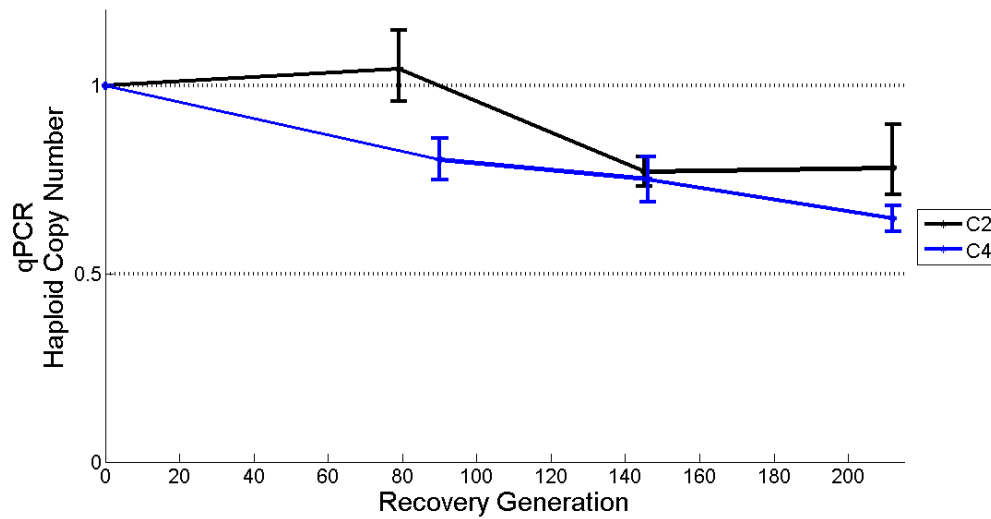


**Additional File 7: Suppl Figure S5**



**Supplemental Figure S5.** Increase in the frequencies of parallel deletion events in two control populations, C2 and C4, containing an overlapping region on Chromosome I. The average copy-number per haploid genome was calculated from qPCR results and is indicated on the vertical axis. The number of recovery generations is indicated on the horizontal axis. The results show a strong decline in average copy-number of these two independent deletions that were initially detected by oaCGH. The deletions have reached fixation when the average copy-number has reached 0.