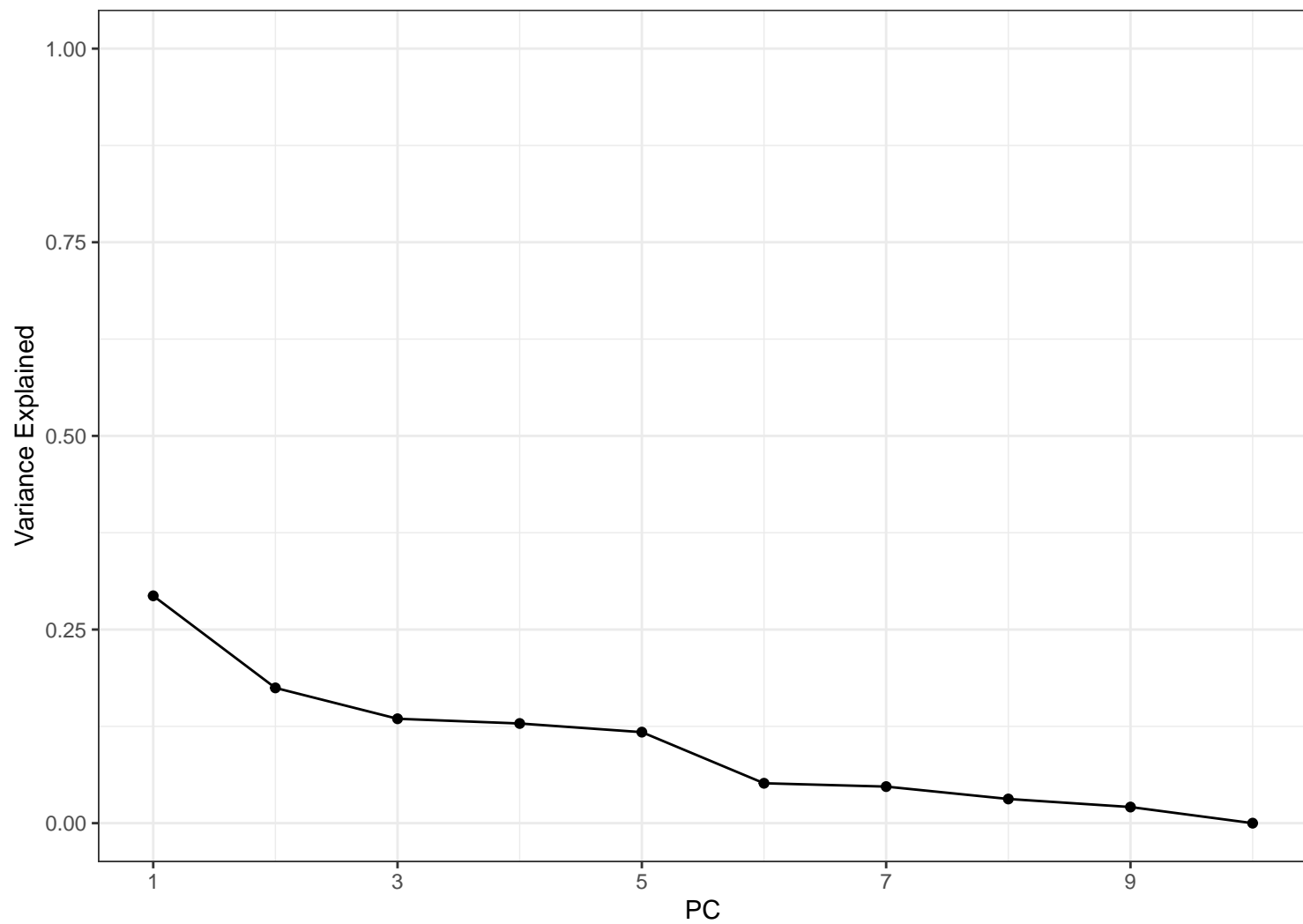
