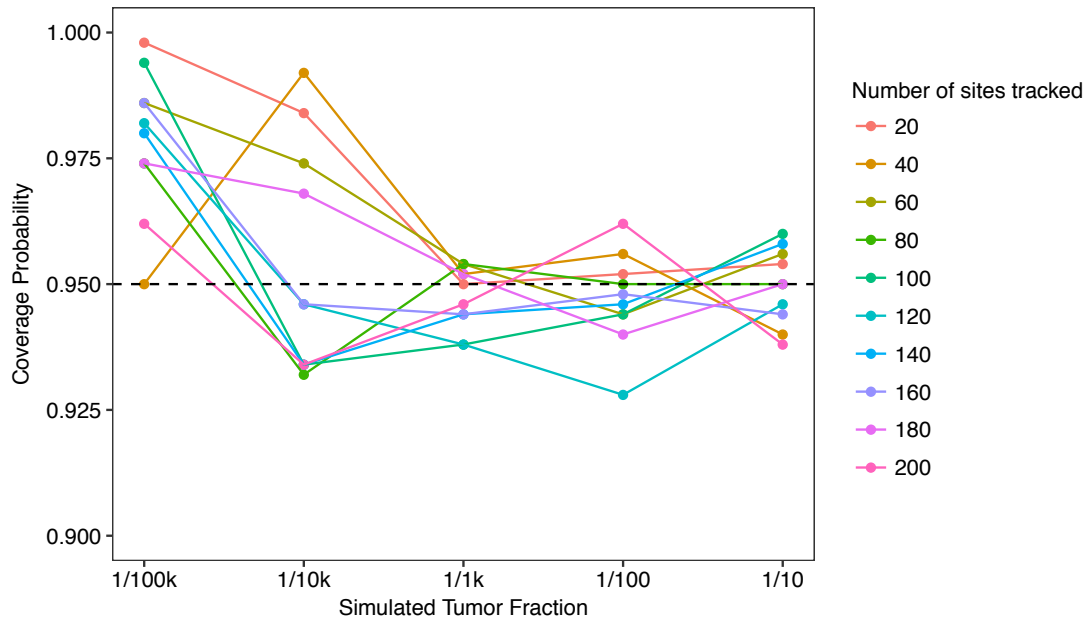
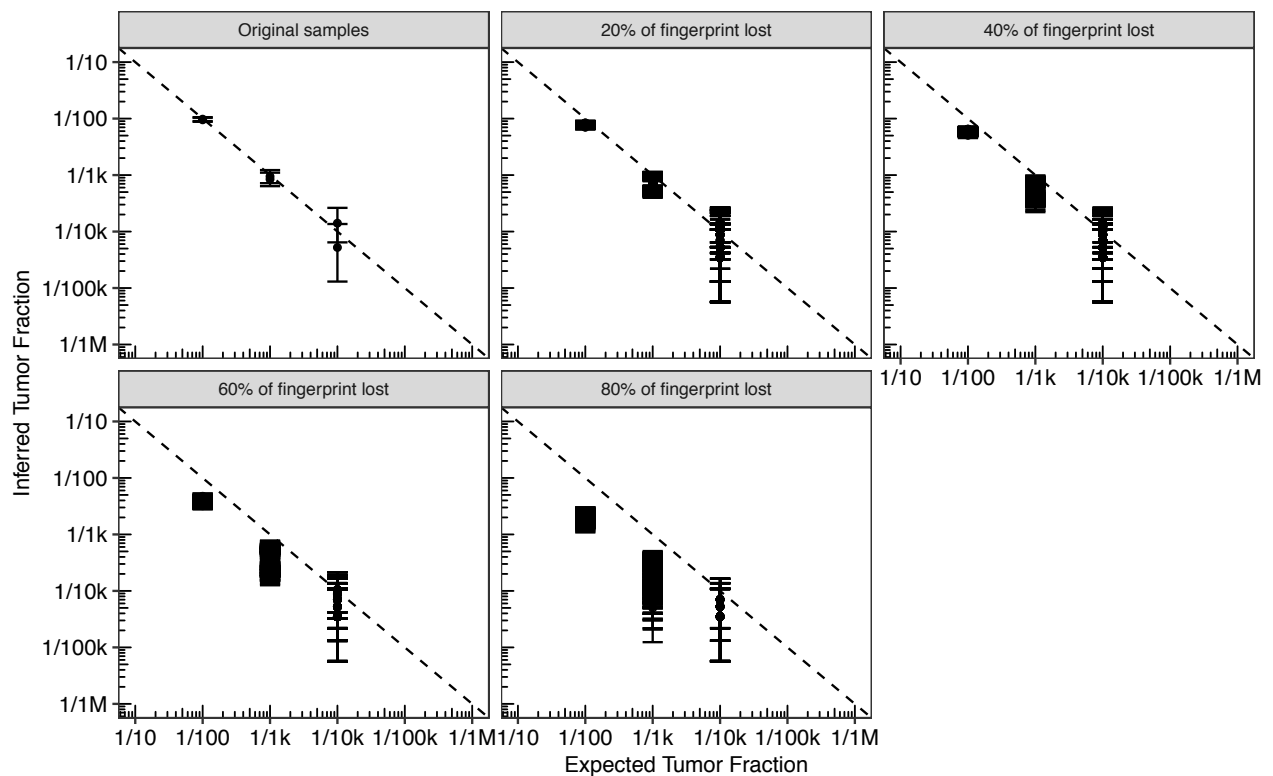


A**B**

Supplementary Figure 6: Validation of tumor fraction confidence interval and effects of biological variability on tumor fraction inference. (A) Confidence intervals inferred from simulated data were aggregated into coverage probability for each condition and compared against theoretical coverage probability of 95% (see Methods). (B) Biological variability of clonal evolution loss was simulated by removing tumor evidence from a fraction of the fingerprint panel. Tumor fraction was calculated for simulated and original samples for comparison.