

# 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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# 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 2.15.0 (2012-03-30), i386-pc-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 2.15.0

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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