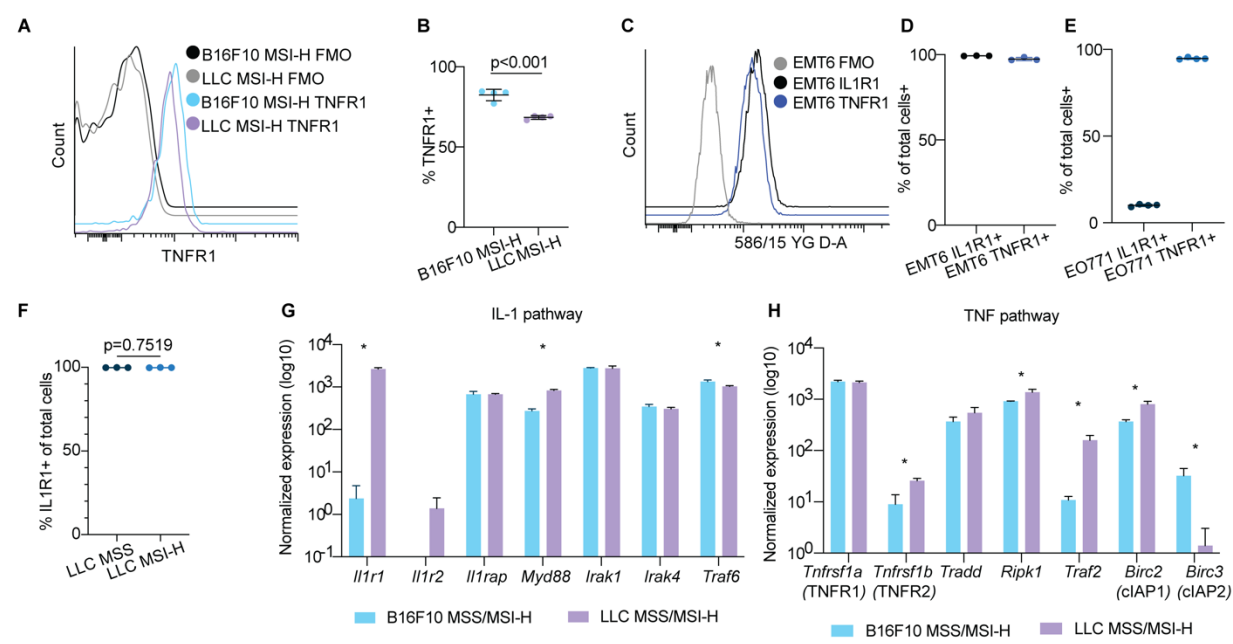


Figure S9



Supplementary Figure S9.

A, Representative histogram of TNFR1 fluorescence staining intensity for indicated cell lines. **B**, Quantification of % TNFR1+ population in indicated cell lines, from one experiment with n=4 biological replicates. Two independent experiments were performed. P value by two-tailed t test. **C**, Representative histogram of IL1R1 and TNFR1 fluorescence staining intensity for EMT6 cell line. **D**, Quantification of % IL1R1+ and % TNFR1+ populations for EMT6 cells, n=3 biological replicates from one representative experiment. Two independent experiments were performed. **E**, Quantification of % IL1R1+ and % TNFR1+ populations for EO771 MSI-H cells, n=4 biological replicates from one representative experiment. Two independent experiments were performed. **F**, Quantification of % IL1R1+ populations for LLC MSS vs. LLC MSI-H cells. n=3 biological replicates from one representative experiment. p value by t test. Two independent experiments were performed. **G**, Quantification of normalized transcript counts for indicated genes belonging to the IL-1 receptor signaling pathway for the indicated cell lines, n=4 biological replicates per condition. Asterisks indicate $p < 0.05$ by Mann-Whitney test. **H**, Quantification as in **(G)** for genes in the TNF receptor signaling pathway. For all applicable panels, error bars represent mean $\pm$ SD.