

Using the FunciSNP data package 'FunciSNP: An R/Bioconductor Tool Integrating Functional Non-coding Datasets with Genetic Association Studies to Identify Candidate Regulatory SNPs'

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Contents

1	Introduction	2
2	Contact information	2
3	sessionInfo	2

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1 Introduction

This is a simple data package, to be used with the *FunciSNP* package. Please refer to the *FunciSNP* vignette for more details.

2 Contact information

Questions or comments, please contact Simon G. Coetzee (scoetzee NEAR gmail POINT com) or Houtan Noushmehr, PhD (houtan NEAR usp POINT br).

3 sessionInfo

- R version 3.1.0 RC (2014-04-02 r65358), i386-w64-mingw32
- Locale: LC_COLLATE=C, LC_CTYPE=English_United States.1252, LC_MONETARY=English_United States.1252, LC_NUMERIC=C, LC_TIME=English_United States.1252
- Base packages: base, datasets, grDevices, graphics, methods, stats, utils
- Loaded via a namespace (and not attached): tools 3.1.0

Our recent paper describing FunciSNP and FunciSNP.data can be found in the Journal Nucleic Acids Research (doi:10.1093/nar/gks542).

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