

SUPPLEMENTARY TABLE S3. OVERALL POINT
MUTATIONS OF THE *RAS* GENES

	<i>Amino acid*</i>	<i>Nucleotide</i>	<i>No. of mutations</i>
<i>NRAS</i> codon 61	Q61R	CAA>CGA	16
	Q61K	CAA>AAA	5
<i>HRAS</i> codon 61	Q61R	CAG>CGG	6
	Q61K	CAG>AAG	1
<i>KRAS</i> codon 12/13	G12R	GGT>CGT	5
	G12V	GGT>GTT	1
<i>KRAS</i> codon 61	Q61R	CAA>CGA	4
<i>NRAS</i> codon 12/13	G12W	GGT>TGG	1
<i>HRAS</i> codon 12/13	G12	GGC	0

*Amino acids: Q, glutamine; R, arginine; K, lysine; G, glycine; V, valine; W, tryptophan.