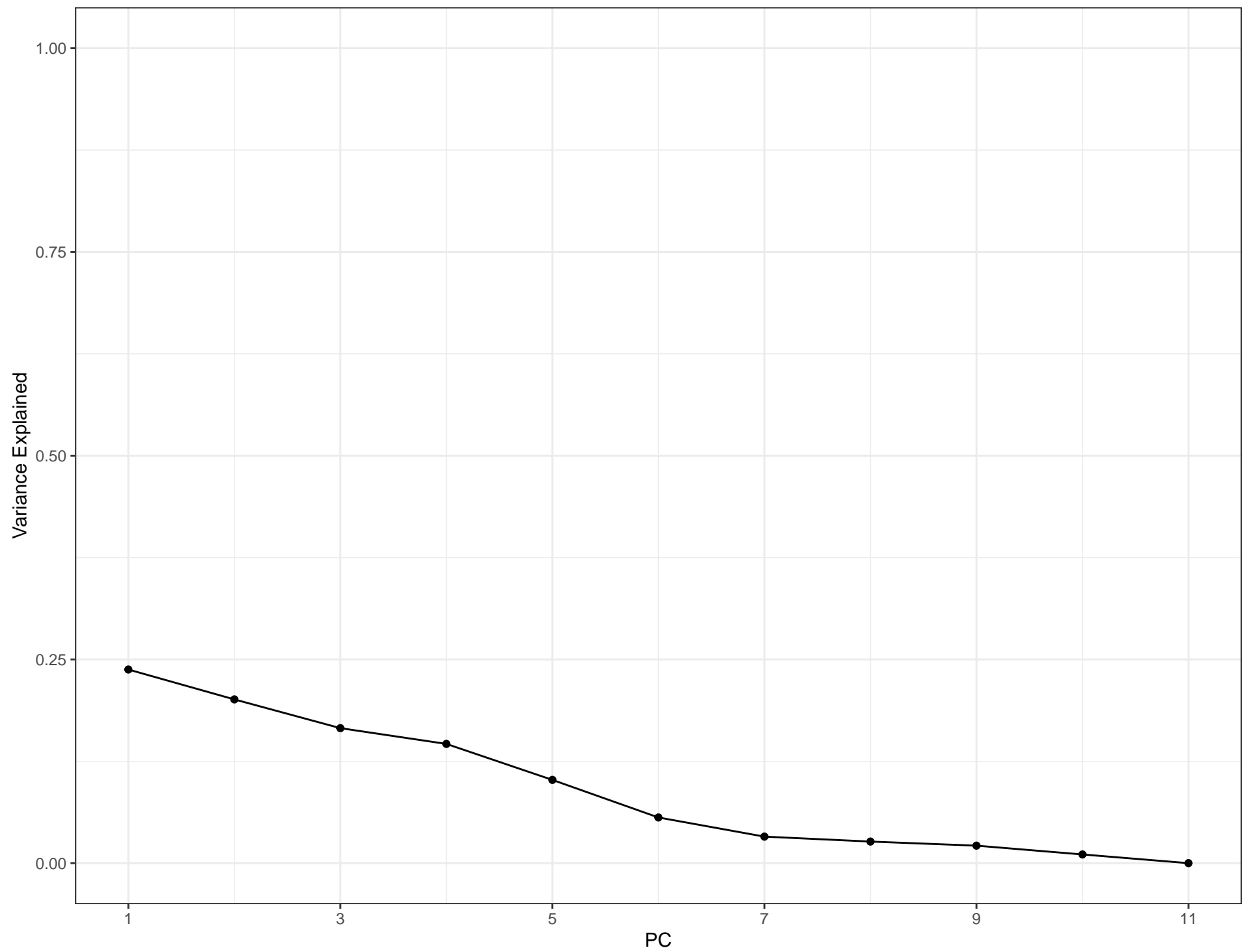
