



Useful links:

- The Workflow of Your Analysis: <http://staff.math.su.se/hoehle/talks/workflow-2018-12-14.pdf>
- Our workflow today roughly follows: https://satijalab.org/seurat/pbmc3k_tutorial_v3.html
- Data for today: https://s3-us-west-2.amazonaws.com/10x.files/samples/cell/pbmc3k/pbmc3k_filtered_gene_bc_matrices.tar.gz
- *Essential guidelines for computational method benchmarking:* <https://arxiv.org/abs/1812.00661>