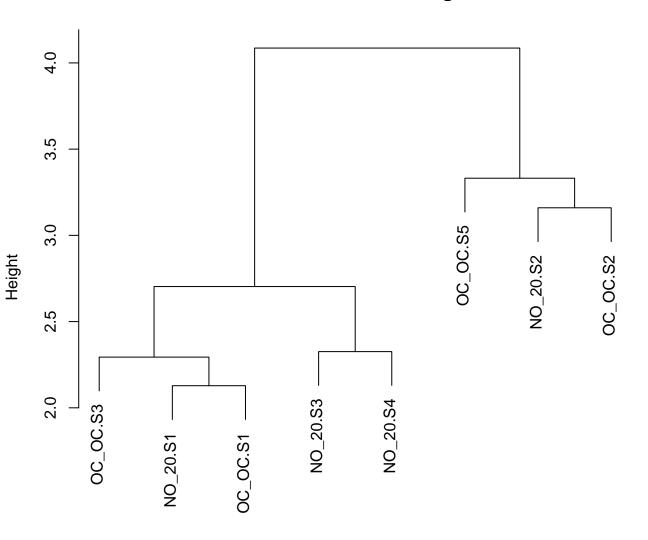
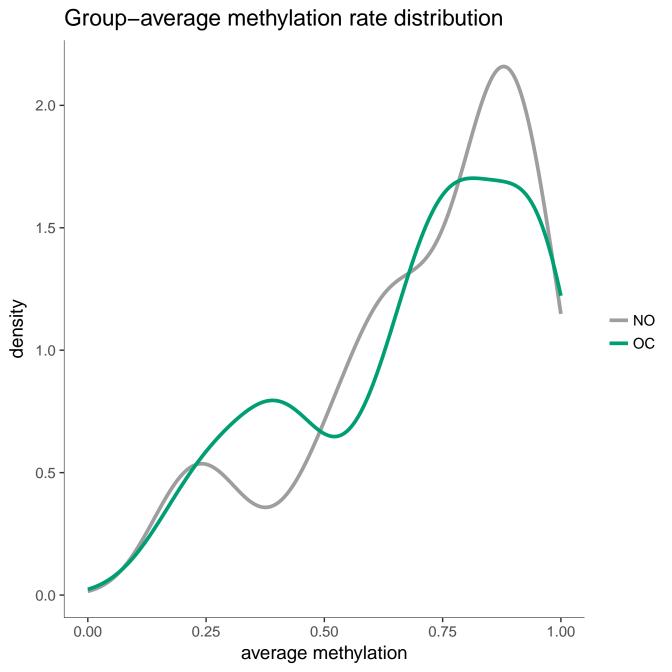


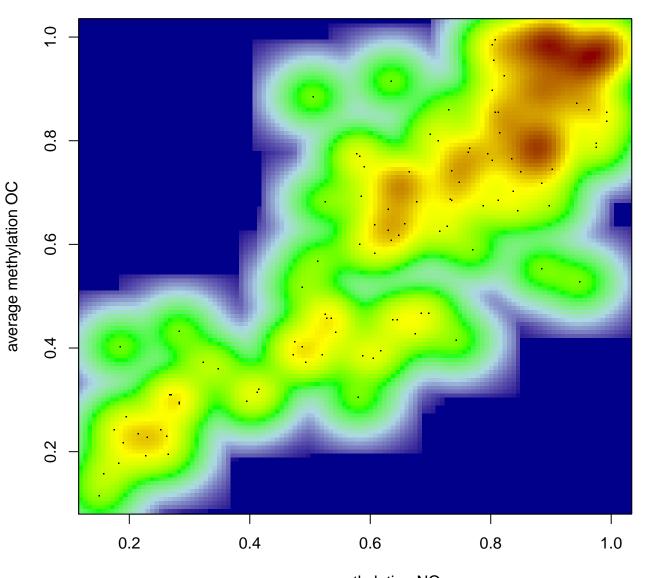
## **Hierachical Clustering**



Binned group-average methylation rate 150 bins 100 -[0,0.2)[0.2,0.4)[0.4,0.6)[0.6, 0.8)[0.8,1] 50 -0 -NO oc



## Scatter plot of group-average methylation rates



average methylation NO adjusted R-squared:0.7742

