

# Package: vtpnet (via r-universe)

June 19, 2026

**Title** variant-transcription factor-phenotype networks

**Version** 0.52.0

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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>

**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-release.r-universe.dev>

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

**RemoteUrl** <https://github.com/bioc/vtpnet>

**RemoteRef** RELEASE\_3\_23

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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).

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