2.0

LazyLoad yes

biocViews Network

Config/pak/sysreqs make libbz2-dev libicu-dev liblzma-dev libpng-dev libxml2-dev libssl-dev libx11-dev xz-utils zlib1g-dev

Repository https://bioc.r-universe.dev

Date/Publication 2026-04-28 12:37:45 UTC

RemoteUrl https://github.com/bioc/vtpnet

RemoteRef HEAD

RemoteSha 1c799737bfa4bd6f5740d4c2e204f1f1d53da711

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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, 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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