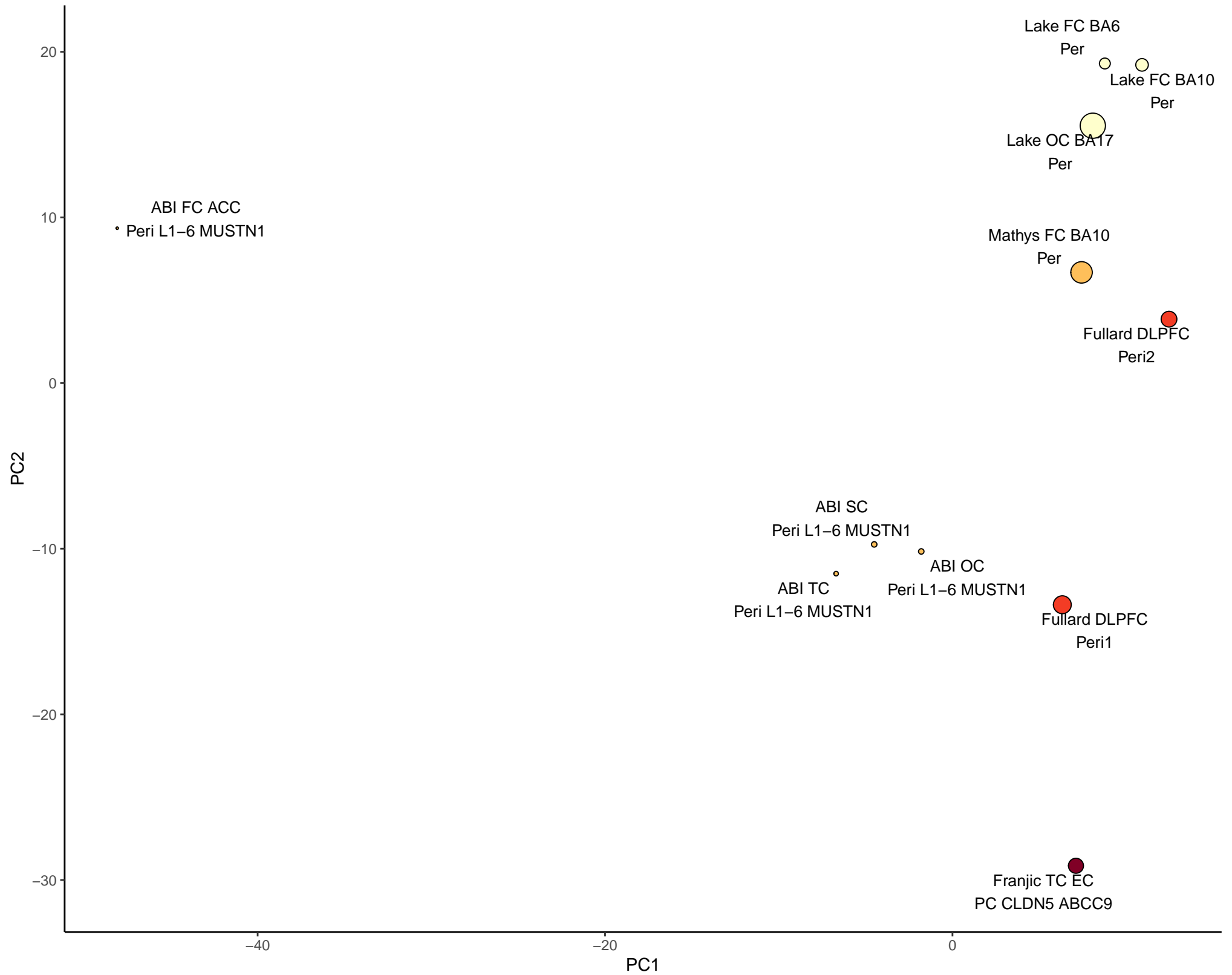


