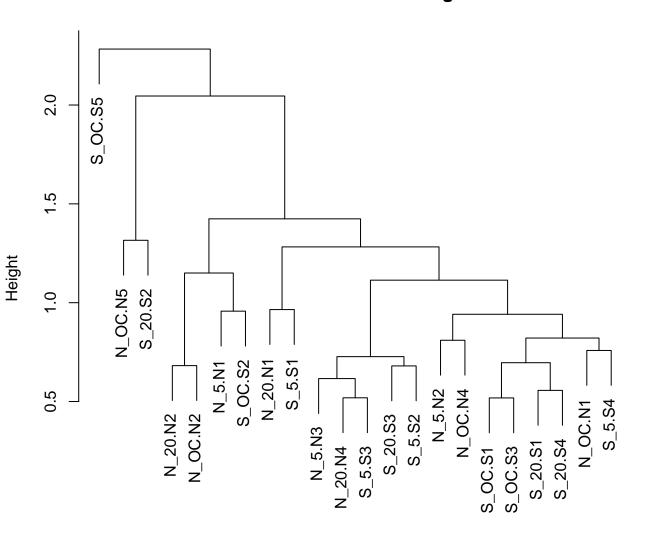
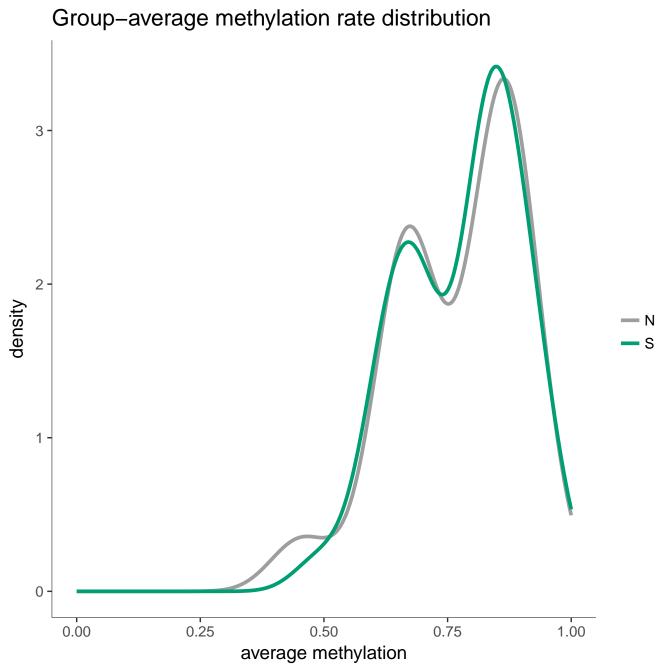


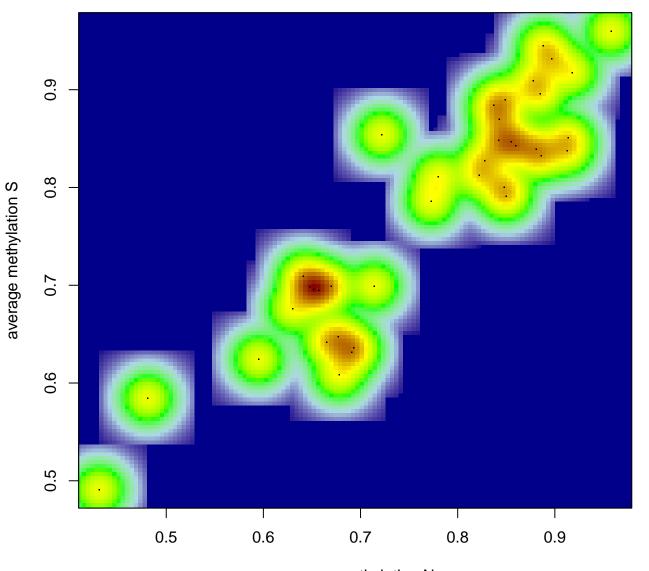
Hierachical Clustering



Binned group-average methylation rate 30 bins conut [0.4,0.6)[0.6, 0.8)[0.8,1] 10 -0 -Ň Ś



Scatter plot of group-average methylation rates



average methylation N adjusted R-squared:0.8462

