

Figure S1

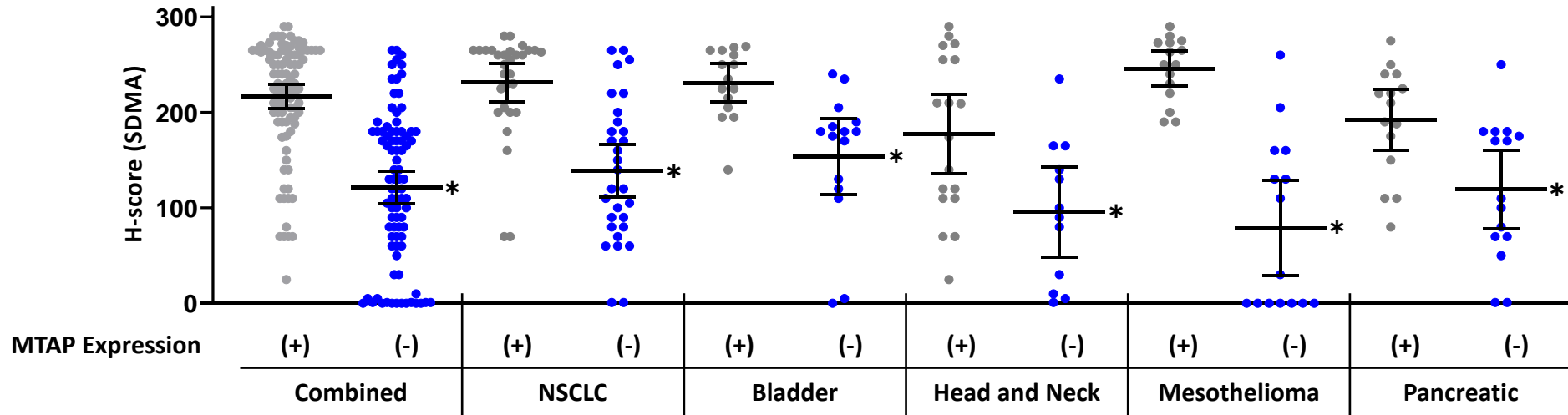


Figure S1. Neoplastic SDMA protein expression levels by immunohistochemistry H-scoring in neoplastic MTAP expressing and neoplastic MTAP non-expressing tumors in tissue microarrays (TMAs) across several cancer types known to have a high percentage of *MTAP* del. Cores from TMAs with no MTAP protein expression in neoplastic cells were identified and compared to a similar number of cores with MTAP protein expression in neoplastic cells and analyzed for neoplastic SDMA H-scores. All cases are shown on the left of the plot and indication-specific analyses are broken out to the right. NSCLC consists of lung adenocarcinoma and lung squamous cell carcinoma cases. Summary statistics shown as mean +/- 95% confidence interval. Statistics were determined using a two-tailed Student's t-test with significance indicated (*, $p < 0.05$).