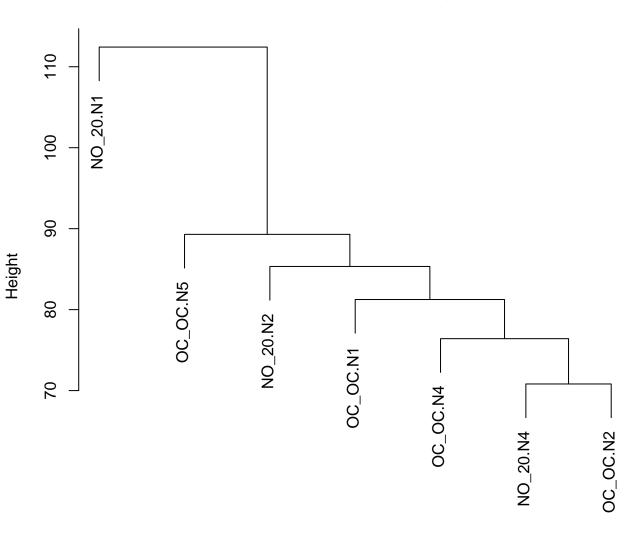


## **Hierachical Clustering**



Binned group-average methylation rate 150000 -100000 bins [0,0.2) [0.2,0.4) [0.4,0.6)[0.6, 0.8)[0.8,1] 50000 -0 -NO oc

Group-average methylation rate distribution 15 -10 density - NO OC 5 -0 -0.25 0.50 0.75 1.00 0.00 average methylation

## Scatter plot of group-average methylation rates

