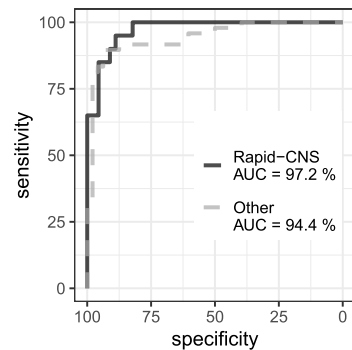
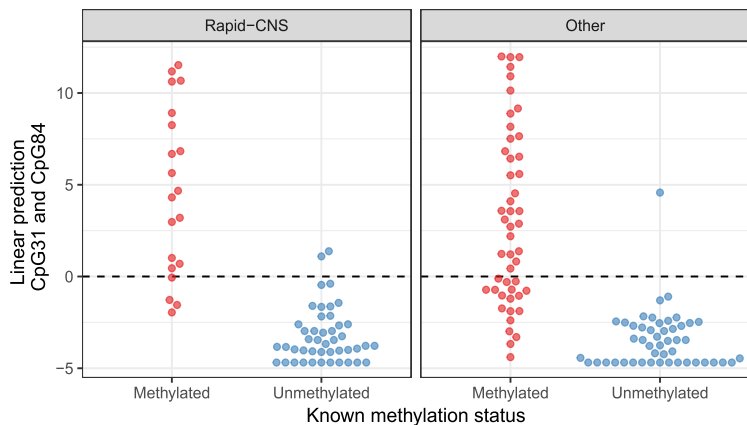


(a)



(b)



(c)

Figure 3: Comparison of nanopore sequencing and the *MGMT*-STOP27 classifier. (a) Methylation percentage of CpG 31 (probe ID: cg12434587) and CpG 84 (probe ID: cg12981137) in the *MGMT* promoter for 65 samples in the Rapid-CNS cohort (left) and remaining samples (right). Ellipses represent 90% confidence intervals. (b) ROC curves for a logistic regression model based on methylation of CpGs 31 and 84 by nanopore sequencing (Rapid-CNS) with classification by the *MGMT*-STOP27 classifier as operator (solid black) and prediction of *MGMT* status of 96 remaining samples (dashed grey). (c) Separation of samples based on the linear predictors extracted from the logistic regression model.