PER Subtypes
Normalized Counts Mean Expression Scree Plot
6 cell types, 10 datasets, 8,703 genes



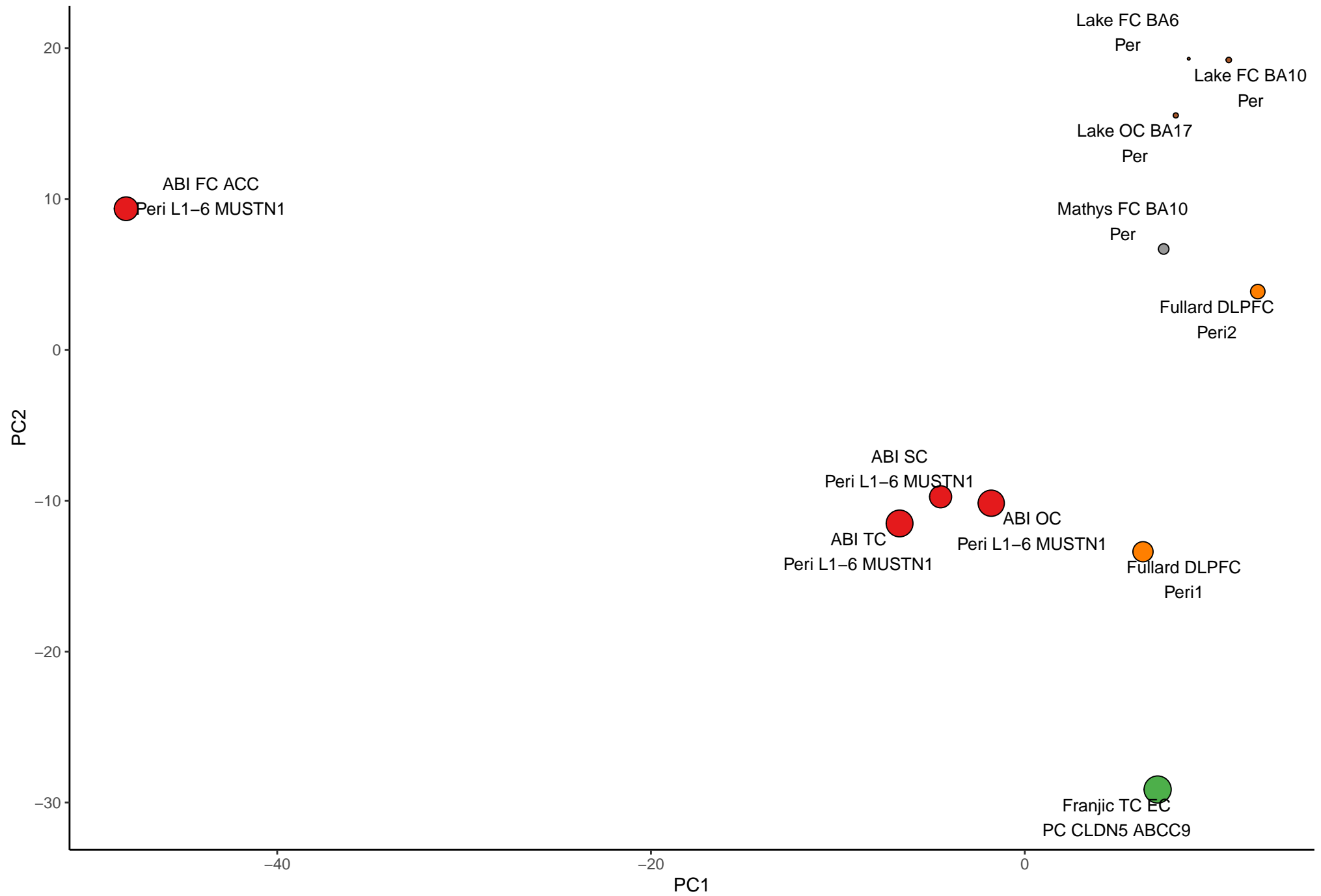
Pericyte Mean Expression

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



