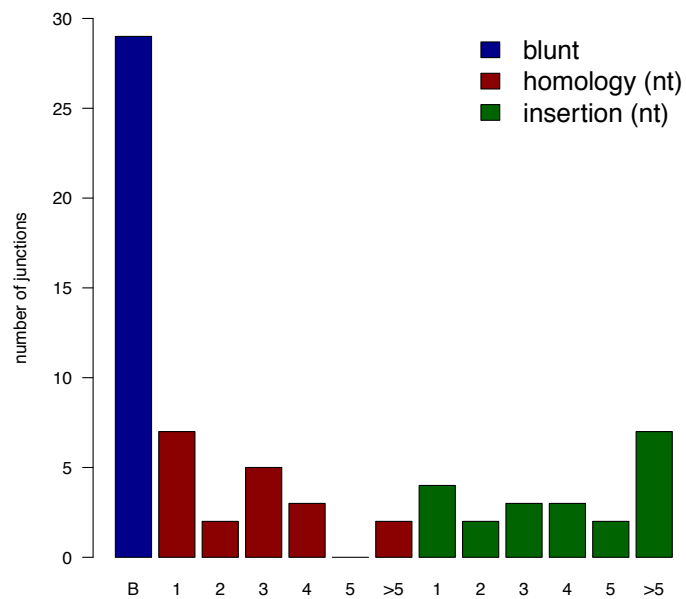
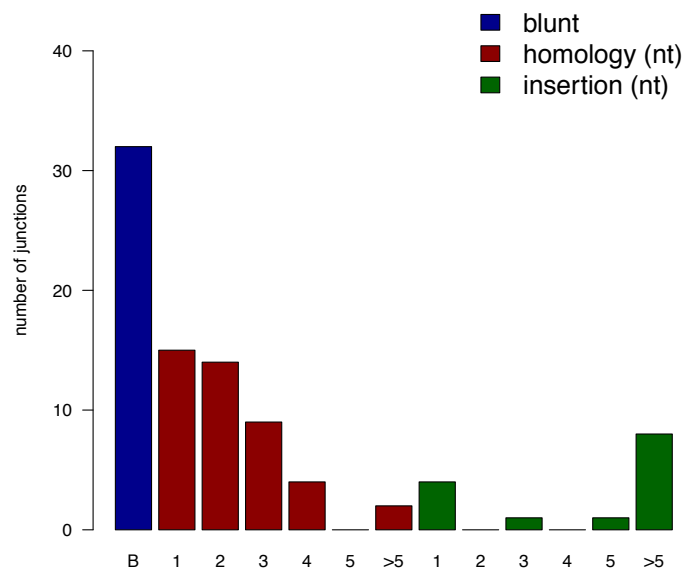


Additional data file 11

A



B



Sequence characteristics for 159 tumor-specific fusion points. (A) Characteristics of fusion points involving chromothripsis clusters (n=69). (B) Characteristics of fusion points outside of chromothripsis clusters (n=90). The distribution across different types of characteristics is similar for fusion points within or outside chromothripsis clusters. For the majority of fusion points we see no or very little microhomology or inserted sequences, which suggests that non-homologous end-joining accounted for the repair process.