Supplementary Table 1. Sequencing quality metrics and run summary statistics for targeted amplicon sequencing of CRISPR-edited PBMCs.

Sample	Library Type	Platform	Read Length (bp)	Total Reads	Total Bases (bp)	GC (%)	Q20 (%)	Q30 (%)
CRISPR-edited PBMCs	Amplicon, paired-end	Illumina	2×301	811,358	244,218,758	46.1	71.9	67.6