

These files contain the sequencing results from a CRISPR Cas9 screen across 5 cell lines grown in culture or as tumors. `zimmerman_5763` contains 2 folders corresponding to the results of 2 sequencing runs. `Zimmerman_5736_190619A5` corresponds to the first run.

`Zimmerman_5736_190620A5` corresponds to the second.

`SampleBarcodeIndexes.xlsx` contains two sheets which contain a key for which sequence barcodes represent which samples.

`sgRNALibrarySequence.xlsx` contains the sgRNA sequences found in the library.

`RawsgRNACounts.xlsx` contains the processed data used in the manuscript.