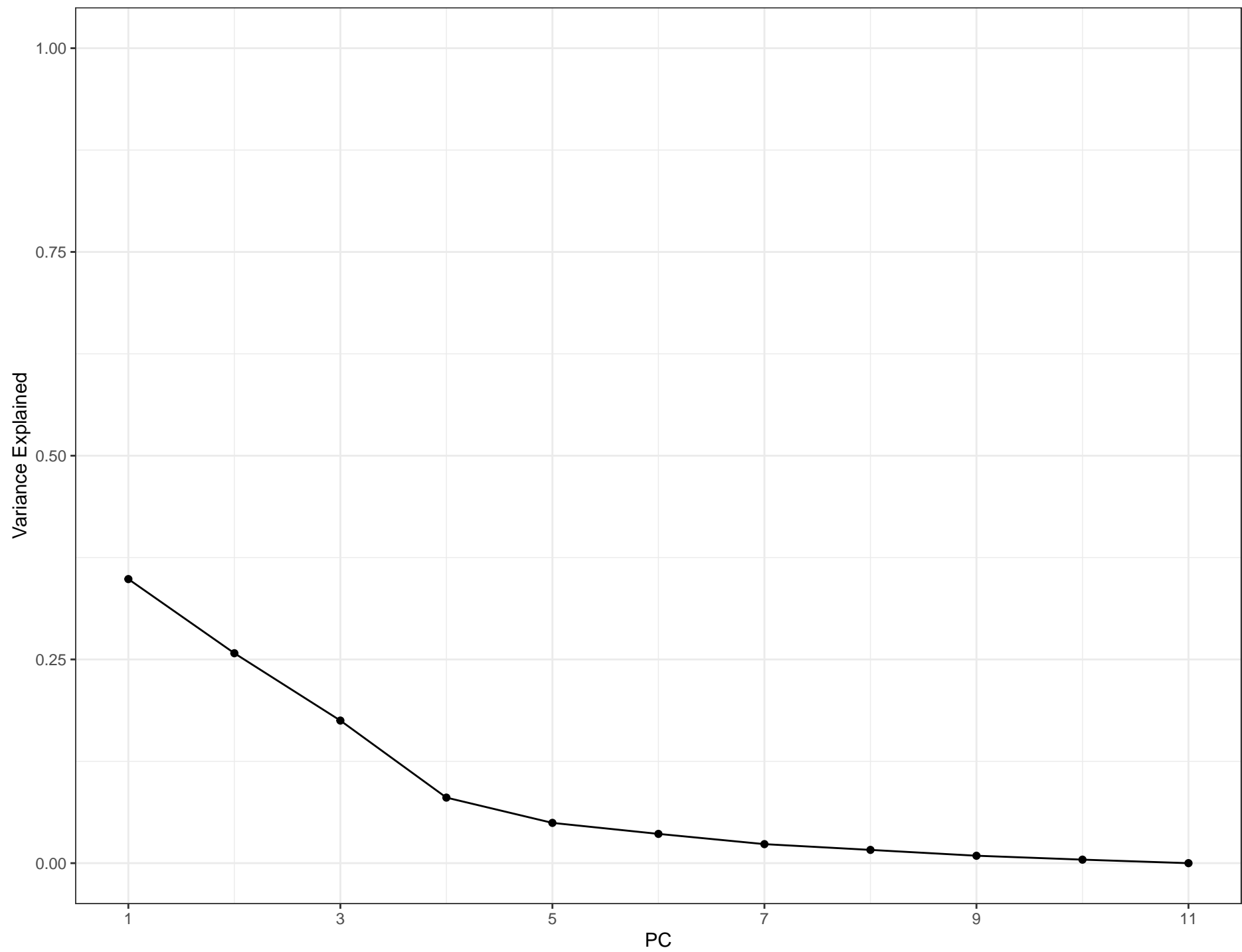
