

# Package ‘vtpnet’

July 8, 2025

**Title** variant-transcription factor-phenotype networks  
**Version** 0.49.0  
**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  
**git\_url** <https://git.bioconductor.org/packages/vtpnet>  
**git\_branch** devel  
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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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