



Figure S4 | Spacer length and PAM distributions for the Lens (chromosome) CRISPR-Cas system. Experimental set-up is as described for supplemental figure 1. All data is the average of three biological replicates. **A)** The distribution of spacer length acquired from the targeted plasmid compared to the wild-type CRISPR-Cas array (n = 64 spacers in the wild-type array). **B)** Quantification of the PAMs for the new protospacers in a stacked bar plot.