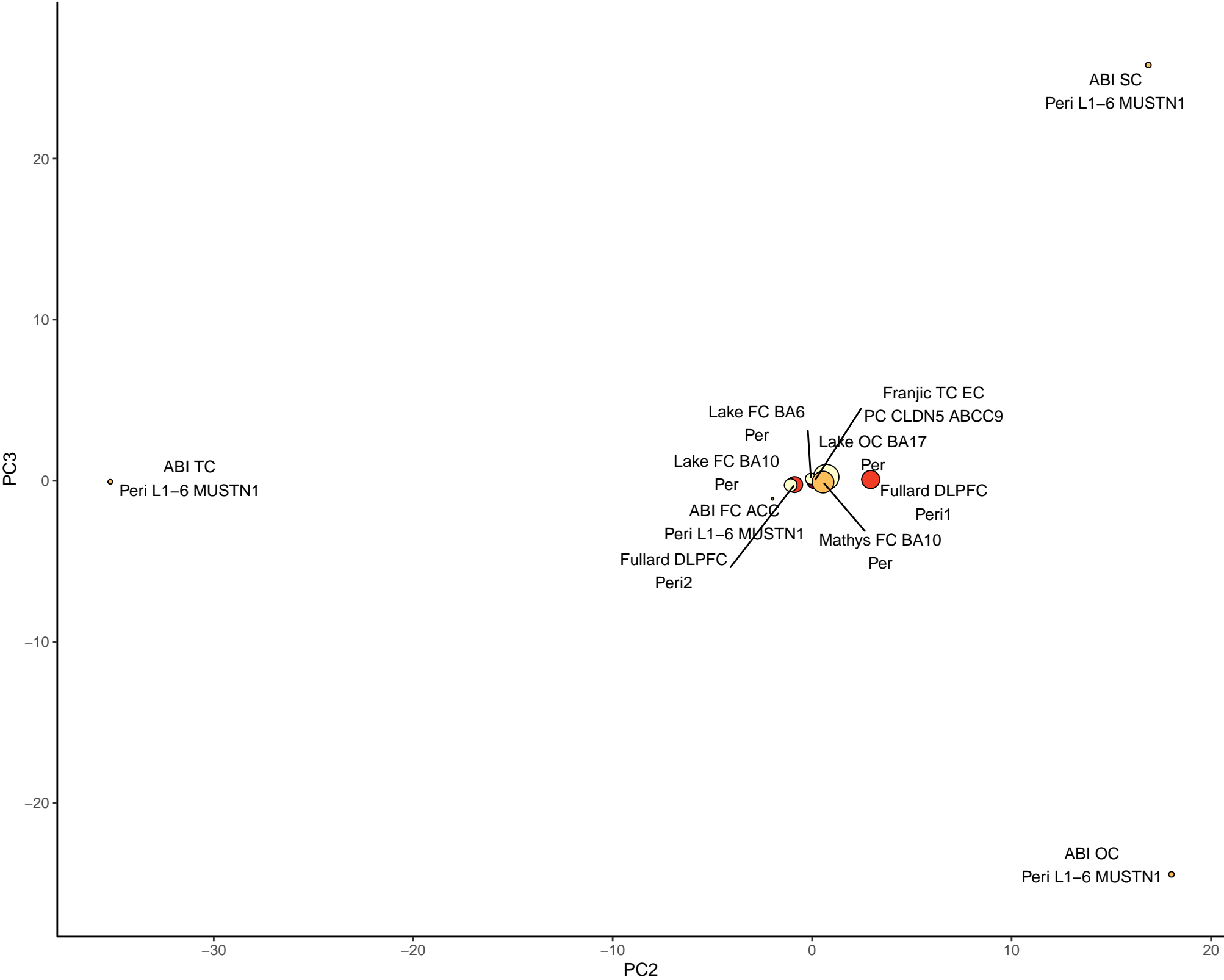


