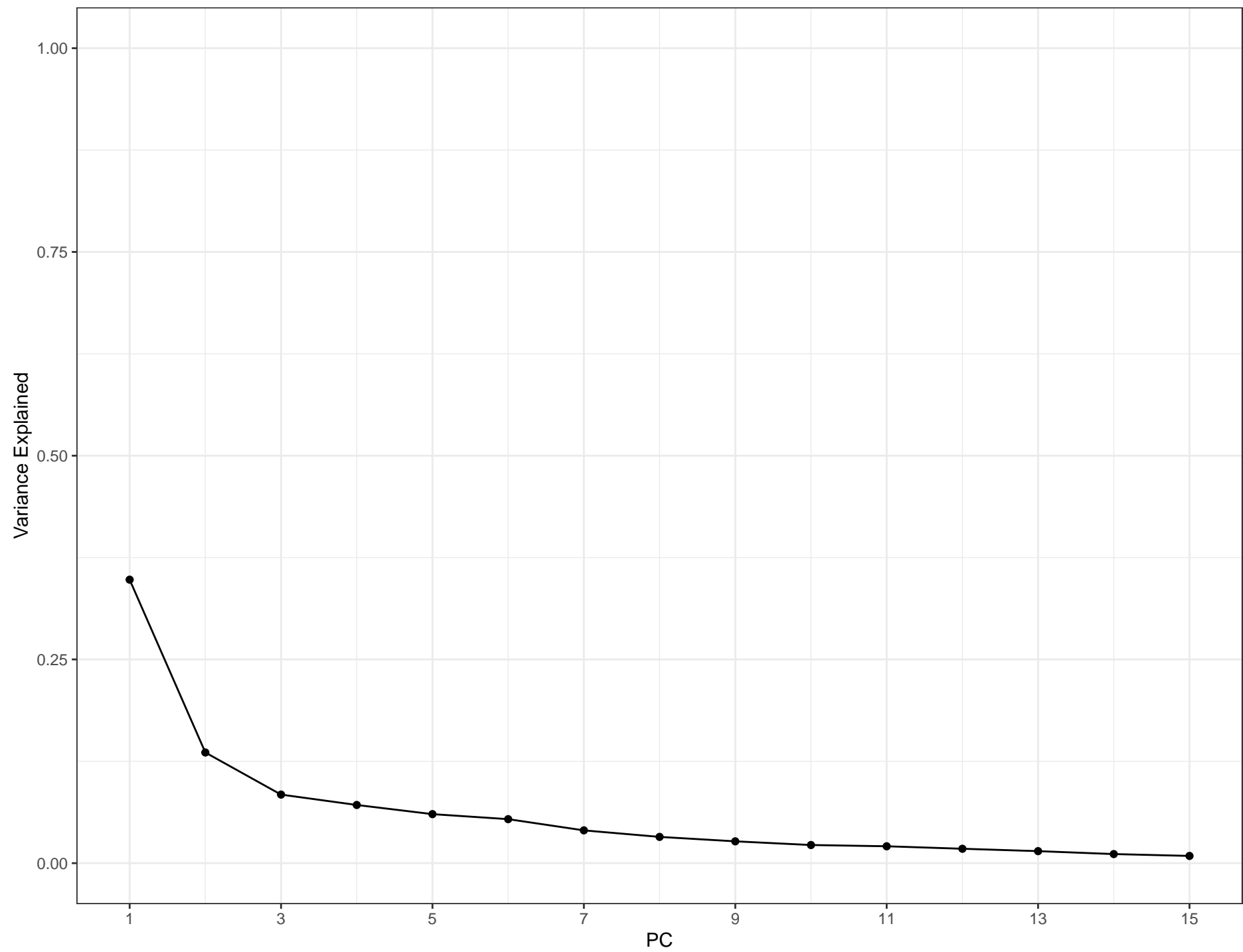