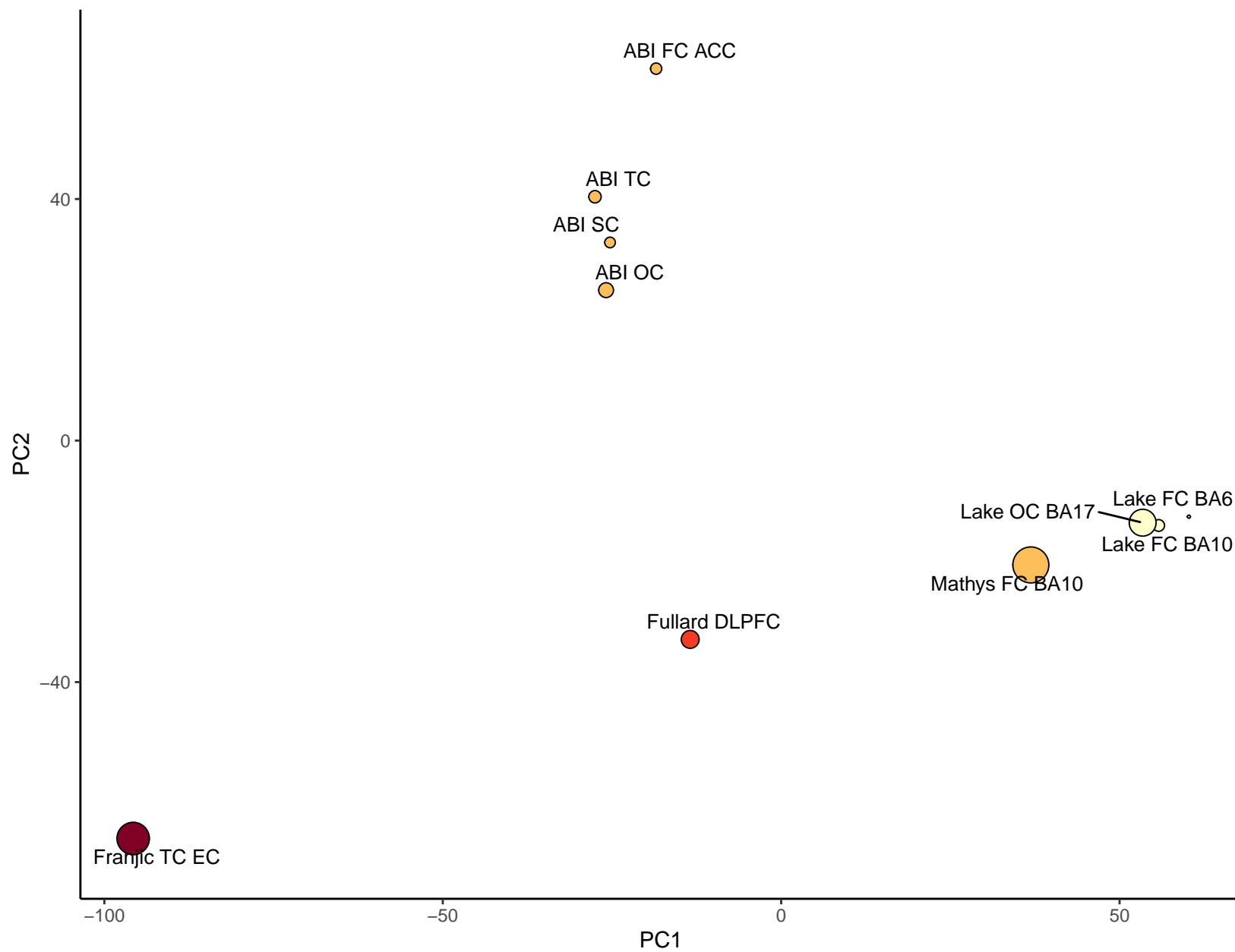