**PER Subtypes**  
**Normalized Counts Mean Expression Scree Plot**  
6 cell types, 10 datasets, 8,703 genes



Pericyte Batch Corrected Mean Expression

6 cell types found in 10 datasets from 5 studies  
8,703 genes



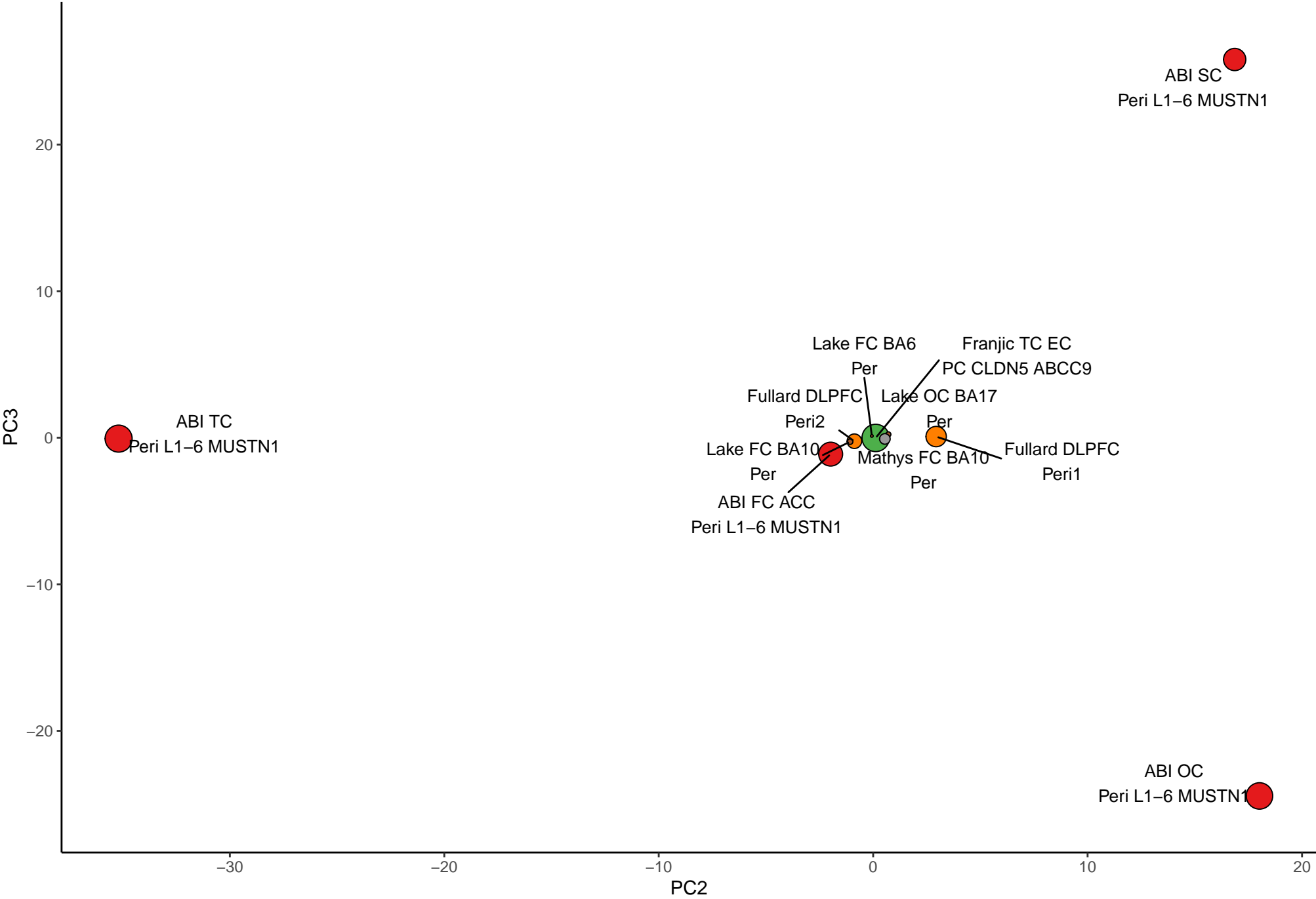
# Nuclei

100

Year 2018 2019 2021 2022

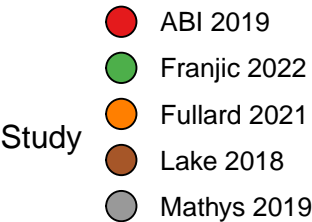
Pericyte Batch Corrected Mean Expression

