

	N° of cases analysed	WT	p.G12R	p.G12C	p.G12S	p.G12V	p.G12A	p.G12D	p.G13D	p.G13C	Codon 12 mutated	Codon 13 mutated	Mutated total
Sanger sequencing	260	152 (58.5%)	1 (0.4%)	11 (4.2%)	6 (2.3%)	28 (10.8%)	4 (1.5%)	33 (12.7%)	24 (9.2%)	1 (0.4%)	83 (31.9%)	25 (9.6%)	108 (41.5%)
Array	235	131 (55.7%)	0 (0%)	12 (5.1%)	7 (3.0%)	30 (12.8%)	3 (1.3%)	32 (13.6%)	19 (8.1%)	1 (0.4%)	84 (35.7%)	20 (8.6%)	104 (44.3%)
Melting curve	190	113 (59.5%)	-	-	-	-	-	-	-	-	-	-	77 (40.5%)
Pyrosequencing	138	87 (63%)	0 (0%)	5 (3.6%)	4 (2.9%)	14 (10.2%)	2 (1.4%)	18 (13.1%)	7 (5.1%)	1 (0.7%)	43 (31.2%)	8 (5.8%)	51 (37%)