SUPPLEMENTARY FIGURE 1 LEGEND.

Genome-wide array CGH plots for two representative self vs. self hybridization experiments. Red and blue tracings each represents separate components of a single dye-swap experiment. An LCV is identified if both red and blue tracings deviate in opposite directions beyond the established thresholds. A single false positive was identified in the second experiment, yielding for these two experiments, a false positive rate of 1 in 5264 clones.