



Supplementary figure 3: ROC curves for One-class classification using SVM and L1 “3 or more” strategy, treating the 4000 random regions as training negatives, treating (A) the Personalis deletion calls and (B) the 1000 Genomes deletion calls as testing positives and treating the 2306 random regions as testing negatives. See original data at <https://plot.ly/266/~parikhhm/>, and <https://plot.ly/274/~parikhhm/>.