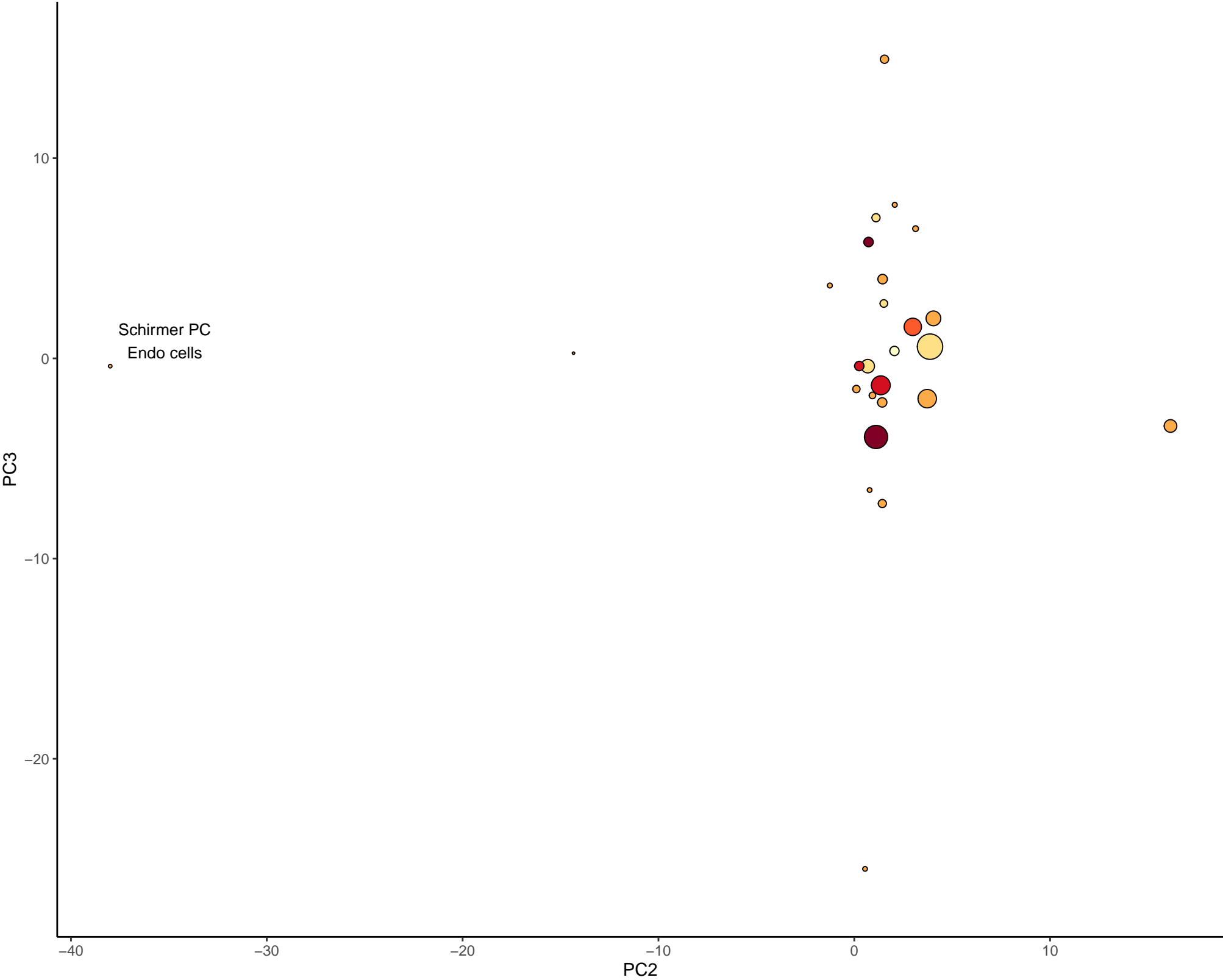


