

**Supplementary Table 3 – Primer pairs used for *KRAS* mutation analysis and correspondent amplicon lengths.**

<b><i>KRAS</i> exon 3</b>	
Forward primer sequence	5' CCA GAC TGT GTT TCT CCC TTC TCA GG 3'
Reverse primer sequence	5' AGA AAG CCC TCC CCA GTC CTC A 3'
Amplicon length	150 bp
<b><i>KRAS</i> exon 4</b>	
Forward primer sequence	5' ACAGGCTCAGGACTTAGCAAGAAGT 3'
Reverse primer sequence	5' AGCATAATTGAGAGAAAACTGA 3'
Amplicon length	158 bp