

Supplementary Table S1. Frequency of mutations in *KRAS* exon 2 by Sanger sequencing

Base change	Amino acid change	No. of cases
<i>KRAS</i> codon 12		
c.34G>A	G12S	8
c.34G>C	G12R	1
c.34G>T	G12C	3
c.35G>A	G12D	55
c.35G>C	G12A	2
c.35G>T	G12C	2
c.35G>T	G12V	19
Total		90
<i>KRAS</i> codon 13		
c.37G>T	G13C	1
c.38G>A	G13D	23
Total		24