END Subtypes
Normalized Counts Mean Expression Scree Plot
17 cell types, 25 datasets, 8,722 genes



Endothelial Batch Corrected Mean Expression

17 cell types found in 25 datasets from 15 studies
8,722 genes



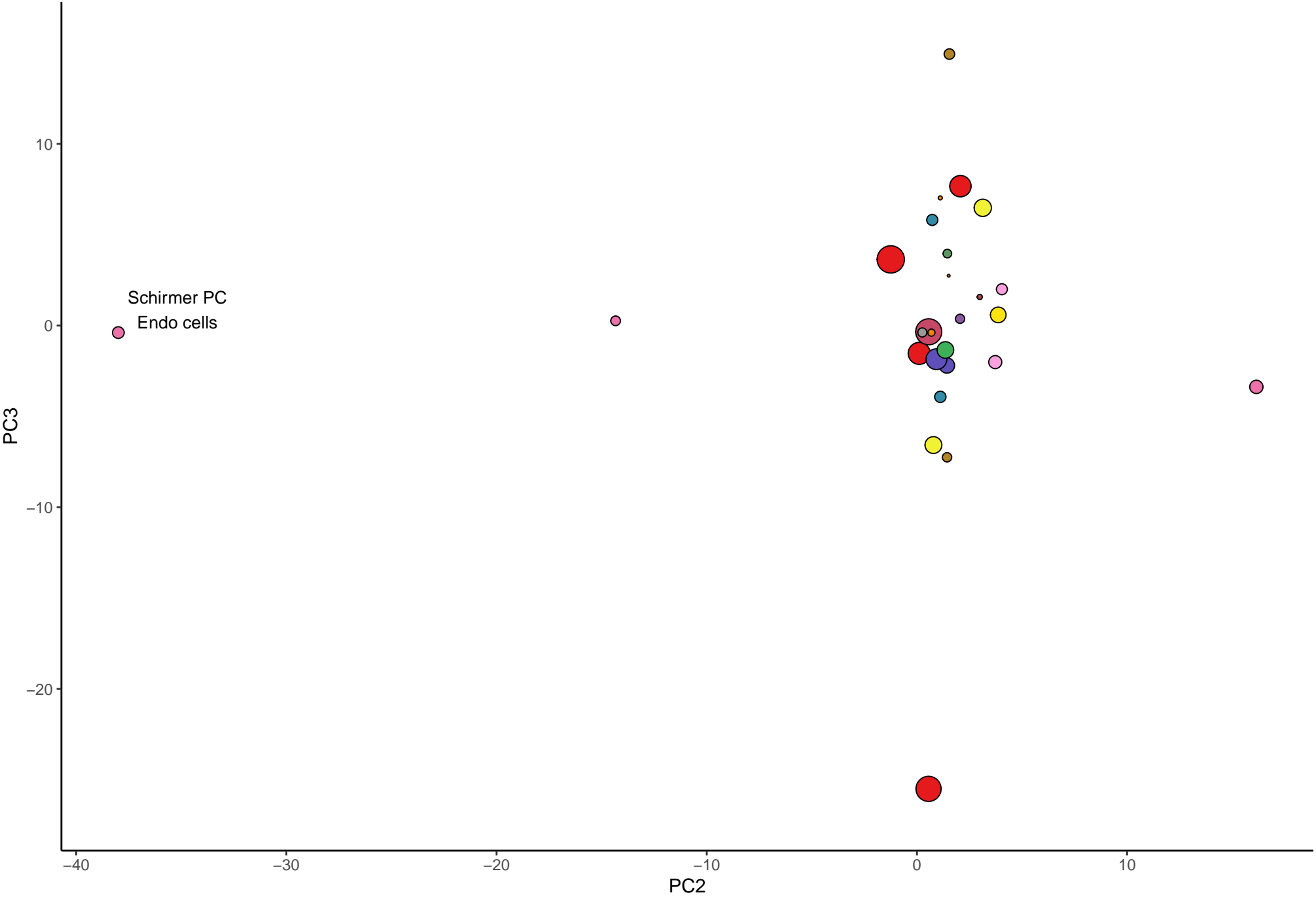
Nuclei

• 1 ● 100

Year 2017 2018 2019 2020 2021 2022

Endothelial Batch Corrected Mean Expression

17 cell types found in 25 datasets from 15 studies
