



**Figure S3 | Spacer length and PAM distributions for the Lens (pLPL) 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 lengths acquired from the targeted plasmid (grey) compared to the wild-type CRISPR-Cas array (n = 53 spacers in the wild-type array). **B)** Quantification of the PAMs for the new protospacers are shown in a stacked bar plot; the (+) strand PAMs are denoted by the hatched bars and the (-) strand PAMs are denoted by the grey bars.