

Expanded View Figures

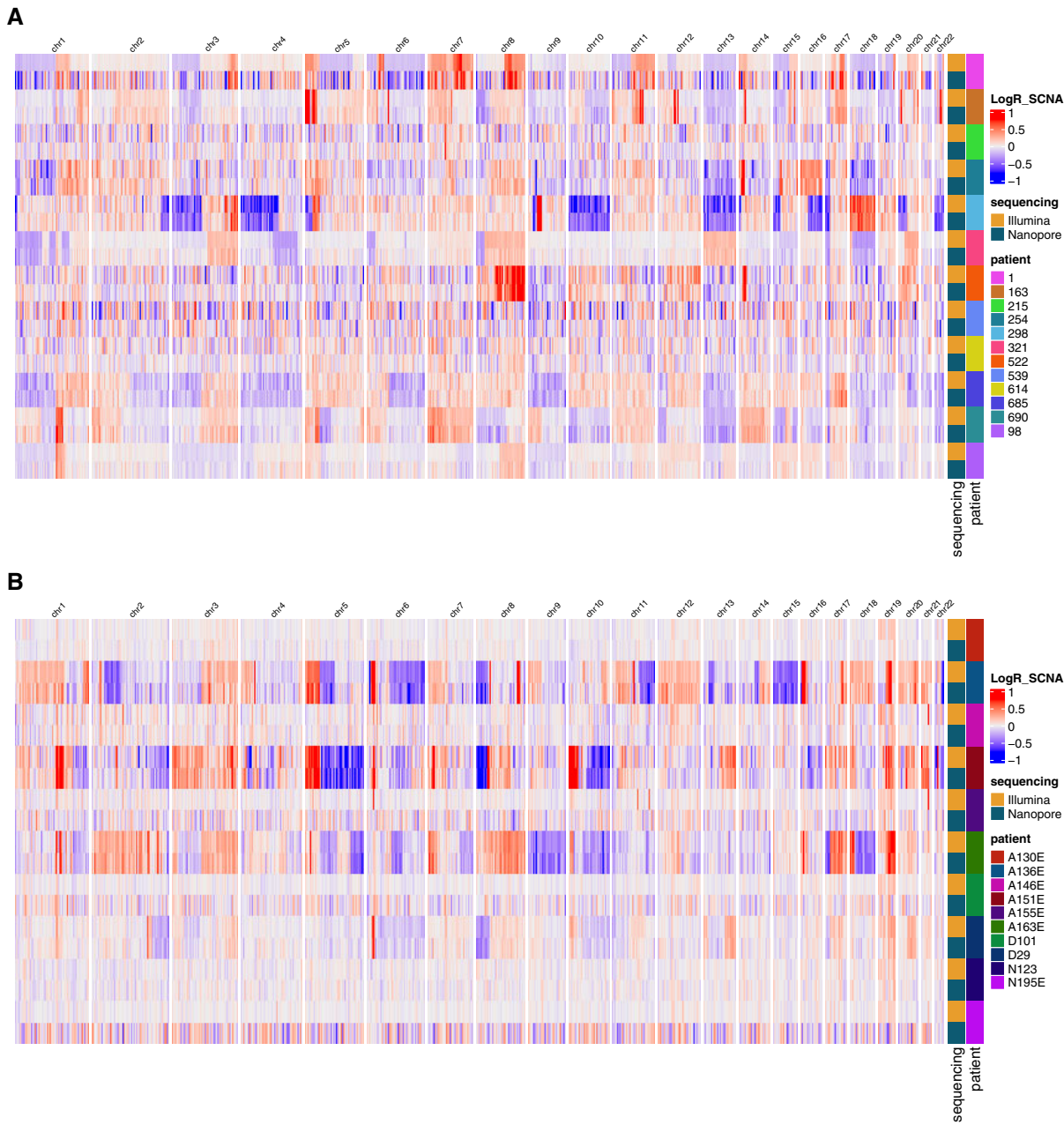


Figure EV1. Copy number aberration heatmaps for the samples included in this study.

A, B (A) Late-stage lung plasma samples and (B) urine samples. Illumina sequencing is downsampled randomly to the same number of reads as the nanopore data.

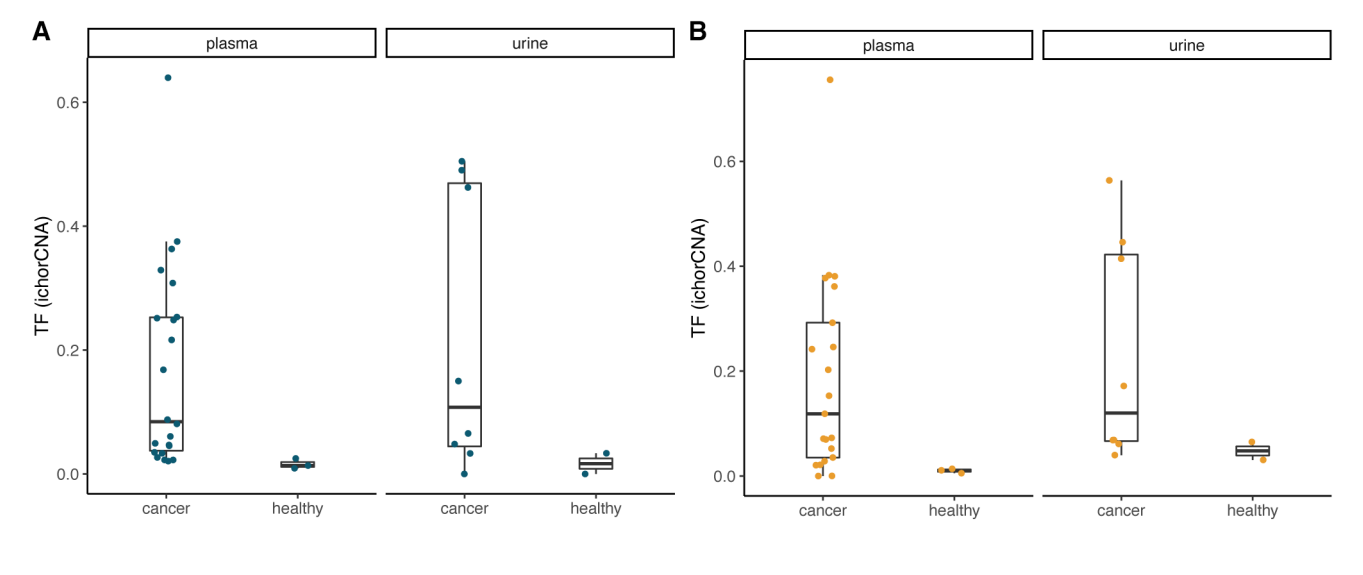


Figure EV2. Tumor fraction by ichorCNA for the cancer and healthy samples included in this study.

A, B (A) From the nanopore data and (B) from the Illumina data.