

Announcements 4/11

- Required reading for scRNA-seq
 - M. Luecken, F. Theis. [Current best practices in single-cell RNA-seq analysis: a tutorial](#) . *Mol Syst Bio.* 15:e8746, 2019
 - L. Maaten, G. Hinton. [Visualizing Data using t-SNE](#) . *J. of Machine Learning Research* 9:2579-2605, 2008
 - B. Sande et al. [A scalable SCENIC workflow for single-cell gene regulatory network analysis](#) . *Nat Protoc.* 15(7):2247-2276, 2020.
- Sign up time slots for project presentations
 - 5/3, 5/5, requiring attendance for both
 - https://docs.google.com/spreadsheets/d/1d4JV4y35UgO7zOzH_98ePteqmZeKe_BQtdsOs1ofaeY/edit#gid=0
 - 10-12 mins presentation plus 2 mins Q&A
 - https://www.biostat.wisc.edu/bmi776/project_presentation.pdf