

1 Introduction

2 Theory

- Hej
- Dã

2.1 ScTransform

3 Method

4 Results/Application

5 Discussion

6 References

Hafemeister, C., & Satija, R. (2019). Normalization and variance stabilization of single-cell RNA-seq data using regularized negative binomial regression. *Genome Biology* 20(296 #This is the article number. Also, Can't find page numbers.), <https://doi.org/10.1186/s13059-019-1874-1>