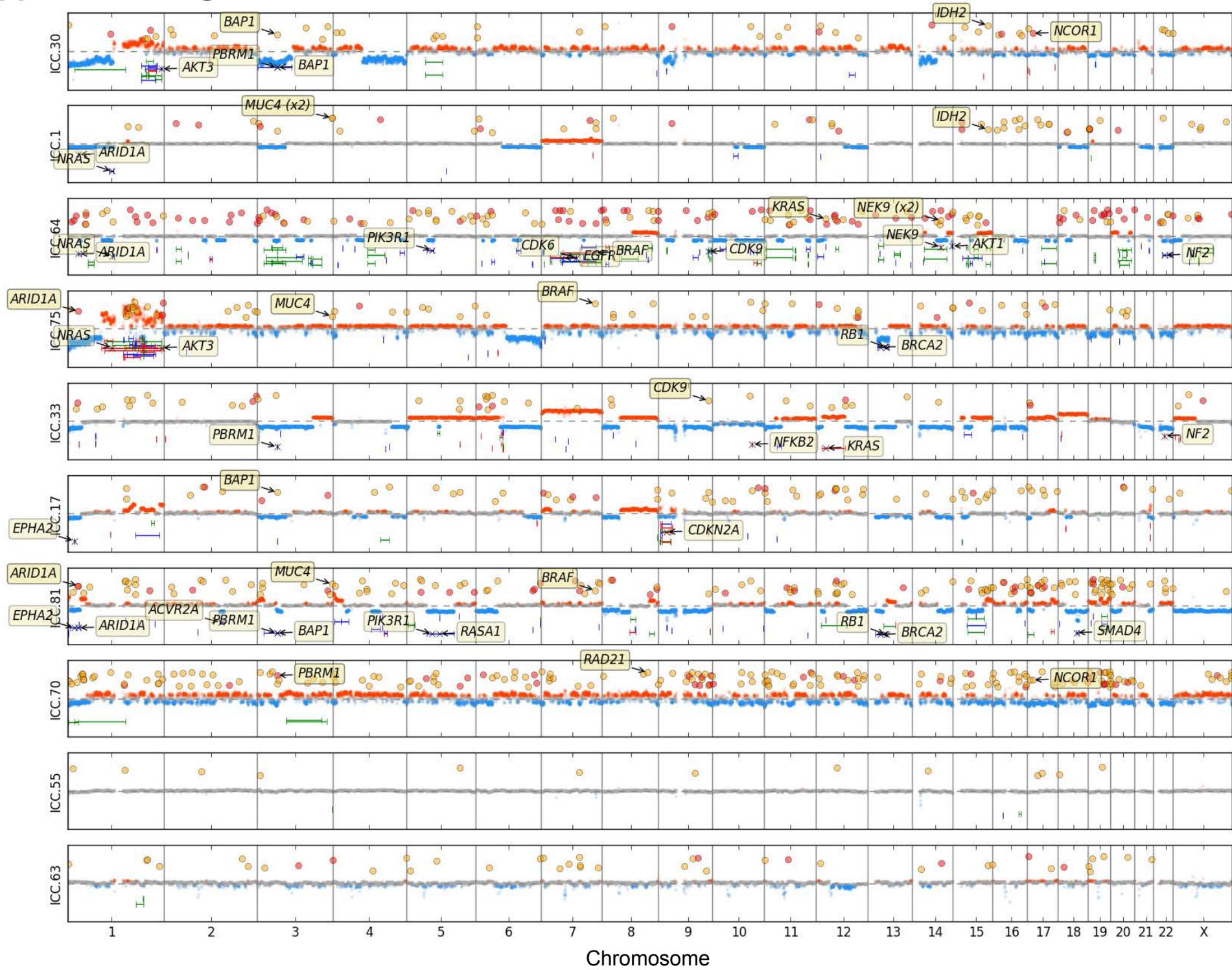


# Supplemental Figure S7



### **Supplemental Figure S7. Whole-genome profiling of ICC samples.**

For each genome, the CNV profile is shown over the dashed horizontal center line. Red CNV points indicate 150kbp bins over a log<sub>10</sub>-fold change of 0.05 and blue points indicate a decrease below a log<sub>10</sub>-fold change of -0.05. Somatic point mutations (SNVs and INDELS) are shown above the horizontal center line, yellow points indicate moderate-impact variants (e.g. non-synonymous changes, in-frame indels) and red points indicate high-impact variants (e.g. frameshifts and stop-gain mutations) as defined by snpEff. Somatic intra-chromosomal structural mutations are shown below the horizontal center line as line segments: blue segments represent deletions, red are duplications, and green are inversions. The vertical positioning of point and structural mutations and is done for ease of visualization and has no biological significance. Selected genes (those mentioned in the text) are highlighted with yellow text boxes and arrows.