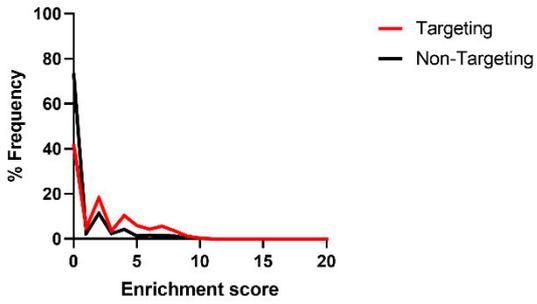
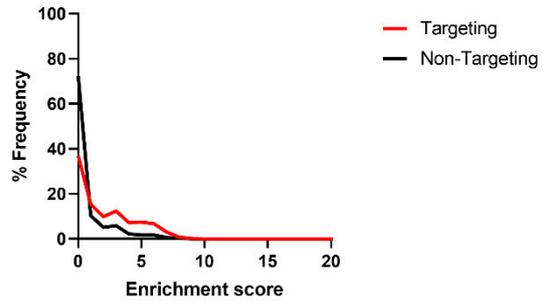


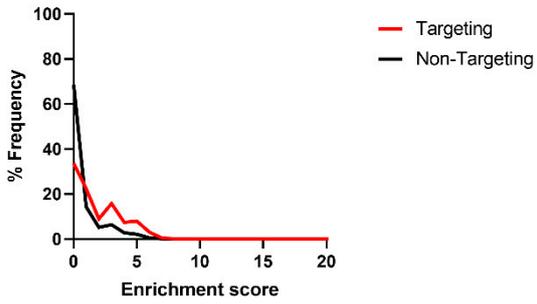
Sample no. 1



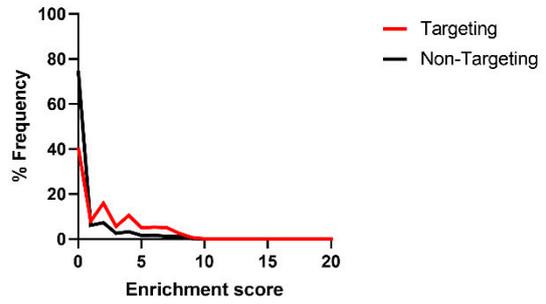
Sample no. 2



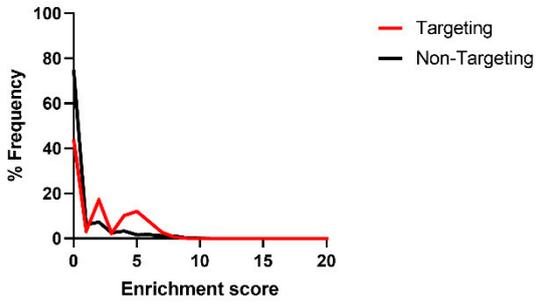
Sample no. 3



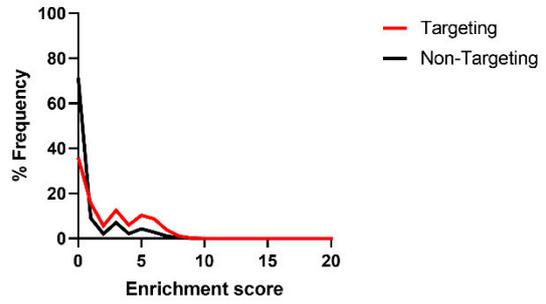
Sample no. 4



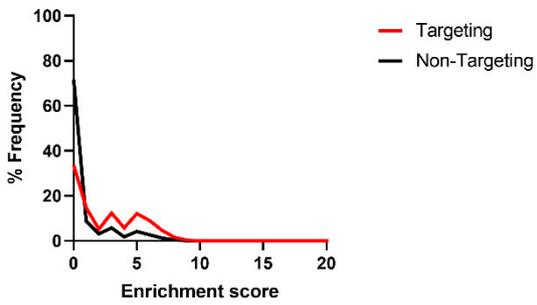
Sample no. 5



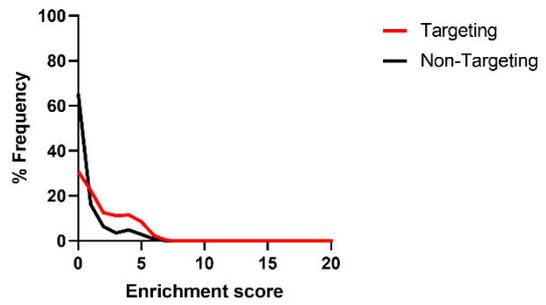
