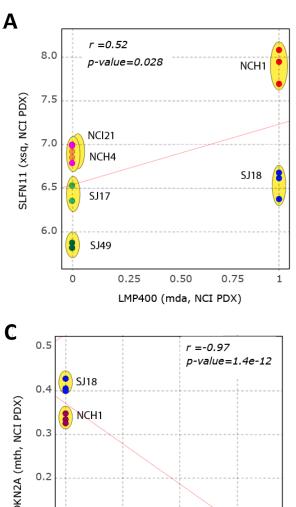
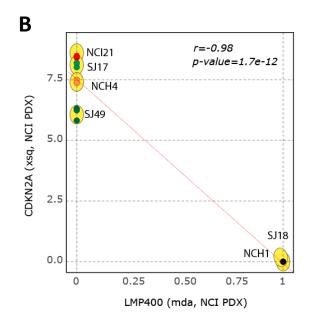
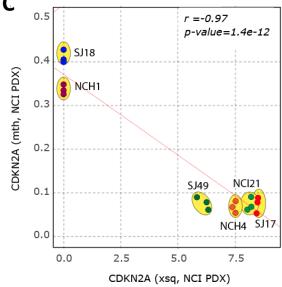
Supplemental Figure S7







Supplemental Figure S7. Representative snapshots of Ewing Sarcoma MinerCDB (https://discover.nci.nih.gov/rsconnect/EwingSarcomaMinerCDB) showing data reproducibility across PDX and insights into SLFN11 and CDKN2A gene expression and LMP400 response. A. Expression of *SLFN11* as a function of LMP400 response. **B.** Expression of *CDKN2A* as a function of LMP400 response (1 and 0 denote >30% and <30% response). C. Epigenetic inactivation of CDKN2A in the NCH1 and SJ18 PDXs. Clustering of the PDX triplicates is indicated as yellow elliptical highlights. Statistical analyses (Pearson correlation coefficients and p-values are annotated with the regression lines shown in red).