8,722 genes



Median # Unique Genes

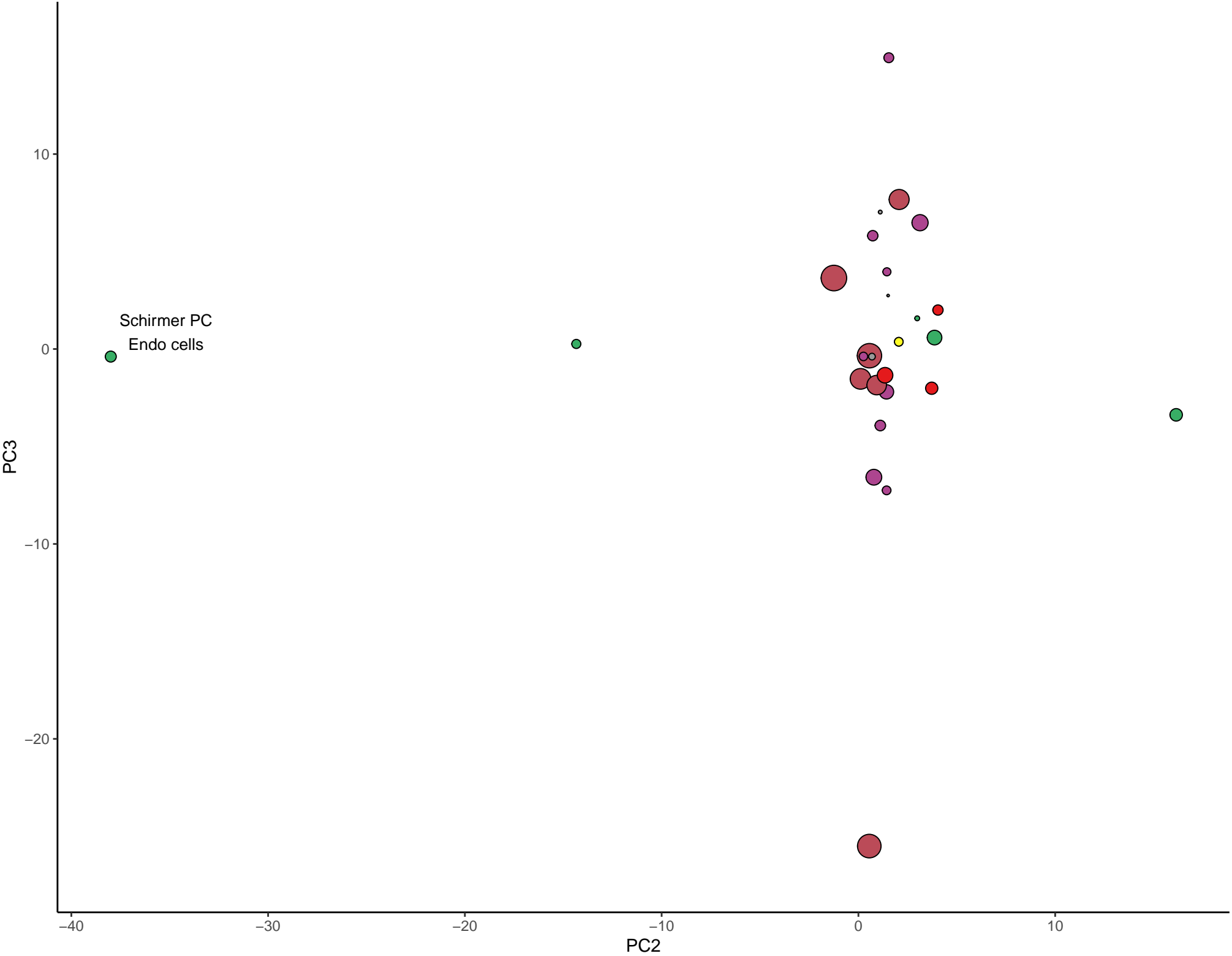
• 400 • 700 • 1,000 • 1,800 • 4,700

Study

- | | | |
|--------------|-------------|----------------|
| ABI 2019 | Hodge 2018 | Schirmer 2019 |
| Bakken 2019 | Lake 2018 | Velmeshev 2019 |
| Franjic 2022 | Li 2018 | Yang 2021 |
| Fullard 2021 | Luo 2019 | |
| Grubman 2019 | Mathys 2019 | |
| Habib 2017 | Nagy 2020 | |