6 cell types found in 10 datasets from 5 studies  
8,703 genes



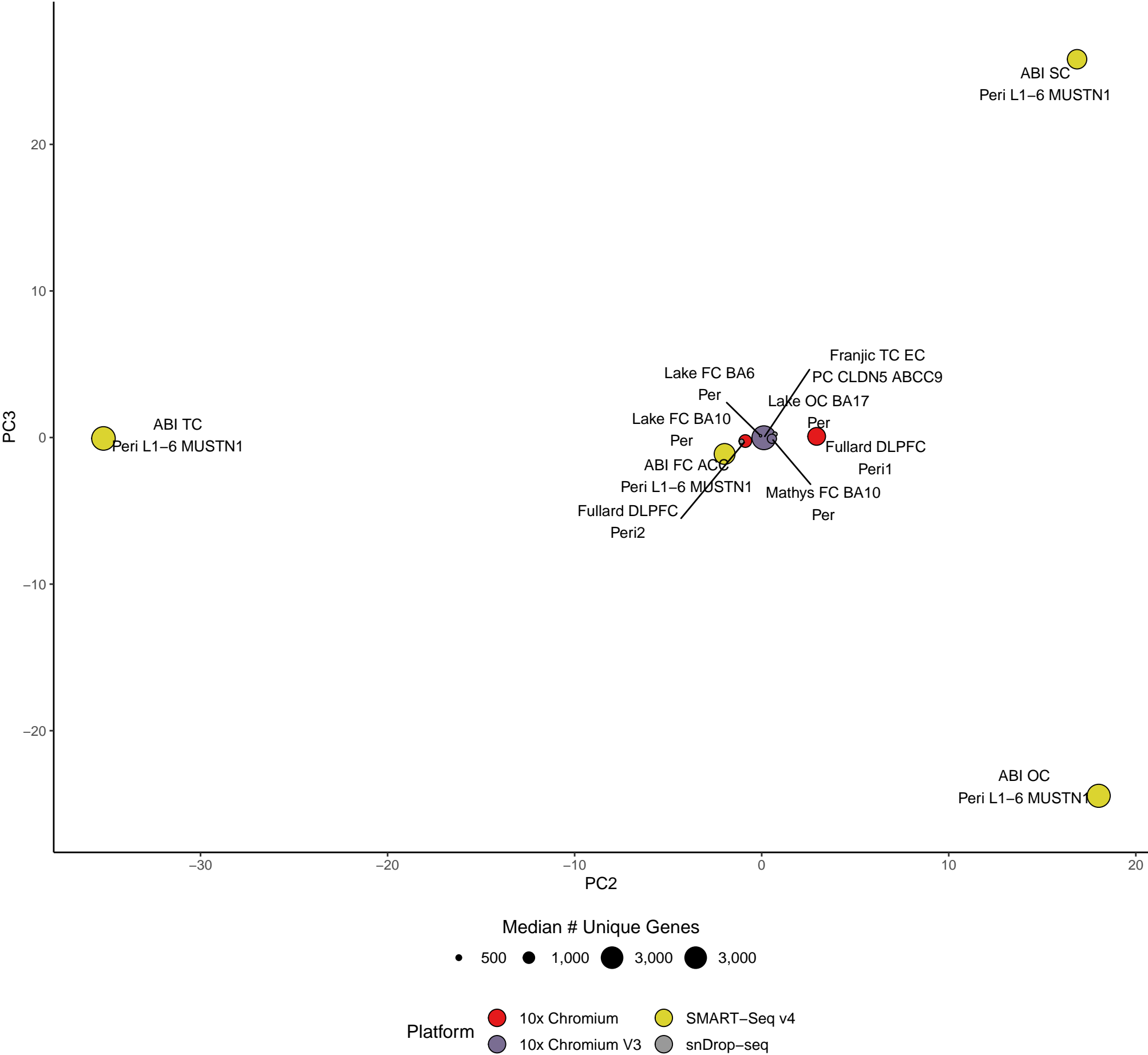
Median # Unique Genes

• 400 • 600 • 2,100 • 3,100 • 3,700

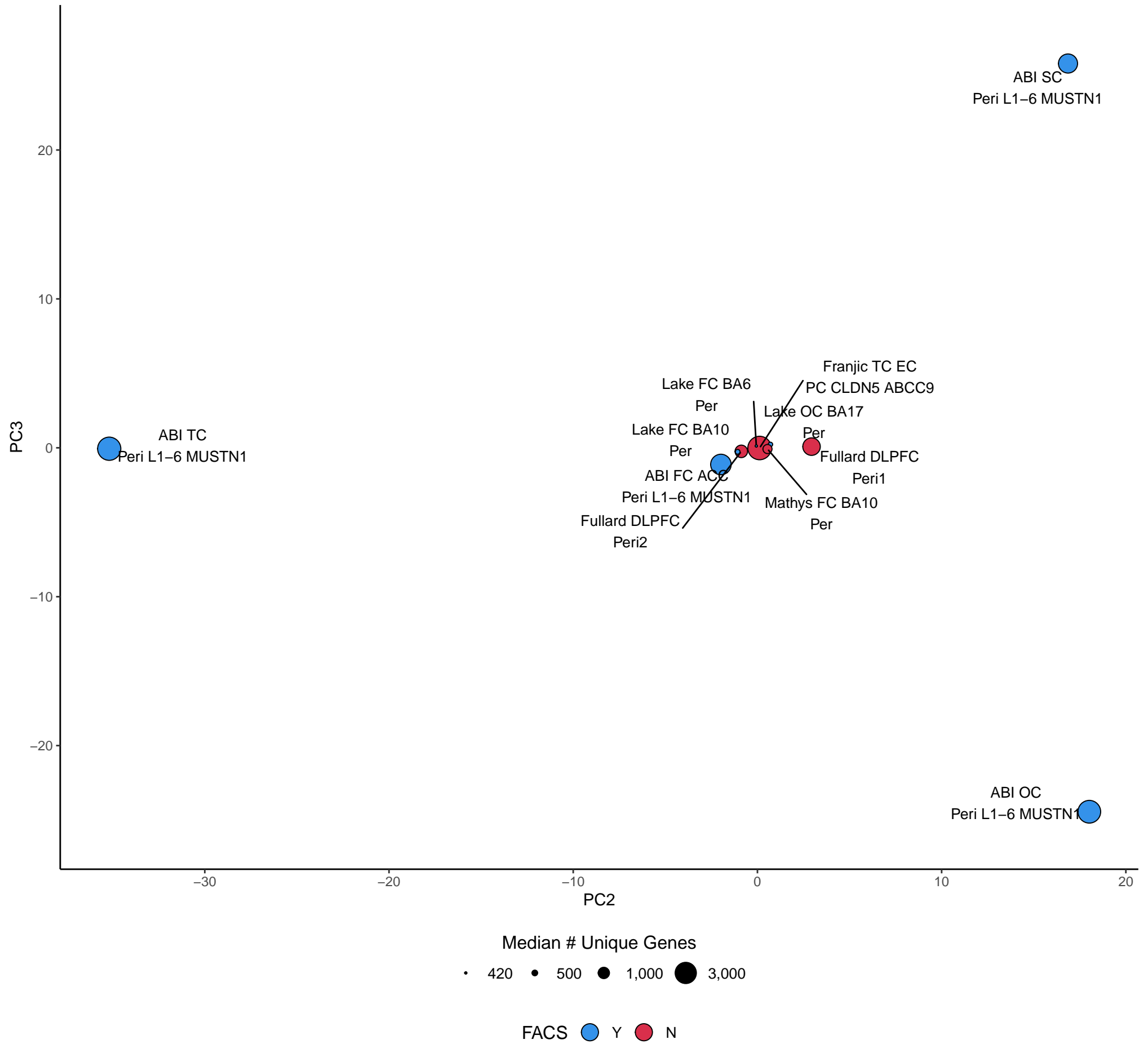


Pericyte Batch Corrected Mean Expression

6 cell types found in 10 datasets from 5 studies  
8,703 genes



6 cell types found in 10 datasets from 5 studies  
8,703 genes



Pericyte Batch Corrected Mean Expression

6 cell types found in 10 datasets from 5 studies  
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