



FIGURE S5. LX-miR HeLa Screen Replicate 2. (A) Scatter plot of the induced versus control read counts for each sgRNA in replicate 2. The dotted line indicates no change between the populations. (B) The kernel density estimate of the \log_2 fold change between induced and control reads per sgRNA in replicate 2. The green line indicates the distribution of the control non-targeting sgRNAs and the blue line indicates the distribution of the miRNA-targeting sgRNAs.