SUPPLUMENTAL MATERIAL

Simultaneous Detection of Common Founder Mutations using a Cost-Effective Deep Sequencing Panel

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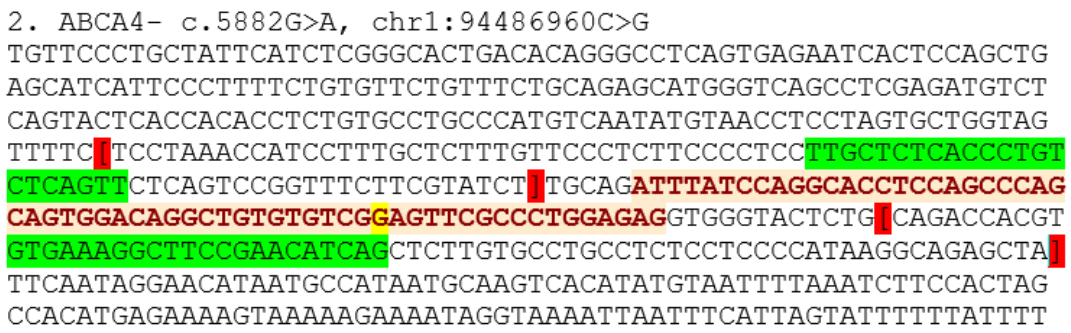
Supp Figure S1



An example of a sequence that was used as an input in FastPCR. In this figure one of the founder mutations from the panel is presented (marked in yellow). The brackets (marked in red) are used to define the preferred primer location. The software designs the primers in the areas bounded by the brackets. The selected primers are marked in green.