

TECHNICAL SHEET IDYLLA™ NRAS-BRAF-EGFR S492R MUTATION ASSAY

The Idylla™ NRAS-BRAF-EGFR S492R Mutation Assay, performed on the Biocartis Idylla™ System, is a molecular assay for the qualitative detection of 18 mutations in codons 12, 13, 59, 61, 117, 146 of the NRAS gene, 5 mutations in codon 600 of the BRAF gene and 2 mutations in codon 492 of the EGFR gene.

The Idylla™ NRAS-BRAF-EGFR S492R Mutation Assay, from sample-to-result, starts with formalin-fixed paraffin-embedded (FFPE) tissue to liberate DNA for subsequent real-time PCR amplification and detection.

FEATURES

G12C (c.34G>T) G12S (c.34G>A) Codon 12 (exon 2) G12D (c.35G>A) G12A (c.35G>C) G12V (c.35G>T) G13D (c.38G>A) Codon 13 (exon 2) G13V (c.38G>T) G13R (c.37G>C) Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A) Q61L (c.182A>T)
Codon 12 (exon 2) G12D (c.35G>A) G12A (c.35G>C) G12V (c.35G>T) G13D (c.38G>A) Codon 13 (exon 2) G13V (c.38G>T) G13R (c.37G>C) Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A)
G12A (c.35G>C) G12V (c.35G>T) G13D (c.38G>A) Codon 13 (exon 2) G13V (c.38G>T) G13R (c.37G>C) Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A)
G12V (c.35G>T) G13D (c.38G>A) Codon 13 (exon 2) G13V (c.38G>T) G13R (c.37G>C) Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A)
G13D (c.38G>A) Codon 13 (exon 2) G13V (c.38G>T) G13R (c.37G>C) Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A)
Codon 13 (exon 2) G13V (c.38G>T) G13R (c.37G>C) Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A)
G13R (c.37G>C) Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A)
Codon 59 (exon 3) A59T (c.175G>A) Q61K (c.181C>A)
Q61K (c.181C>A)
Q61L (c.182A>T)
Codon 61 (exon 3)
Q61R (c.182A>G)
Q61H (c.183A>C; c.183A>T)
Codon 117 (exon 4) K117N (c.351G>C; c.351G>T)
Codon 146 (exon 4) A146T (c.436G>A) (c.437C>T)

NRAS Total (acting as Sample Processing Control)



BRAF mutation detection		
Codon 600	BRAF V600E BRAF V600D	(c.1799T>A; c.1799_1800delinsAA) (c.1799_1800delinsAC)
	BRAF V600K BRAF V600R	(c.1798_1799delinsAA) (c.1798_1799delinsAG)

BRAF Total (acting as Sample Processing Control)

EGFR mutation detection			
Codon 492	S492R S492R2	(c.1476C>A) (c.1474A>C)	
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EGFR Total (acting as Sample Processing Control)

Specimen requirements	
Sample Type	FFPE tissue sections (5 to 10 μm)
Neoplastic cells	≥10%, if less macrodissection is required
Tissue area	50-600 mm² (5 μm) 25-300 mm² (10 μm)
Total turnaround time	
Time	120 minutes
Result Reporting	
Report	Qualitative genotype call



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