PER Subtypes  
Normalized Counts Mean Expression Scree Plot  
10 datasets, 8,596 genes



# Dataset Pericyte Mean Expression Projections

8,596 protein coding genes



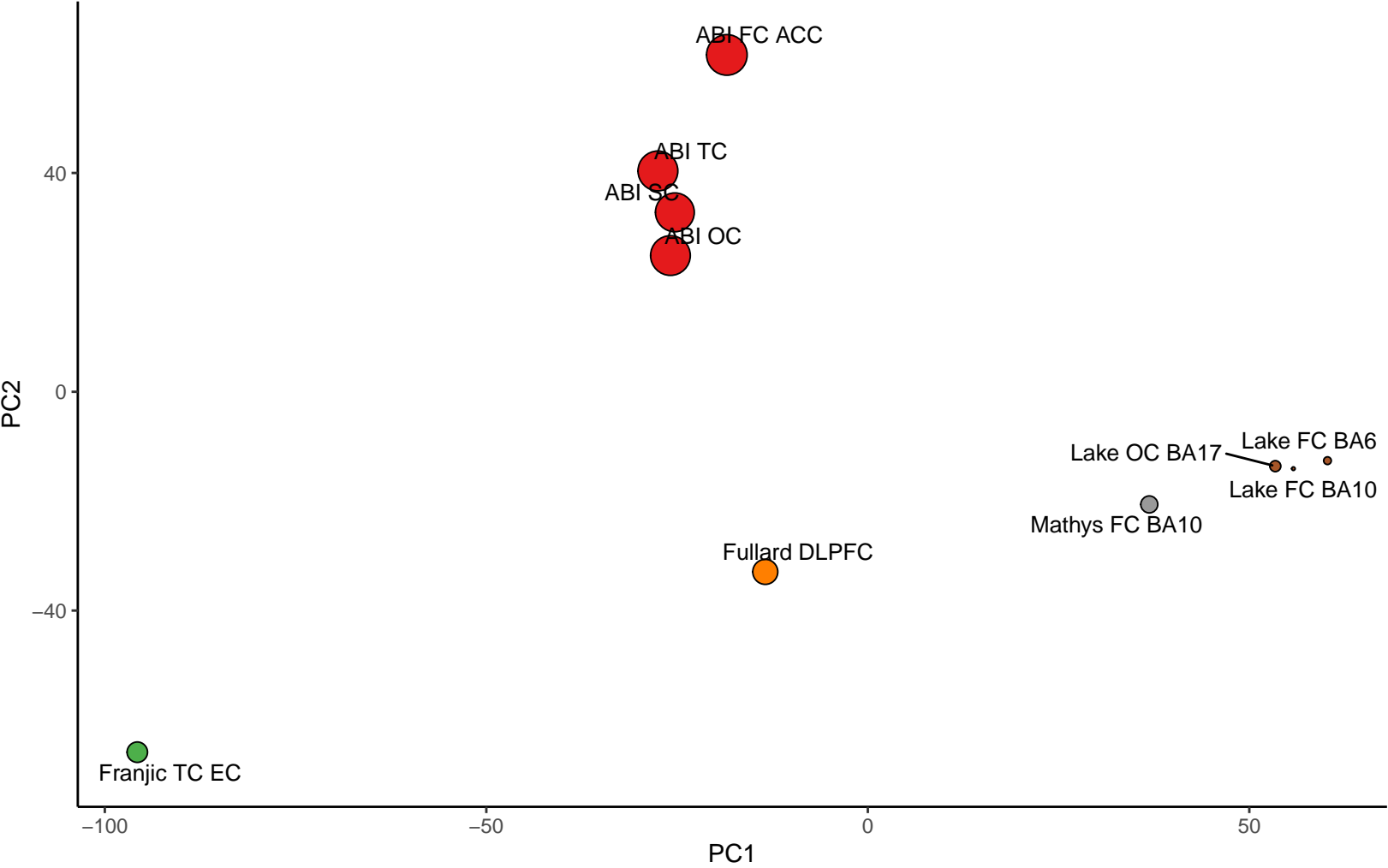
# Nuclei

• 4,100 • 5,900 • 7,100 • 19,300 • 33,300

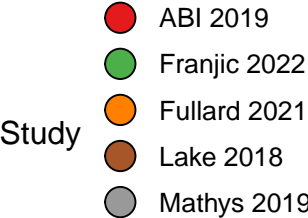
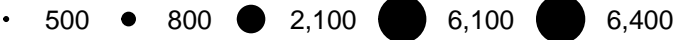
Year 2018 2019 2021 2022

