systemPipeRdata: sample data for NGS workflows

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1 Introduction

This package provides sample data for running the NGS workflows of the *systemPipeR* package (Girke, 2014). The source and usage of the data is described in the overview vignette here.

2 Version Information

toLatex(sessionInfo())

- R version 3.2.0 (2015-04-16), x86_64-apple-darwin13.4.0
- Locale: en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
- Base packages: base, datasets, graphics, grDevices, methods, stats, utils
- Other packages: knitr 1.10.5
- Loaded via a namespace (and not attached): BiocStyle 1.6.0, evaluate 0.7, formatR 1.2, highr 0.5, magrittr 1.5, stringi 0.4-1, stringr 1.0.0, tools 3.2.0

3 Funding

This project was supported by funds from the National Institutes of Health (NIH).

4 References

Thomas Girke. systemPipeR: NGS workflow and report generation environment, 28 June 2014. URL https://github.com/tgirke/systemPipeR.