Supplementary Material

Genetic insights: High germline variant rate in an indigenous African cohort with early-onset colorectal cancer

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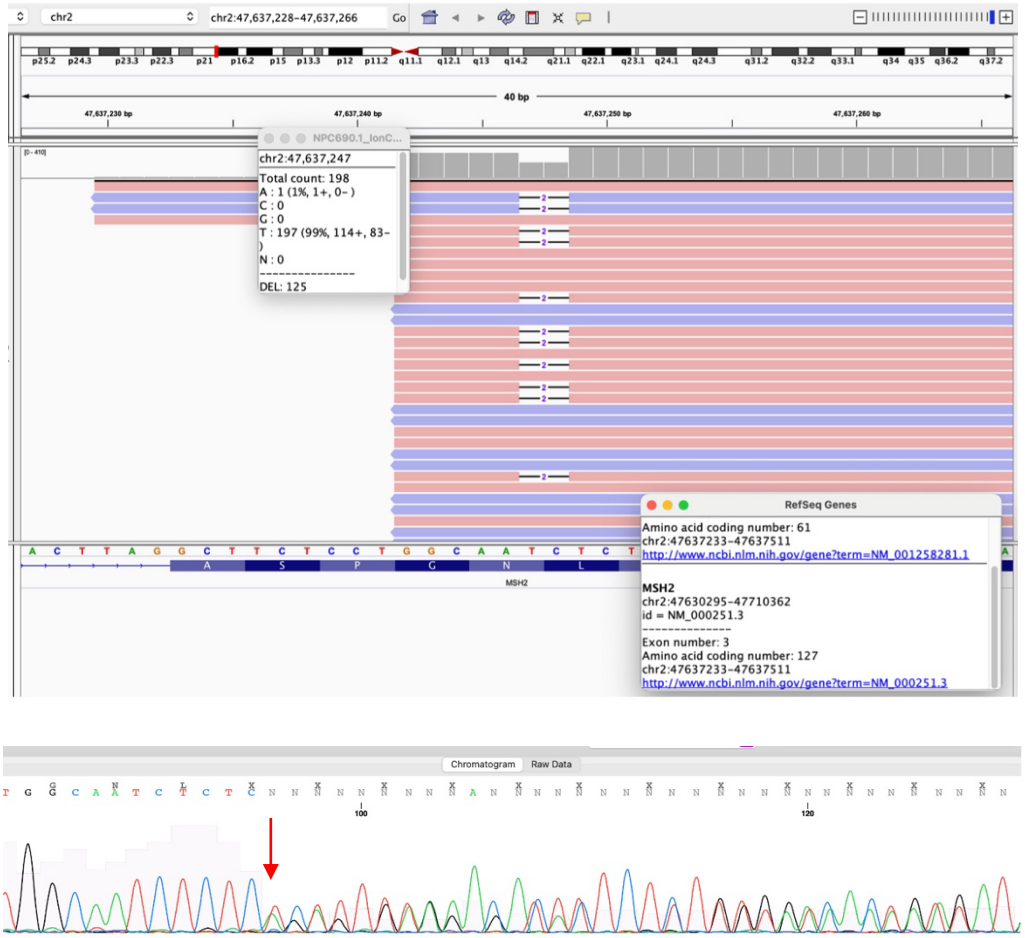

Supplementary Figure 1. Evidence for the two base-pair deletion in *MSH2* in patient 17, confirmed after manual inspection using the Integrative Genomics Viewer software (top). The presence of the variant *MSH2*:c.387_388delTC was then verified in patient 17 by Sanger sequencing (bottom).