



Supplementary Figure 9. Enrichment for SV breakpoints at sites of PRDM9-associated recombination hotspots in samples expressing *PRDM9*, as shown by odds ratios > 1 for each cancer type, and excluding SV breakpoints falling in Alu or THE1 repeat elements (blue: $p < 0.05$ with Bonferonni correction for the number of cancer types tested; bars indicate 95% confidence intervals).