Dataset Pericyte Mean Expression Projections

8,596 protein coding genes

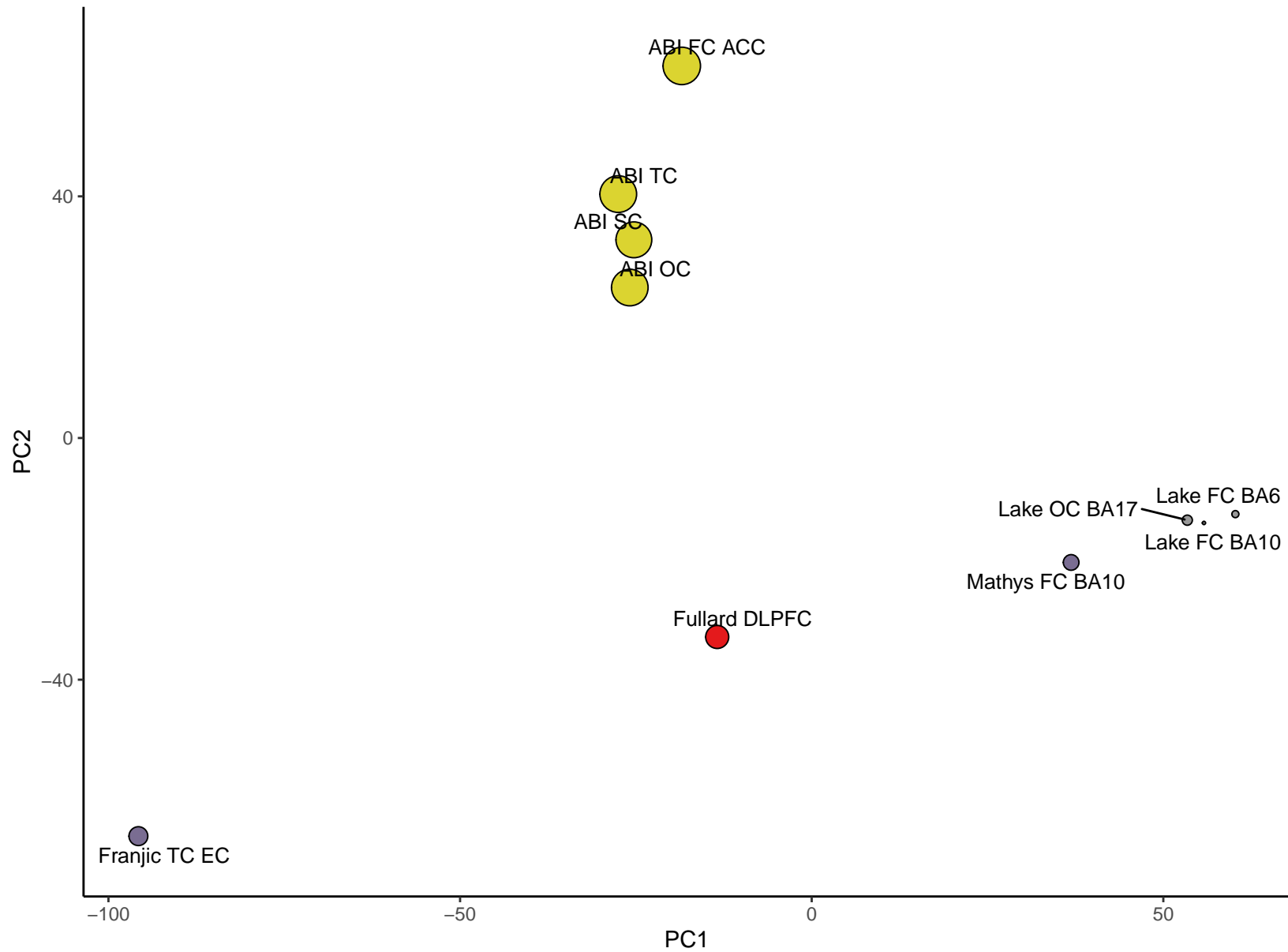


Median # Unique Genes

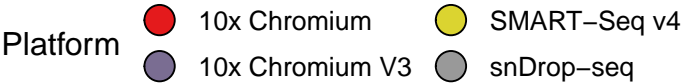
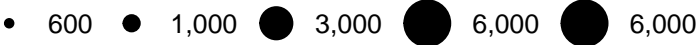