Endothelial Batch Corrected Mean Expression

17 cell types found in 25 datasets from 15 studies
8,722 genes



Median # Unique Genes

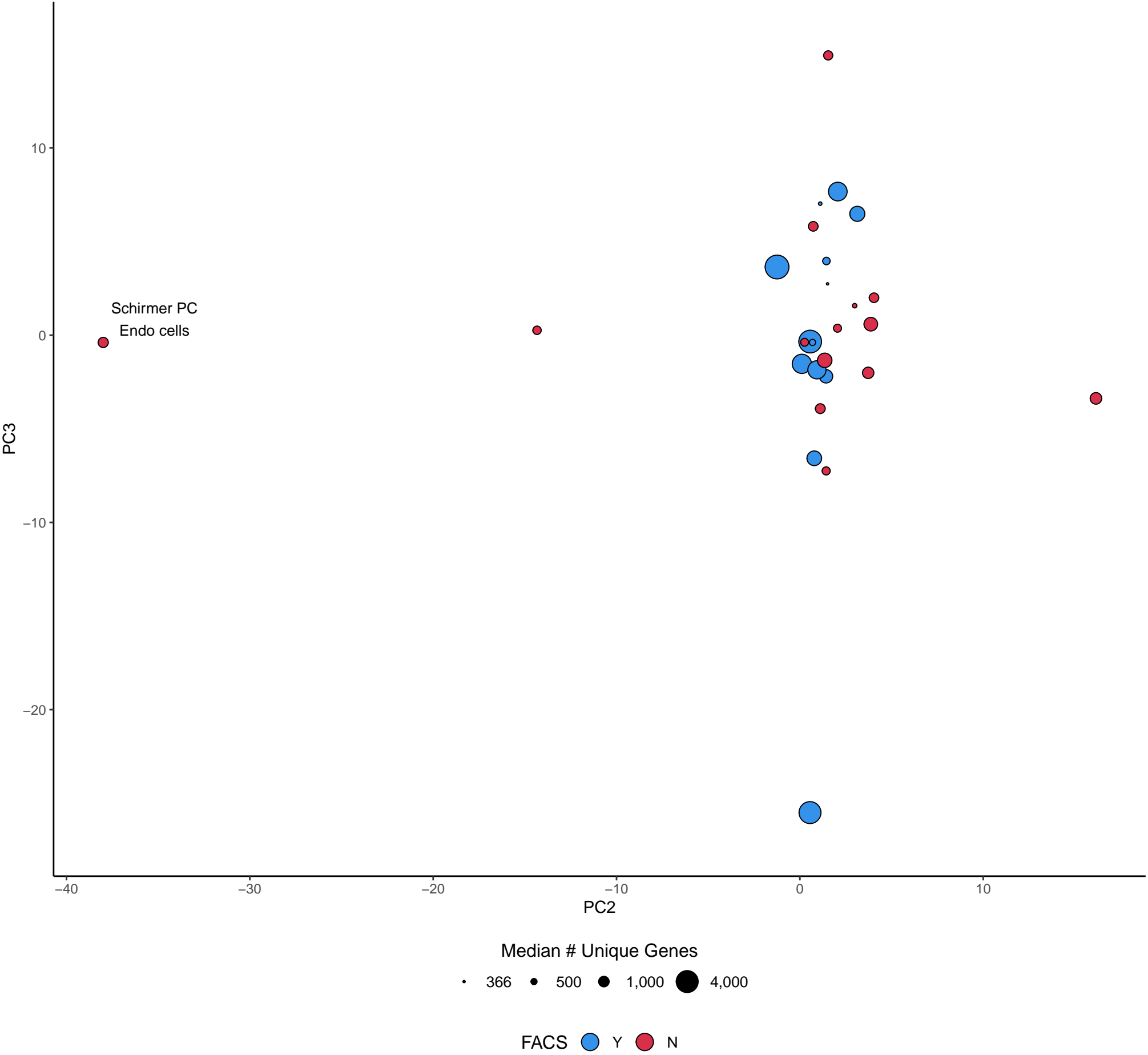
• 400 • 1,000 • 3,000 • 4,000

Platform

10x Chromium	10x Chromium V3	SMART-Seq v4
10x Chromium V2	DroNc-seq	snDrop-seq

Endothelial Batch Corrected Mean Expression

17 cell types found in 25 datasets from 15 studies
8,722 genes



Endothelial Batch Corrected Mean Expression

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