

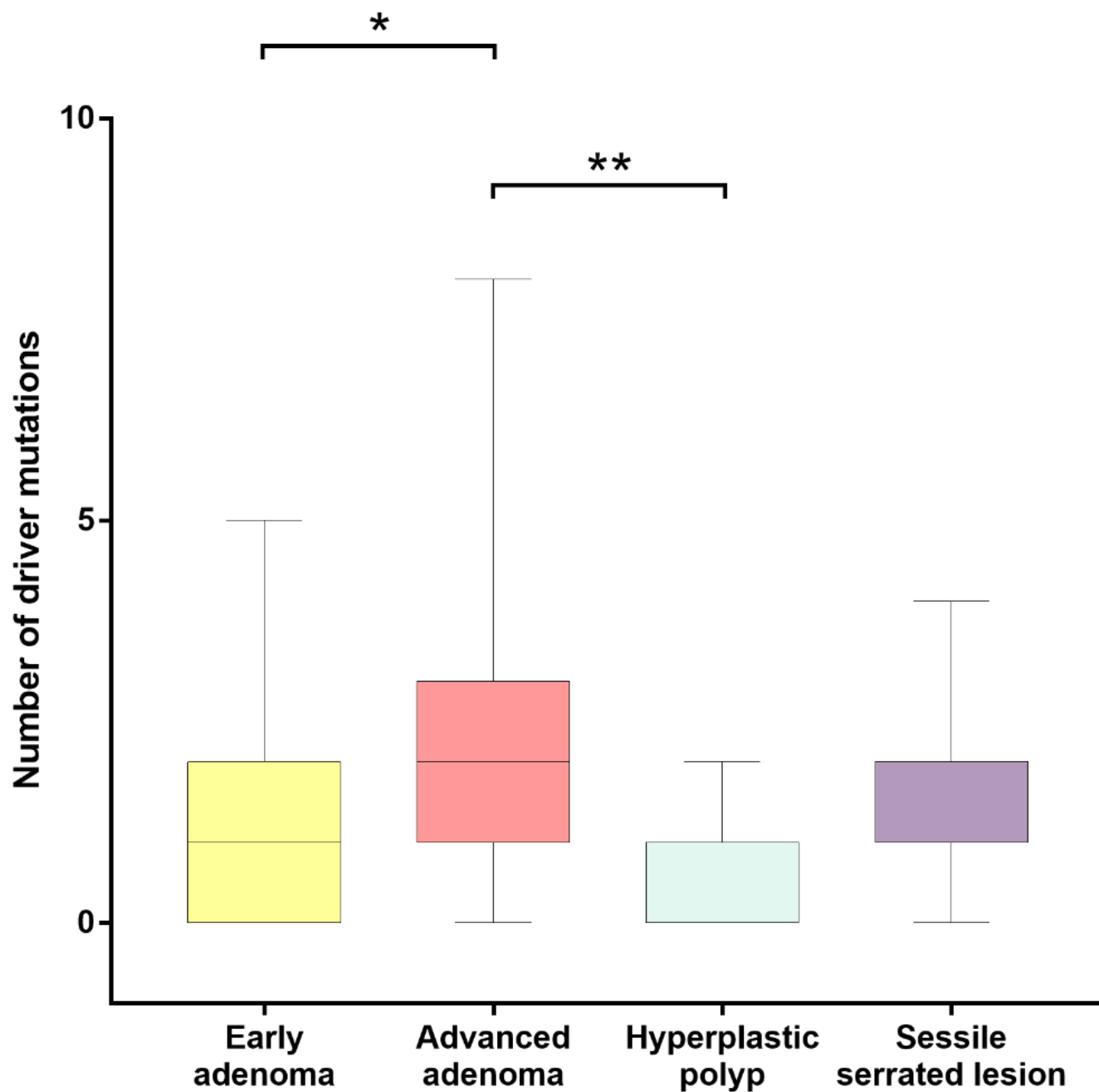


Figure S3. Number of driver mutations detected in the sequencing differs between the four classes of precursor lesions (Kruskal-Wallis test $p < 0.001$). The boxplot shows the median number of mutations observed across the early adenomas, advanced adenomas, hyperplastic polyps and sessile serrated lesion. Mann-Whitney test was used to determine the statistical significance with Bonferroni multiple comparisons correction: * $p < 0.05$; ** $p < 0.001$.