Sample no. 6

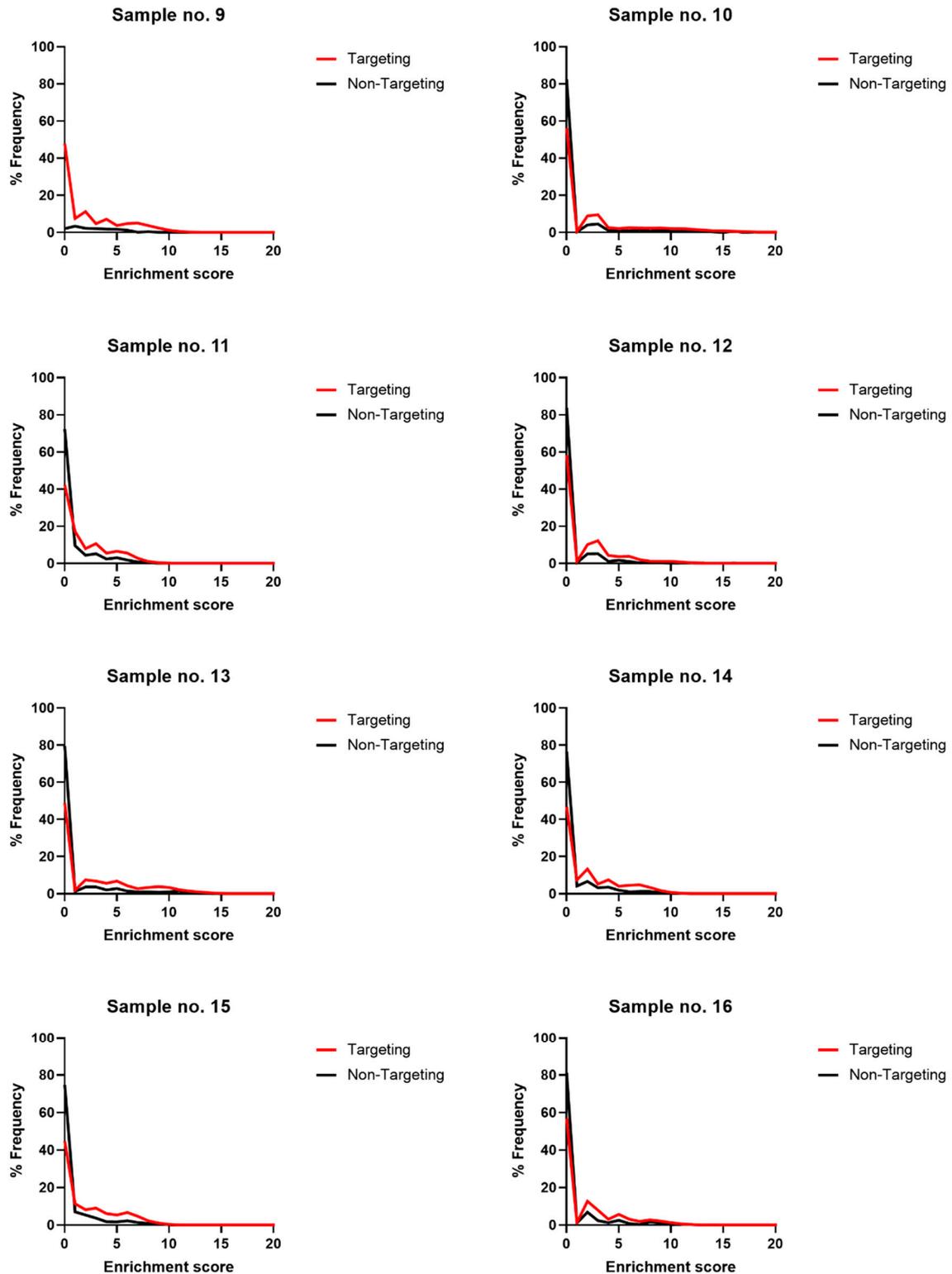


Sample no. 7



Sample no. 8





Supplementary Figure S1. Panels show density plots of normalized abundances for all sixteen samples for targeted (red) and non-targeted (black) sgRNAs, respectively, proving normalization performance and showing that for all cases there is a set of genes where enrichment score (x-axis) exceeds levels for non-targeting controls