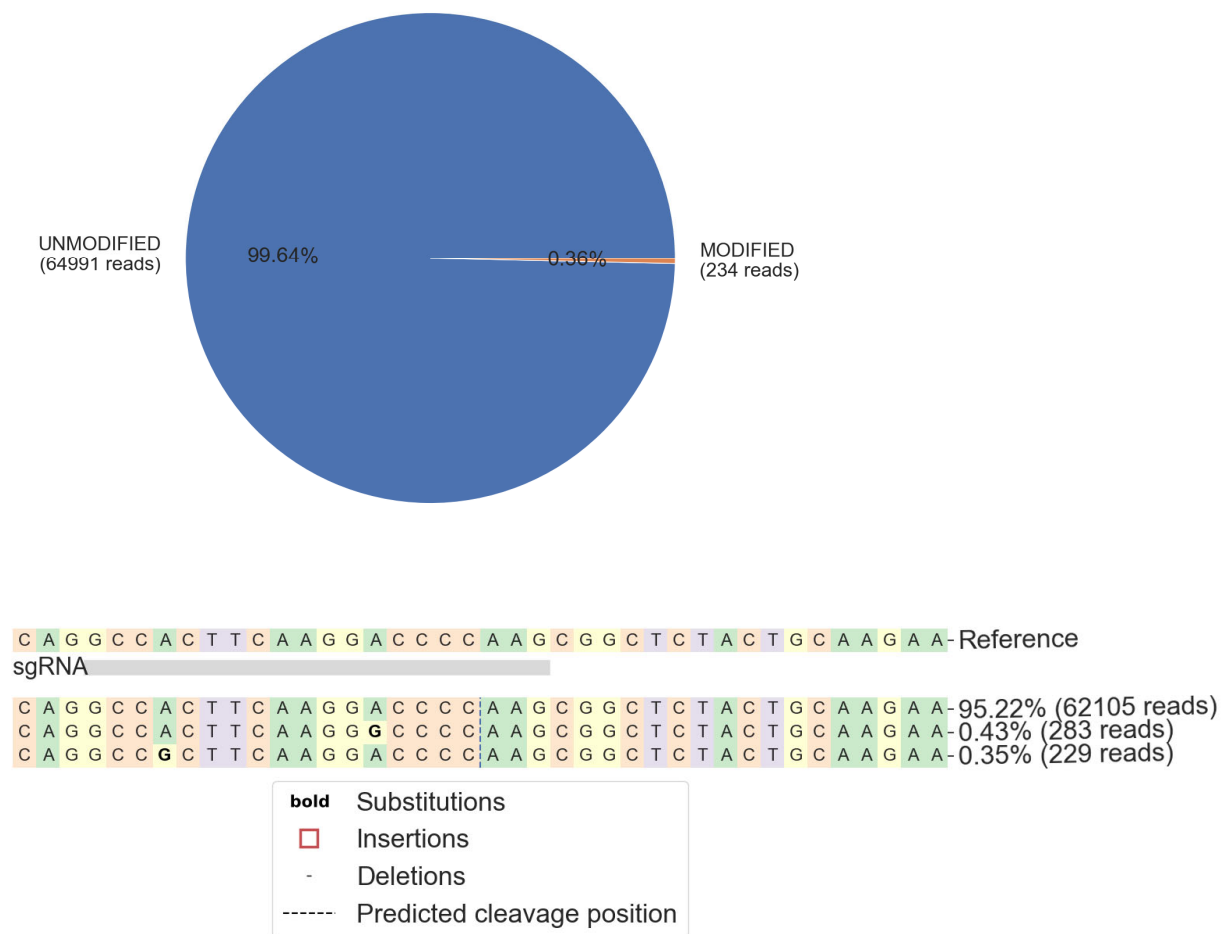




Supplemental Figure 3. Top panel, percentage of sequence modifications generated by the loxP sgRNA around the cleaving site of the Fgf2 sgRNA in skin fibroblasts. Bottom panel, sequence modifications and their read counts. Note that unlike indels induced by the sgRNA targeting Fgf2, changes detected here are substitutions, probably constituting base call errors during sequencing.