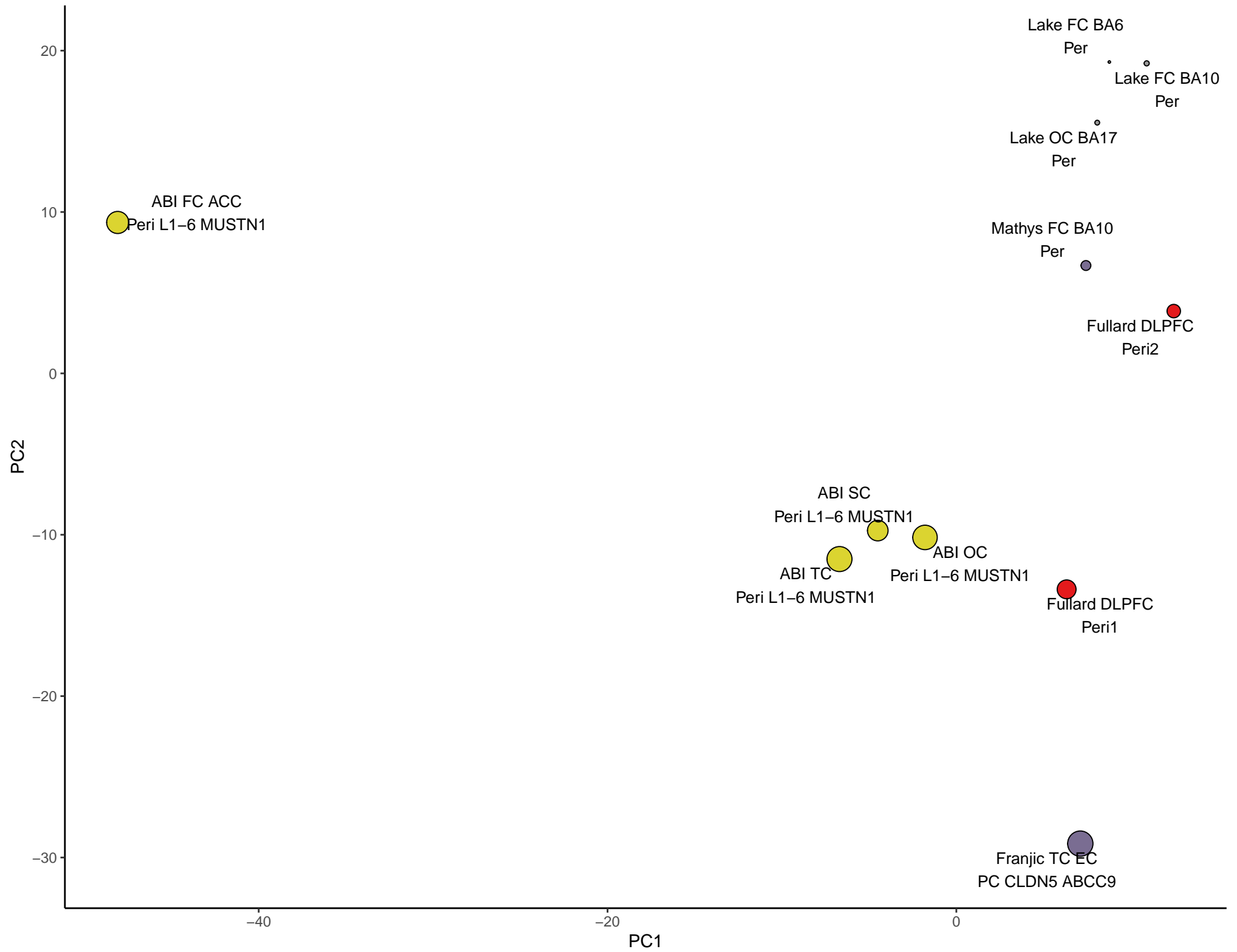
Pericyte Mean Expression

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



Pericyte Mean Expression

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



Median # Unique Genes

