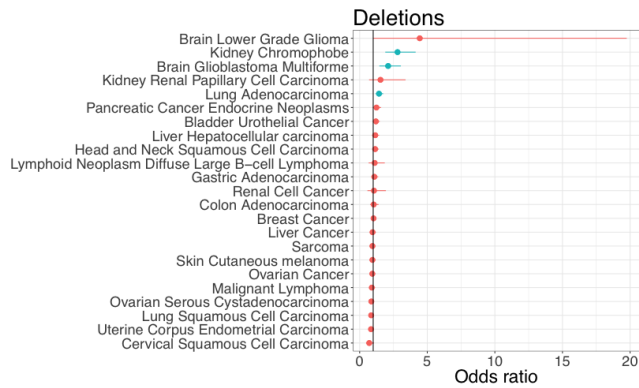
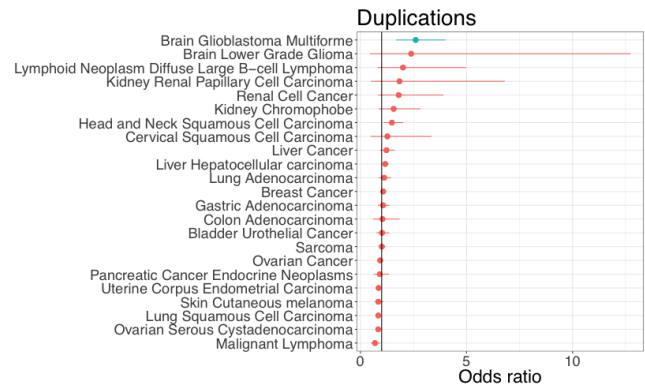
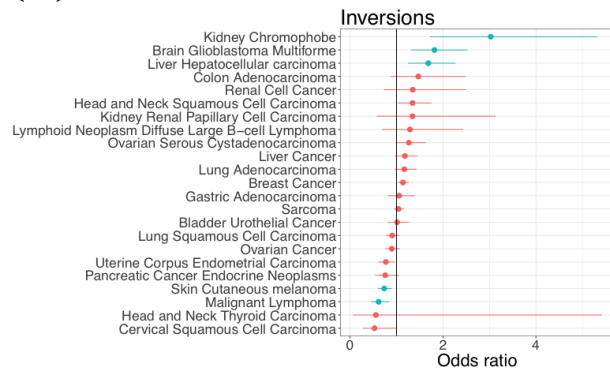
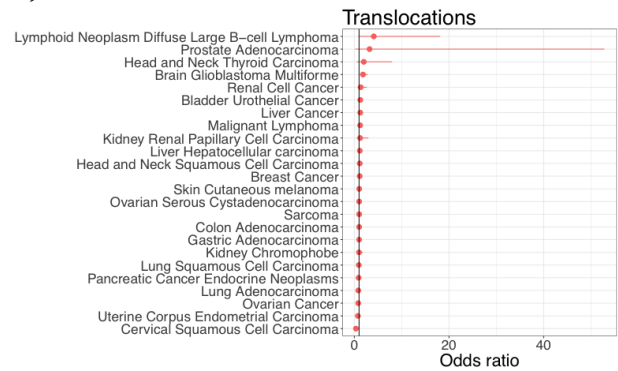


**(A)****(B)****(C)****(D)**

**Supplementary Figure 10.** Enrichment for each SV, namely deletions (A), duplications (B), inversions (C), and translocations (D), at sites of PRDM9-associated recombination hotspots in samples expressing *PRDM9*, as shown by odds ratios > 1 for each cancer type (blue:  $p < 0.05$  with Bonferroni correction for multiple testing for the number of cancers tested; bars indicate 95% confidence intervals).