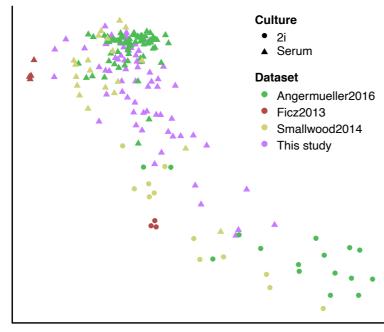
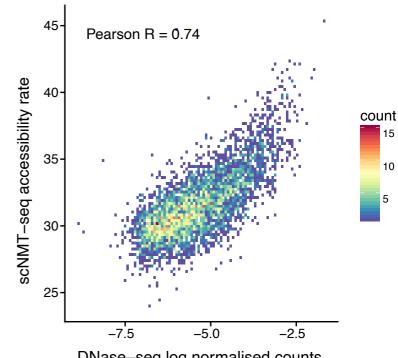
## **DNA** methylation



Principal Component 2 (4%)

Principal Component 1 (51%)

## Chromatin accessibility



DNase-seq log normalised counts