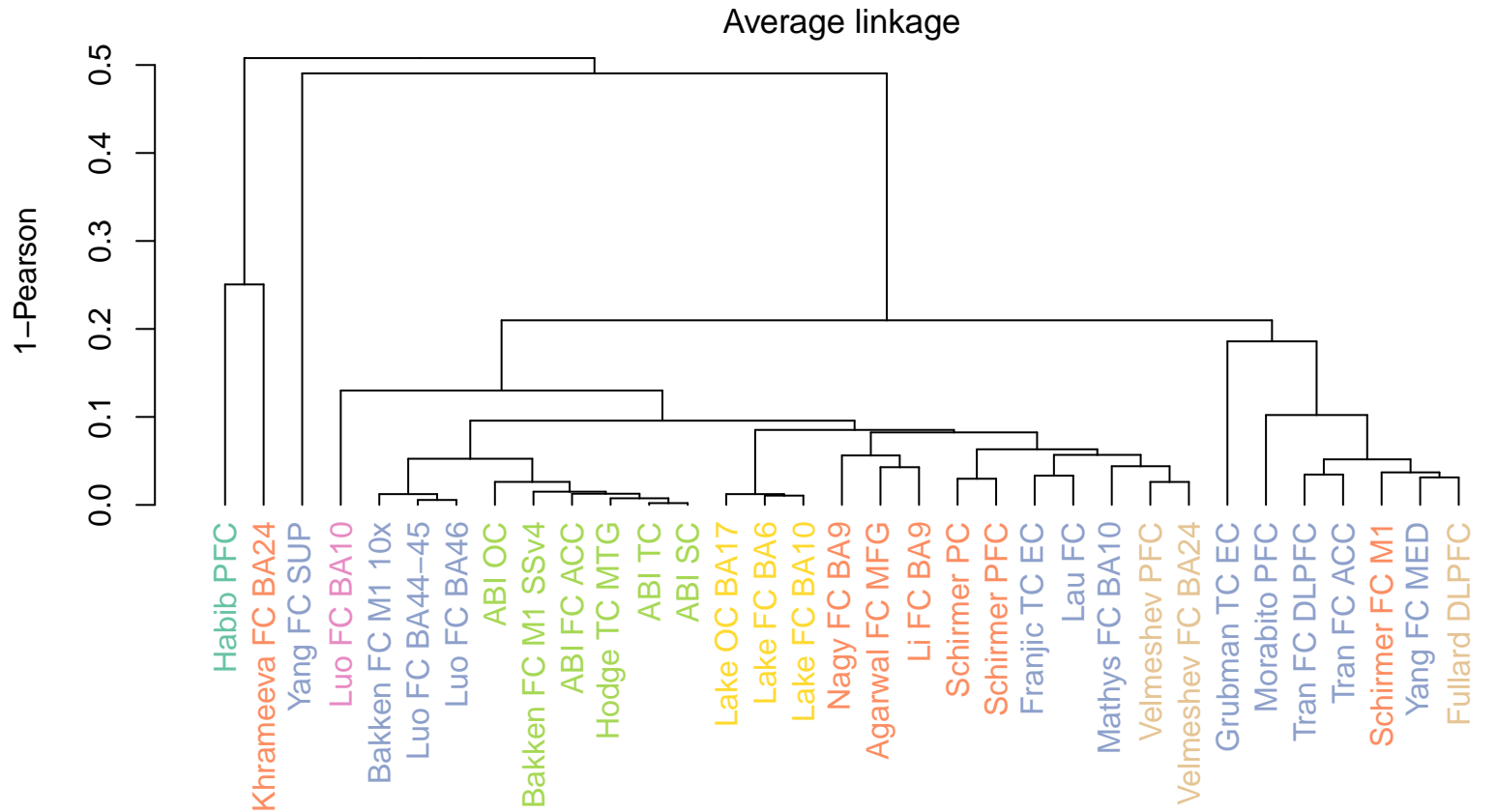


# Dataset Mean Expression Clustering

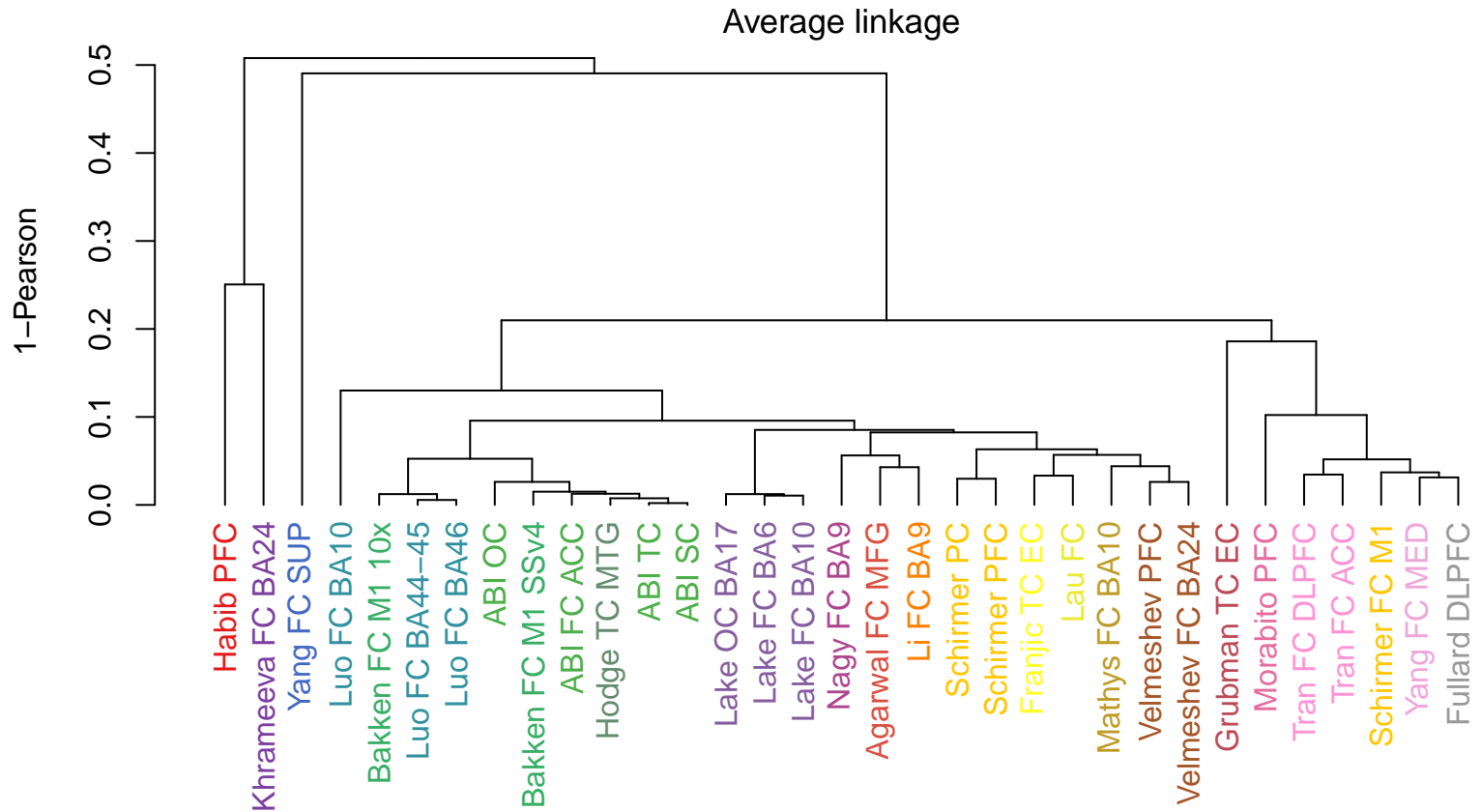


33 datasets from 21 studies  
8,596 protein coding genes

## Platform

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

# Dataset Mean Expression Clustering



33 datasets from 21 studies  
8,596 protein coding genes

## Study

- |                  |                |                 |                  |                |
|------------------|----------------|-----------------|------------------|----------------|
| ● Habib 2017     | ● ABI 2019     | ● Li 2018       | ● Velmeshev 2019 | ● Fullard 2021 |
| ● Khrameeva 2020 | ● Hodge 2018   | ● Schirmer 2019 | ● Grubman 2019   |                |
| ● Yang 2021      | ● Lake 2018    | ● Franjic 2022  | ● Morabito 2021  |                |
| ● Luo 2019       | ● Nagy 2020    | ● Lau 2020      | ● Tran 2020      |                |
| ● Bakken 2019    | ● Agarwal 2020 | ● Mathys 2019   | ● Yang 2021      |                |