Supplementary Figure S1. Genomic profiling and clinical history of late relapse patients.

a) *Bona fide* Breast cancer drivers

b) Late relapses – Clinical details

Supplementary Figure S1. Genomic profiling and clinical history of late relapse patients. a) High depth profiling of ER+ BC late relapses using a custom targeted panel. The heatmap shows the mutations of *bona fide* breast cancer drivers that passed the filter for Allele depth >=20, Alternate F1R2+F2R1 >= 4, Allele frequency >= 0.1 and consequence level of moderate or high. Time to relapse, recurrence in the dataset, allele frequency and relapse site are indicated across x and y axes. Significant genes are indicated based on dN/dS analysis from the q-value of neutrality test at gene level (*q_global_cv <= 0.1*) and confidence intervals for the dN/dS ratios per gene (*CI for missense and truncating mutations do not span through value of 1*). b) Table depicting the time to relapse (in years), site of relapse and cellularity (%) of late relapse samples (n=49).