Dataset Pericyte Mean Expression Projections

8,596 protein coding genes

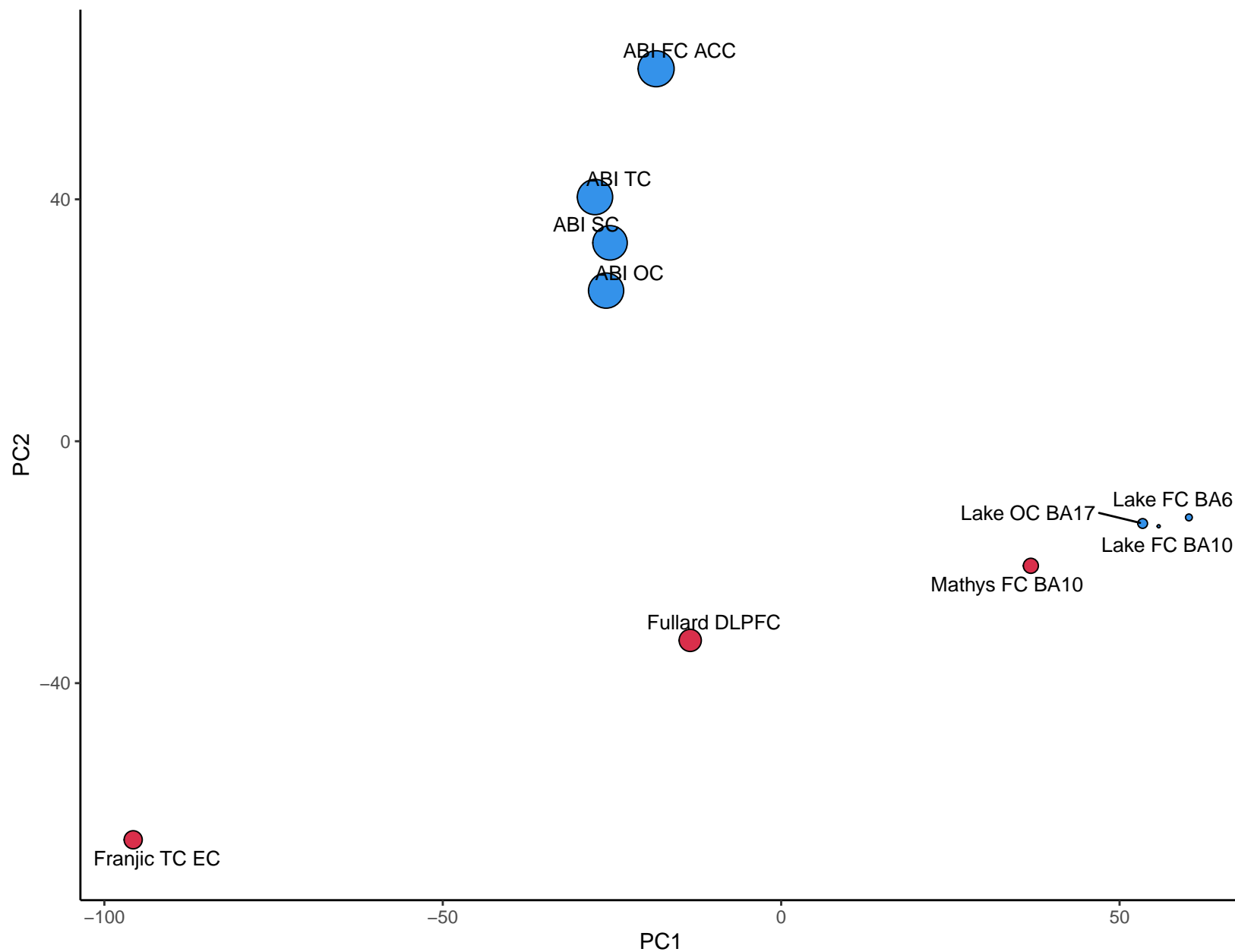


Median # Unique Genes



# Dataset Pericyte Mean Expression Projections

8,596 protein coding genes