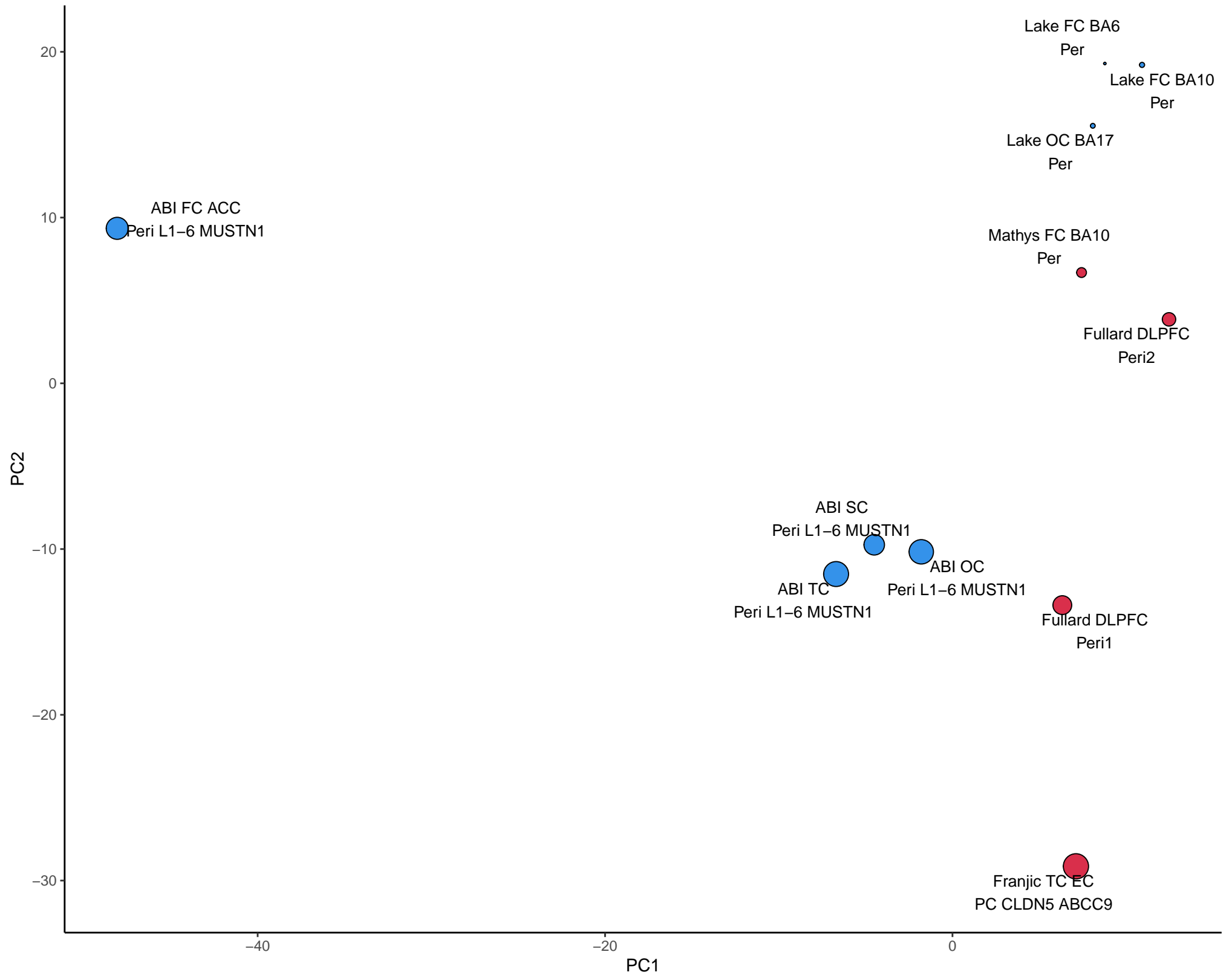
• 500 • 1,000 • 3,000 • 3,000

Platform

10x Chromium SMART-Seq v4
10x Chromium V3 snDrop-seq

Pericyte Mean Expression

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



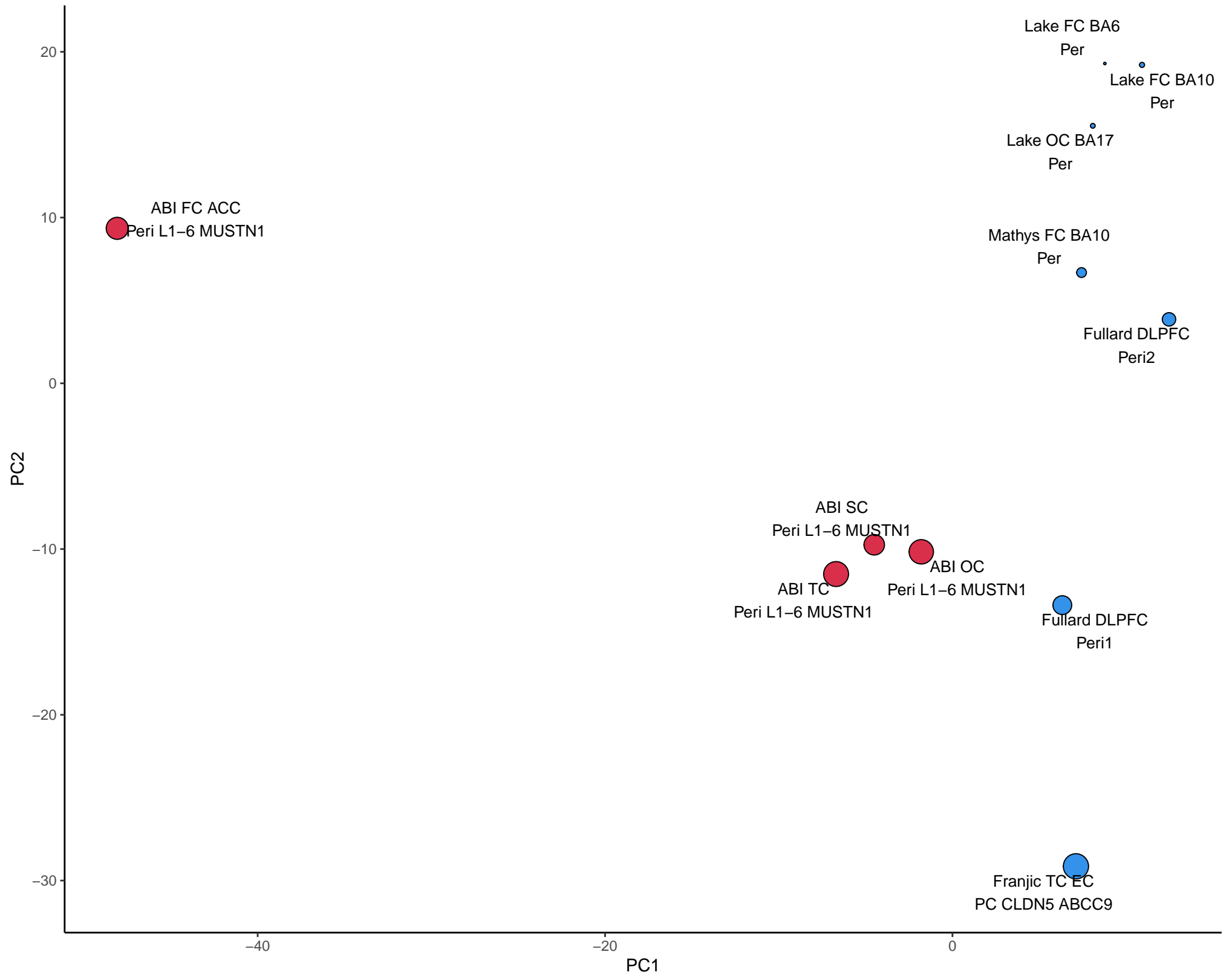
Median # Unique Genes

• 420 • 500 • 1,000 • 3,000

FACS Y N

Pericyte Mean Expression

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



Median # Unique Genes

• 420 • 500 • 1,000 • 3,000

Unbiased Sampling Y N