
Chromosome evolution screens recapitulate tissue-specific tumor aneuploidy patterns

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Supplementary Table 1 Acquired mutations during *in vitro* evolution experiments across all sequenced clones.

Supplementary Table 2 Mutations present in parental hTERT-HMEC population, from which all clones were derived.

Supplementary Table 3 All SVs acquired during *in vitro* evolution experiments across all sequenced clones.

Supplementary Table 4 Curated Notch Activation and Repression gene sets.

Supplementary Movie 1 Simulation of Notch-ON/Notch-OFF pattern formation in 40x40 cell lattice. Left panel: WT 1q control mono-culture experiment (50% WT 1q + 50% WT 1q homogenous population). Middle panel: WT 1q vs +1q co-culture experiment (50% WT 1q + 50% +1q mixed population). Right panel: +1q mono-culture experiment (50% +1q + 50% +1q homogenous population)