

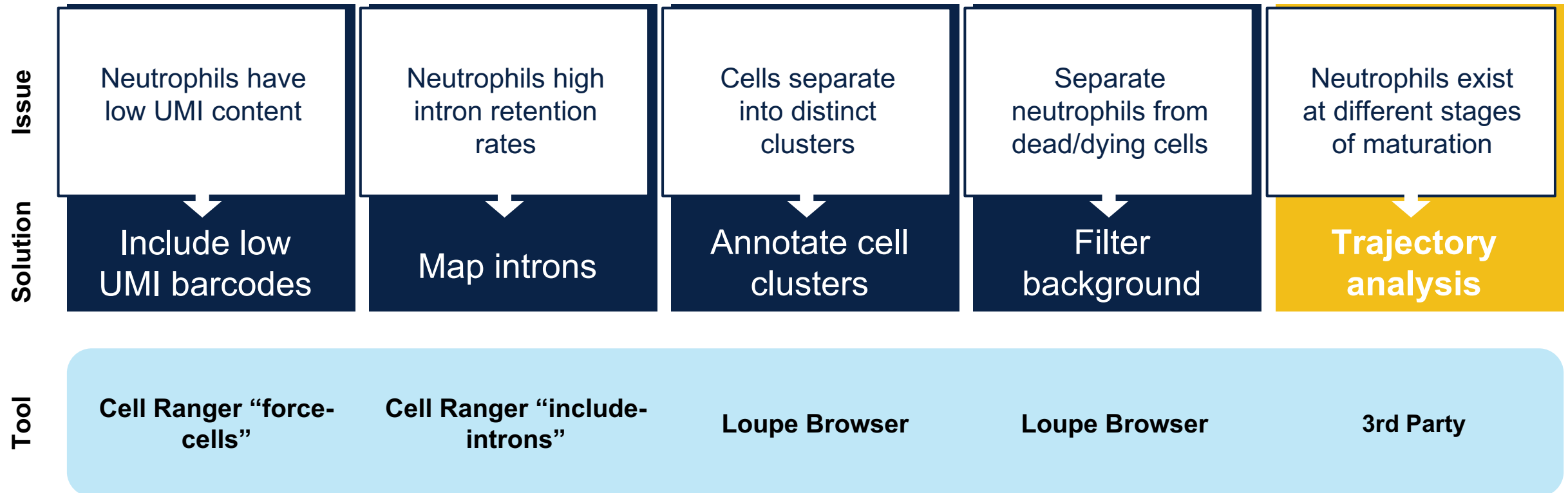
Introduction to RNA Velocity Analysis

7/20/2022

Hands on tour of the single cell analysis journey

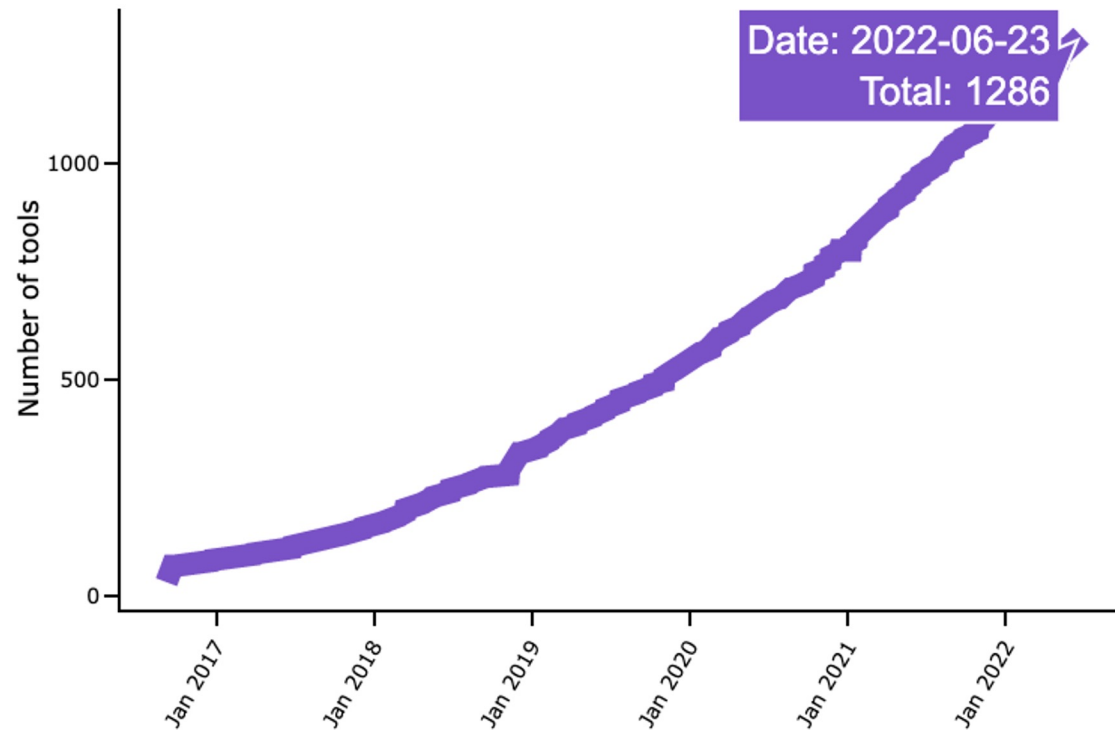
Recap: Neutrophil analysis plan

Our journey through analysis



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.

