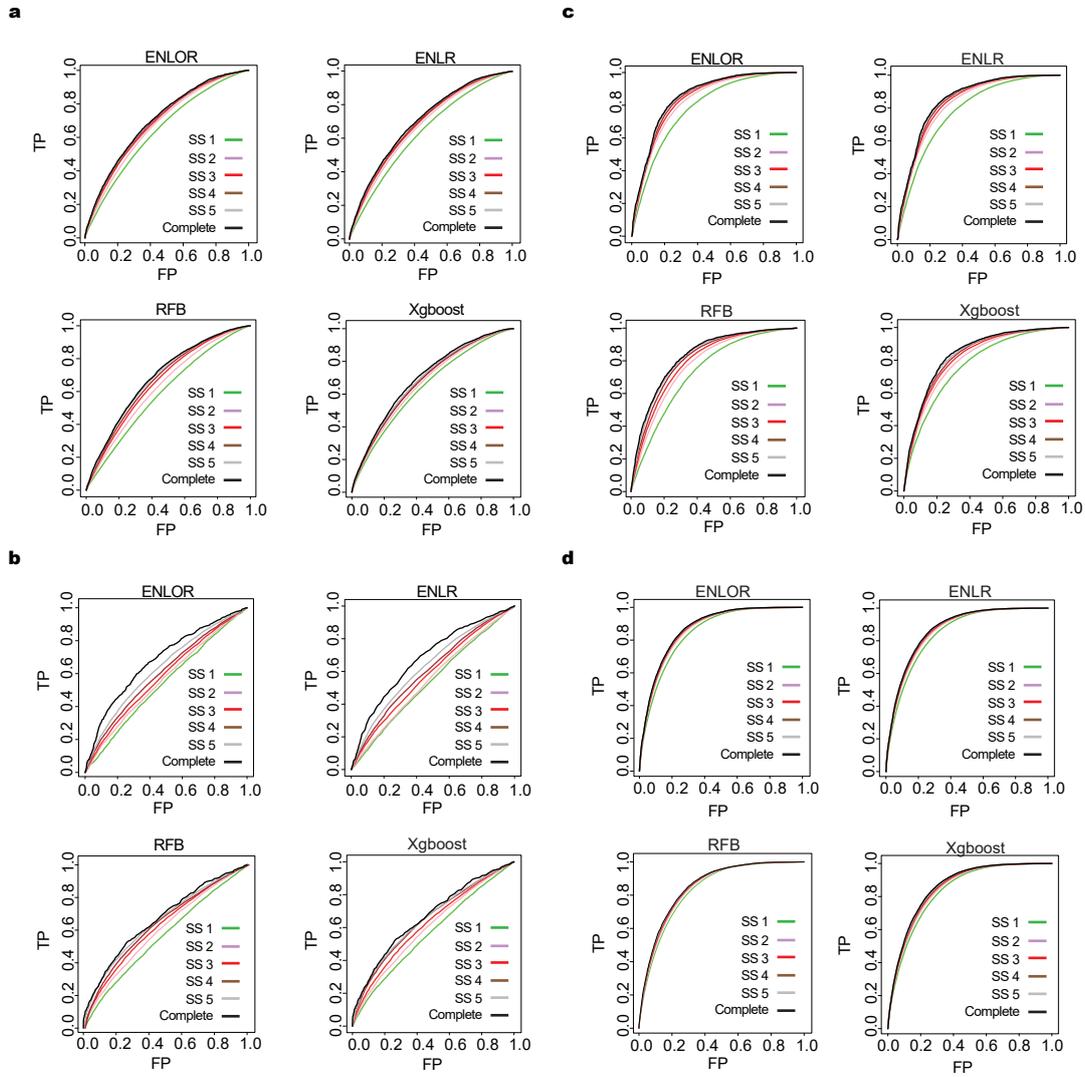


Supplementary Figure 3



Supplementary Figure 3: Performance analysis using subsets of the training dataset. a-d ROC curves depicting the algorithm performance on each subset (SS1..5) as well as the complete training datasets for CRISPRi (**a**), CRISPRa (**b**), Cas9 (**c**), and Cas12a (**d**). The *x-axis* represents the average False Positive rate (FP) whereas the *y-axis* represents the average True Positive rate (TP)