

# Introduction to RNA Velocity Analysis

7/20/2022

Hands on tour of the single cell analysis journey

## Recap: Neutrophil analysis plan

introns"

Our journey through analysis

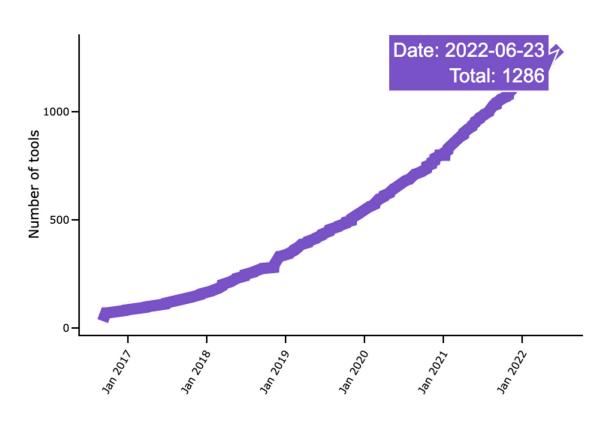
cells"

Neutrophils have Neutrophils high Cells separate Neutrophils exist Separate low UMI content intron retention into distinct neutrophils from at different stages dead/dying cells of maturation clusters rates Solution Filter Include low Annotate cell **Trajectory** Map introns UMI barcodes background analysis clusters **Cell Ranger "force-**Cell Ranger "include-**Loupe Browser Loupe Browser 3rd Party** 



## **Approaching the Analysis Ecosystem**

More than 1,000 tools!



https://www.scrna-tools.org/

- How do you choose?
  - Leverage your research question
  - Look to the literature
    - Citations
    - Reviews
  - Look at GitHub sites
    - Check for regular updates
    - Check for issues and responses
- 10x Analysis Guides
  - Introductions
  - Tutorials
  - Informatics blogs



## **RNA Velocity**

Single-cell RNA-seq provides only static snapshots of cellular states at the moment of the measurement.

RNA velocity (<u>La Manno et al, 2018</u>; <u>Bergen et al. 2020</u>) can predict the direction and speed of movement of cells in transcriptome space.

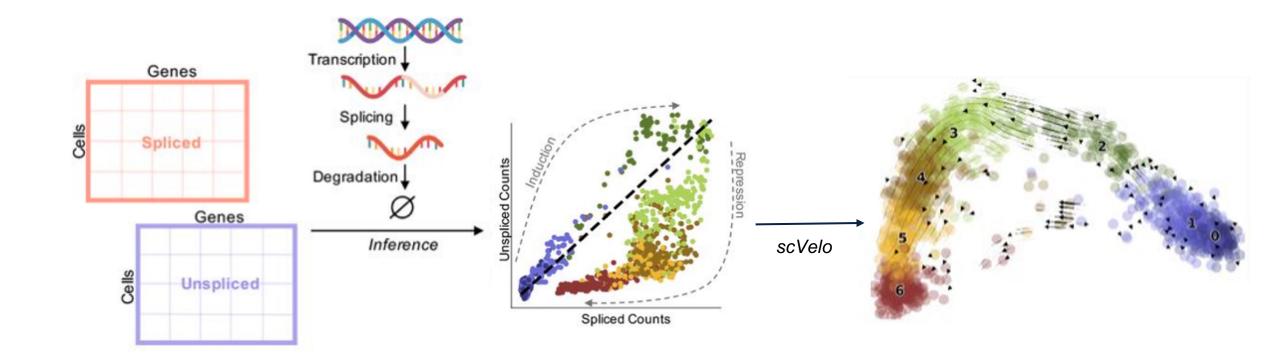
Application: analysis of cell dynamics → developmental biology, tissue regeneration, disease progression





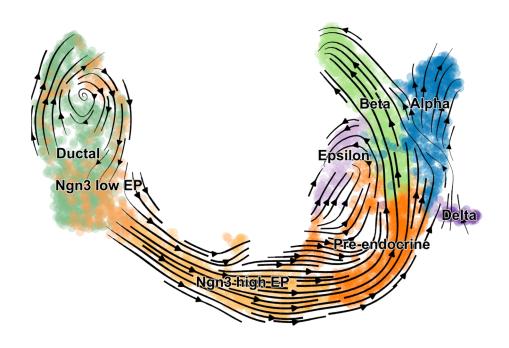
## How does RNA velocity work?

Estimation basis: the relative abundance of nascent (**unspliced**) and mature (**spliced**) mRNA → rate of RNA splicing and degradation



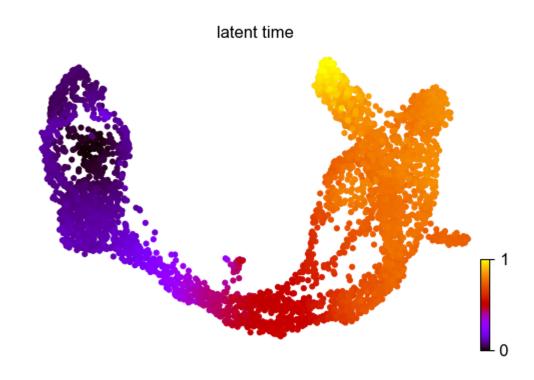


### **Example of scVelo results**



Endocrine development in the pancreas with lineage commitment to four major fates:  $\alpha$ ,  $\beta$ ,  $\delta$  and  $\epsilon$ -cells.

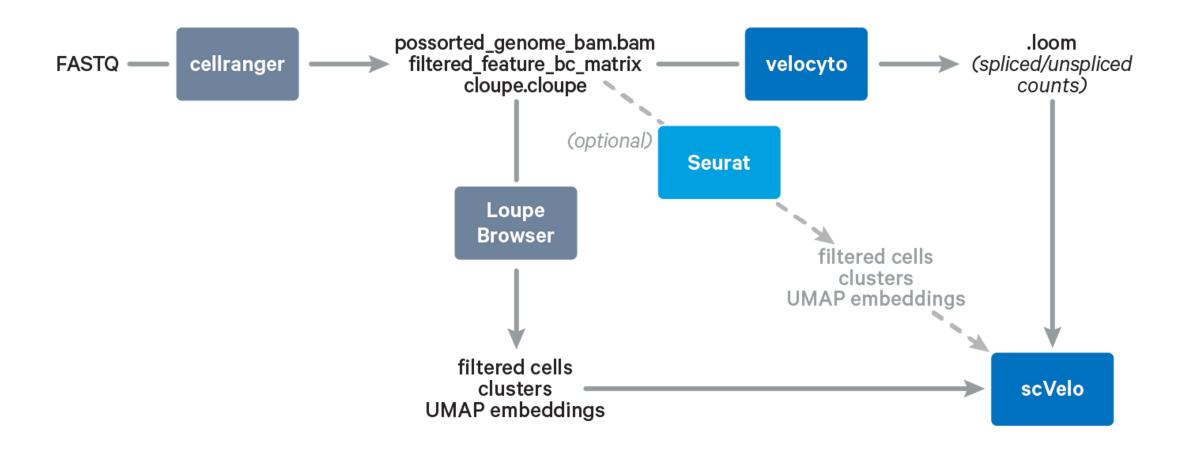
https://scvelo.readthedocs.io/



The <u>latent time</u> represents the **cell's internal clock** and approximates the real time experienced by cells as they differentiate, based only on its transcriptional dynamics.



## Velocity analysis using 10x data



https://www.10xgenomics.com/resources/analysis-guides/trajectory-analysis-using-10x-Genomics-single-cell-gene-expression-data