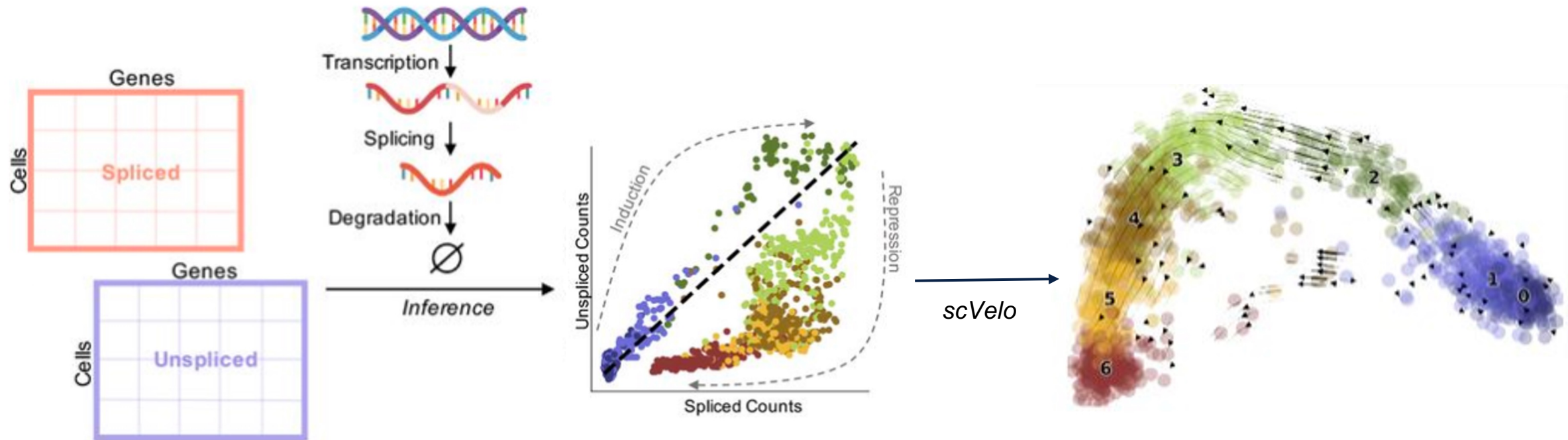
RNA velocity ([La Manno et al, 2018](#); [Bergen et al. 2020](#)) can predict the direction and speed of movement of cells in transcriptome space.

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

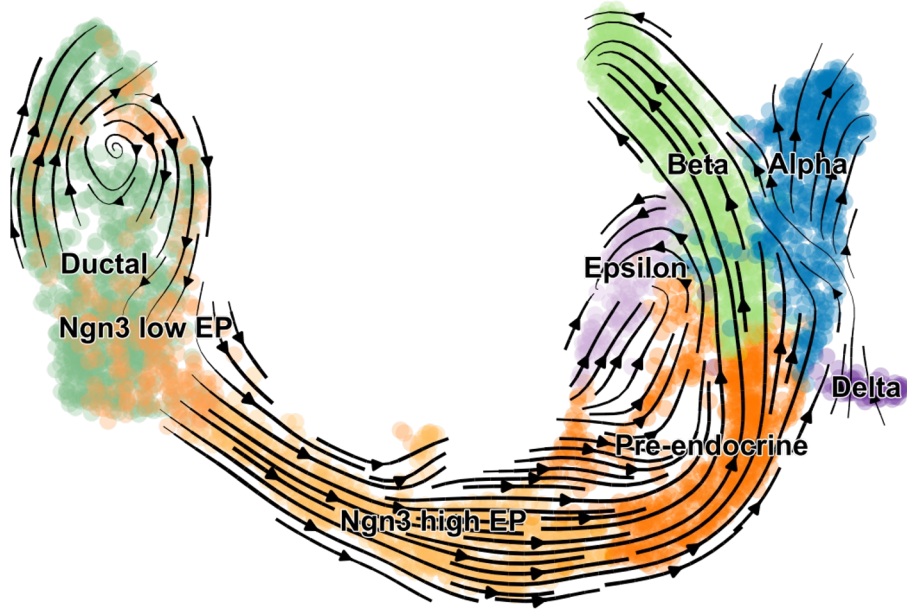


How does RNA velocity work?