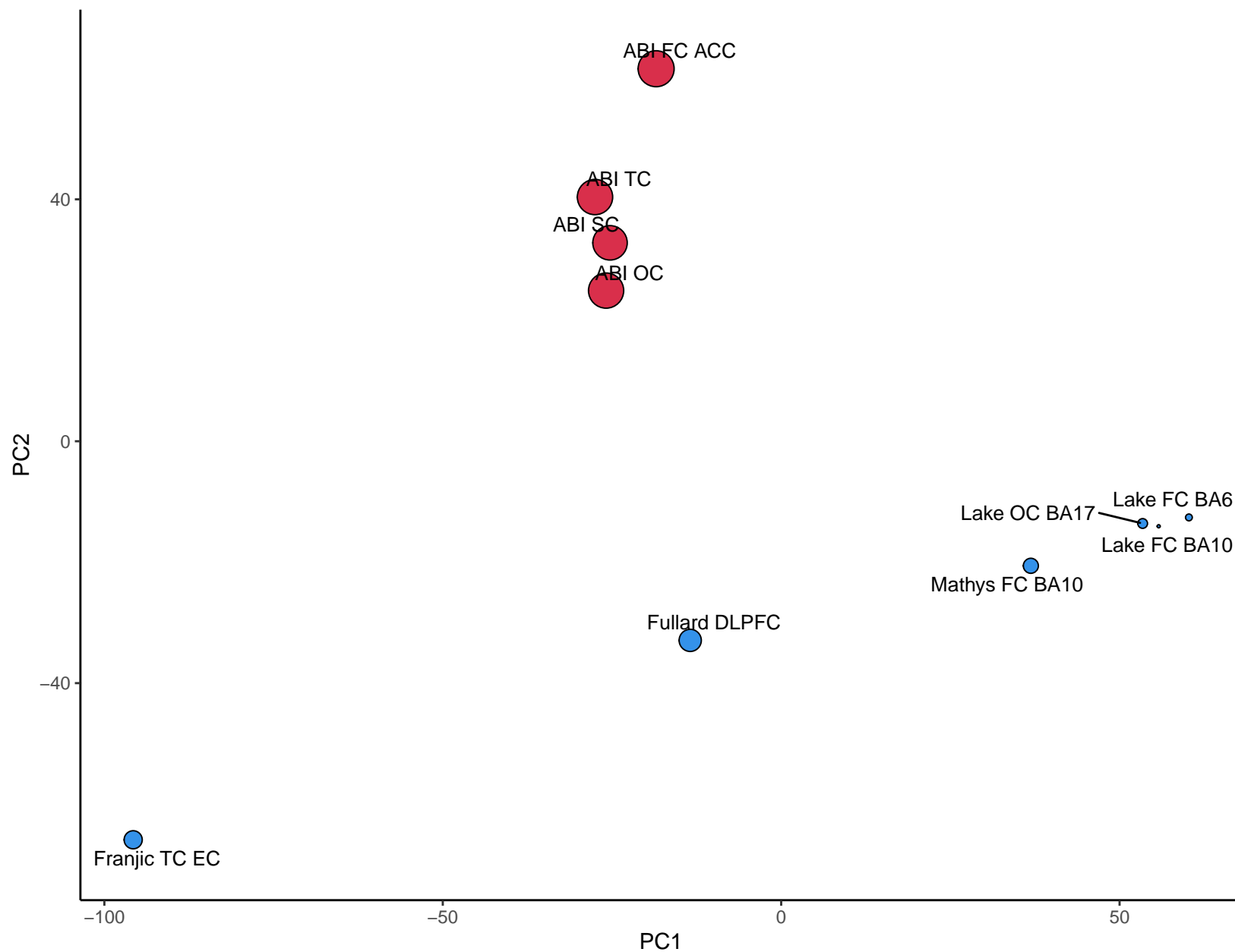
Median # Unique Genes

• 539 • 1,000 • 5,000 • 6,000

FACS Y N

# Dataset Pericyte Mean Expression Projections

8,596 protein coding genes



Median # Unique Genes

• 539 • 1,000 • 5,000 • 6,000

Unbiased Sampling Y N