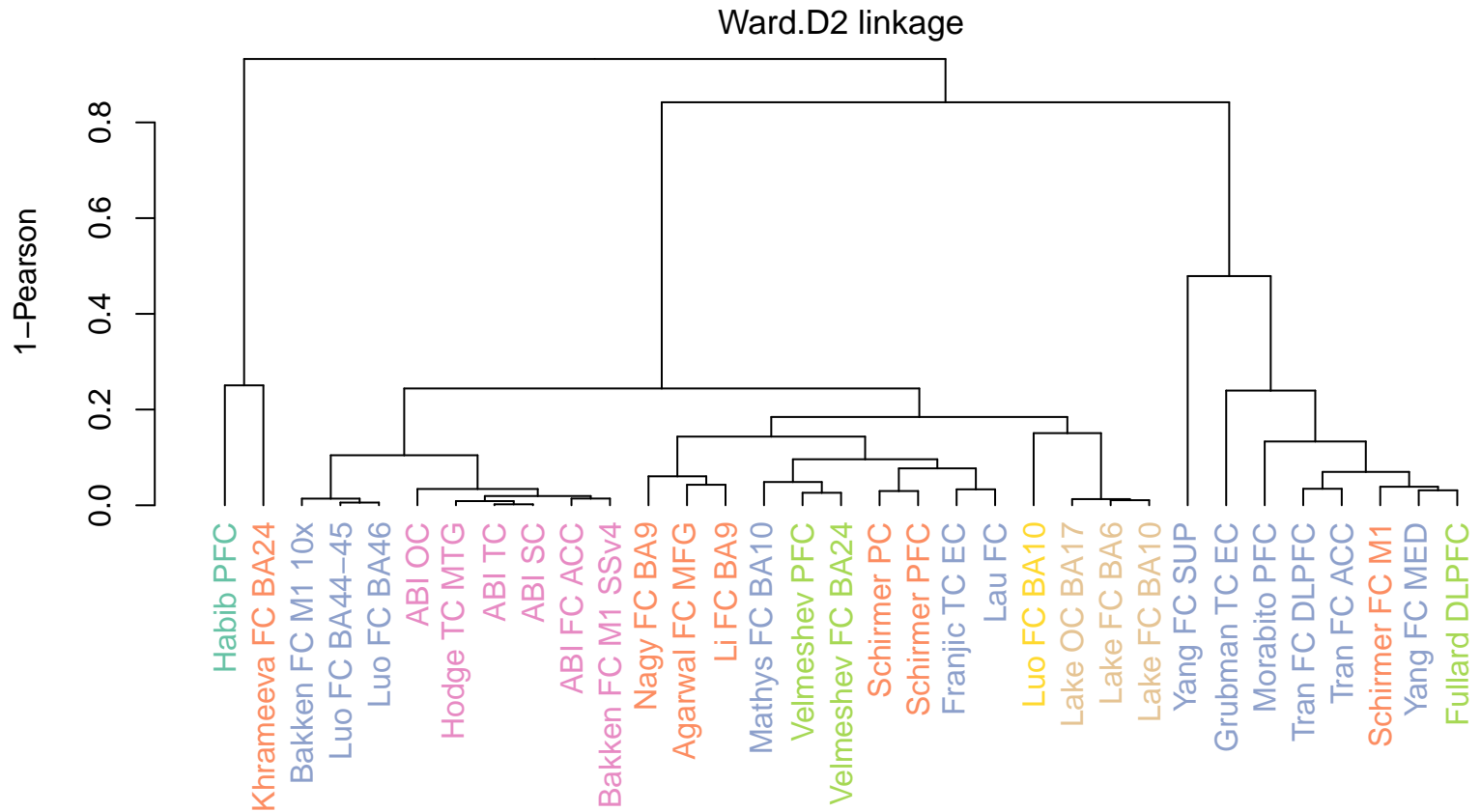


# Dataset Mean Expression Clustering

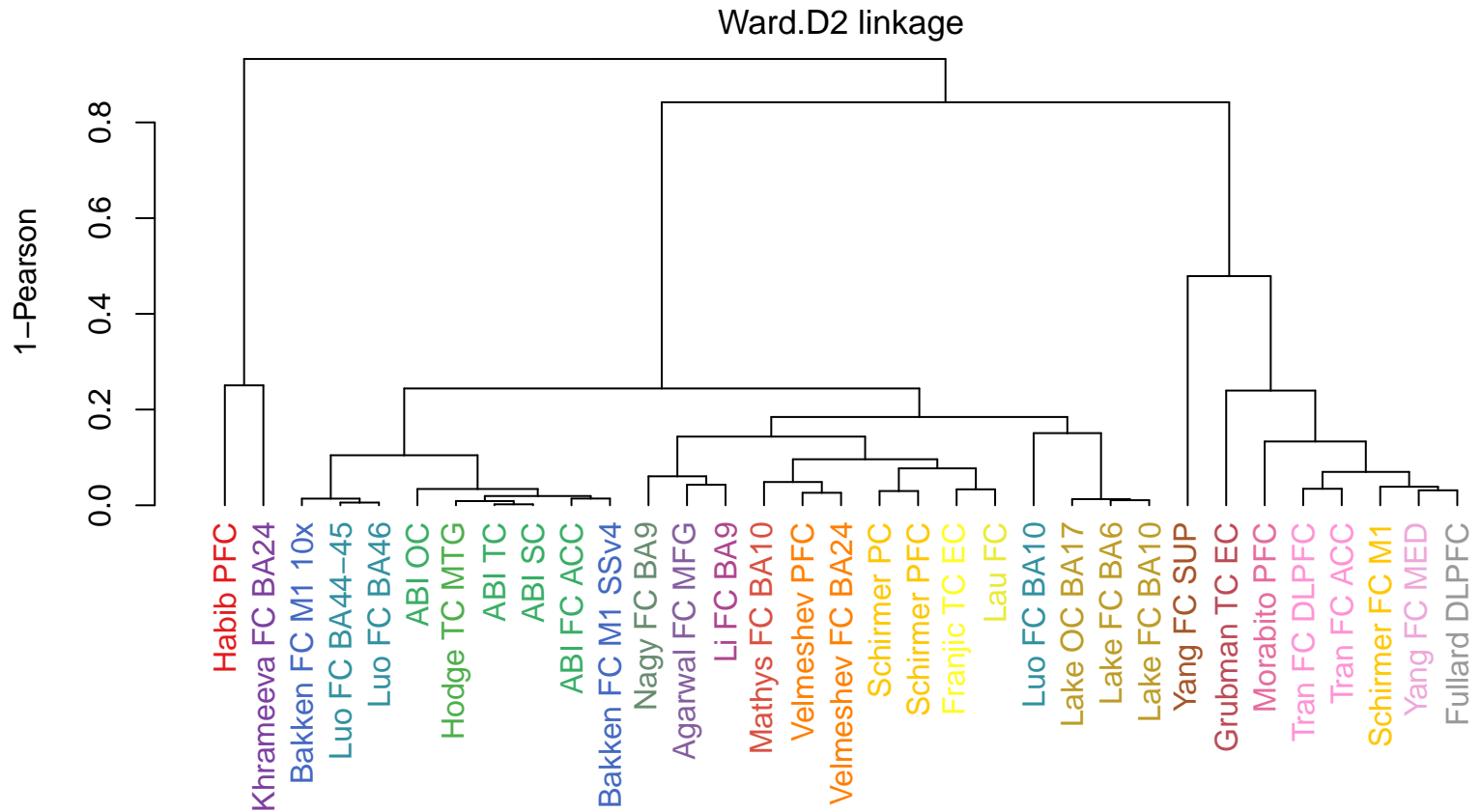


33 datasets from 21 studies  
8,596 protein coding genes

## Platform

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

# Dataset Mean Expression Clustering



33 datasets from 21 studies  
8,596 protein coding genes

## Study

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