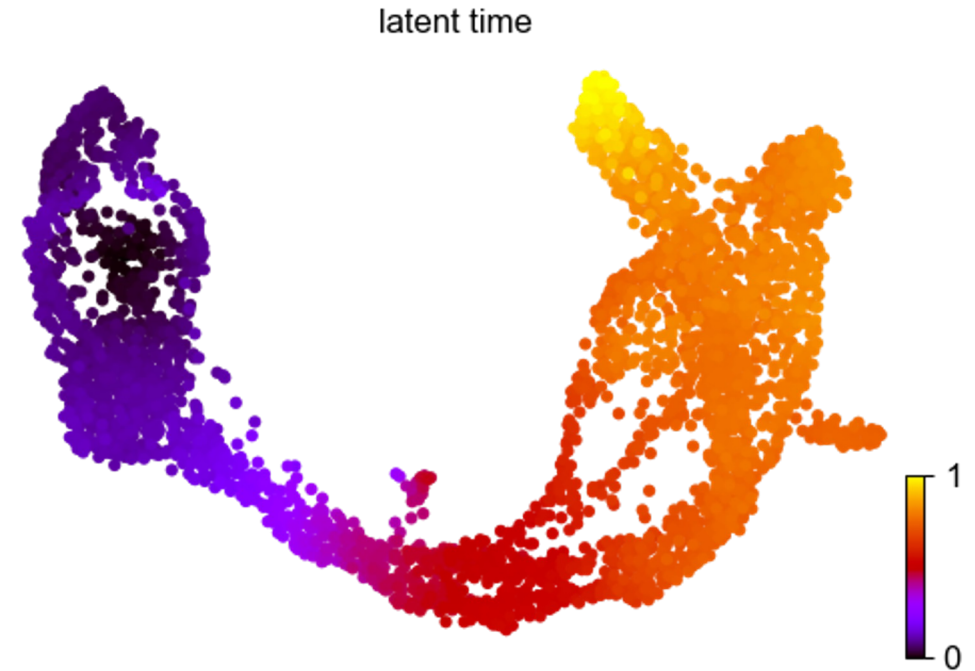
Estimation basis: the relative abundance of nascent (**unspliced**) and mature (**spliced**) mRNA → rate of RNA splicing and degradation



Example of scVelo results

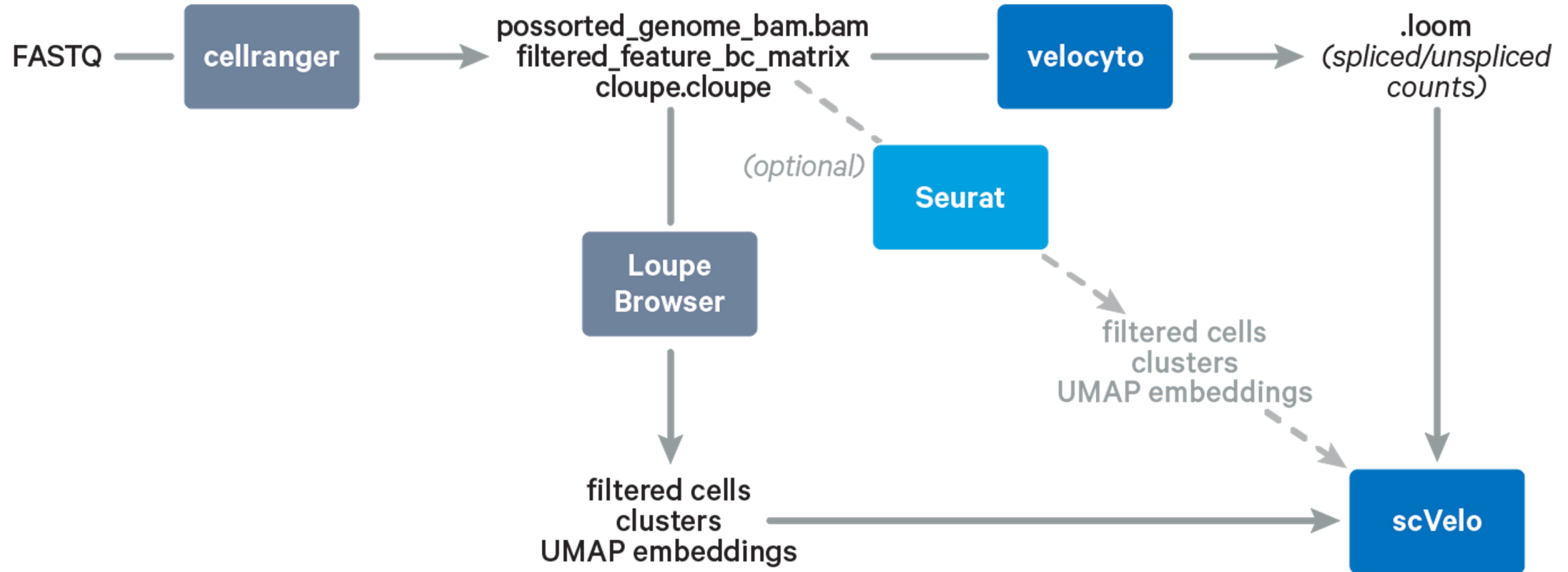


Endocrine development in the pancreas with lineage commitment to four major fates: α , β , δ and ϵ -cells.



The **latent time** 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>