

TECHNICAL SHEET IDYLLA™ KRAS MUTATION TEST

The **Idylla™ KRAS Mutation Test**, performed on the Biocartis Idylla[™] system, is an *in vitro* diagnostic Test for the qualitative detection of 21 mutations in **codons 12, 13, 59, 61, 117 and 146** of the *KRAS* gene. The Idylla[™] KRAS Mutation Test, from **sample-to-result**, starts with formalin-fixed paraffin-embedded (FFPE) human tissue from metastatic colorectal cancers to liberate DNA for subsequent real-time PCR amplification and detection.

FEATURES

KRAS mutation detection		
	G12C	(c.34G>T)
	G12R	(c.34G>C)
Coden 12 (even 2)	G12S	(c.34G>A)
	G12A	(c.35G>C)
	G12D	(c.35G>A)
	G12V	(c.35G>T)
Codon 13 (exon 2)	G13D	(c.38G>A)
	A59E	(c.176C>A)
Codon 59 (exon 3)	A59G	(c.176C>G)
	A59T	(c.175G>A)
	Q61K	(c.181C>A; c.180_181delinsAA)
Codon 61 (exon 3)	Q61L	(c.182A>T)
	Q61R	(c.182A>G)
	Q61H	(c.183A>C; c.183A>T)
Codon 117 (exon 4)	K117N	(c.351A>C; c.351A>T)
	A146P	(c.436G>C)
Codon 146 (exon 4)	A146T	(c.436G>A)
•••••••••••••••••••••••••••••••••••••••	A146V	(c.437C>T)

KRAS Total (acting as Sample Processing Control)

Specimen requirements	
Sample Type	FFPE tissue sections (5 to 10 $\mu\text{m})$
Neoplastic cells	≥10%, if less macrodissection is required
Tissue area	50-600 mm² (5 μm) 25-300 mm² (10 μm)
Performance	

Analytical Sensitivity

LOD ≤5% for all KRAS mutations



	100% agreement for 5% KRAS G12D
Between Laboratory Reproducibility (480 results at 3 sites)	100% agreement for 5% KRAS G12S
	100% agreement for 5% KRAS G12V
	100% agreement for 50% KRAS G13D
	100% agreement for 5% KRAS G12A
Detuces Let Desve ducibility	100% agreement for 5% KRAS G12D
(375 results on 3 lots)	100% agreement for 5% KRAS G12S
	100% agreement for 5% KRAS G12V
	100% agreement for 5% KRAS G13D

Total turnaround time

Time

120 minutes

ACCURACY - CLINICAL PERFORMANCE EVALUATION

96.7% overall percent agreement was obtained during the clinical performance evaluation comparing Idylla[™] with a reference method based on RT-PCR.

96.7% overall concordance							Referen	ce test		
		G12A	G12C	G12D	G12R	G12S	G12V	G13D	No mutation	Total
ldylla™	G12A	6								6
KRAS Mutation	G12C		6							6
Test	G12D			25						25
	G12R				З				1*	4
	G12S					6				6
	G12V					1	15		1*	17
	G13D					1		16	З*	20
	No mutation	1*							97	98
	A59E/G/T								1	1
	Q61H/H2								3	3
	Q61K/K2									0
	Q61L/R									0
	K117N/N2		1						2	3
	A146T/V/P						1		4	5
	Totals	7	7	25	3	8	16	16	112	194

Note: the Reference test is not designed to pick up mutations in codon 59, 61, 117 and 146

* Due to limitations in available material (n=2) or insufficient DNA quality (n=3), only one of the 6 discordant results could be resolved; NGS confirmed the G12V result called by Idylla[™]

Discordant analysis by NGS

100% ove		R	eference te	est and fu	rther analy	sis by NGS				
		G12A	G12C	G12D	G12R	G12S	G12V	G13D	No mutation	Total
Idylla™ KRAS Mutation	G12A	6	••••••	• ••••••	•••••••	•••••••••••••••	••••••••••	•••••		6
	G12C		6							6
Test	G12D			25						25
	G12R				З					3
	G12S					6				6
	G12V					1	16			17
	G13D					1		16		17
	No mutation								97	97
	A59E/G/T								1	1
	Q61H/H2								3	3
	Q61K/K2									0
	Q61L/R									0
	K117N/N2								2	3
	A146T/V/P		1				1		4	5
_	Totals	6	7	25	3	8	17	16	107	189

MULTI-CENTER EVALUATION OF THE FULLY-AUTOMATED PCR-BASED IDYLLA™ KRAS MUTATION ASSAY FOR RAPID KRAS MUTATION STATUS DETERMINATION ON FORMALIN-FIXED PARAFFIN-EMBEDDED TISSUE OF HUMAN COLORECTAL CANCER.

Solassol J. et al. PLOS ONE 2016.

95.9% overall concordance				Routine reference methods*						
		Codon 12	Codon 13	Codon 59	Codon 61	Codon 117	Codon 146	No mutation	Total	
ldylla™	Codon 12	138					•	З	141	
KRAS Mutation	Codon 13		38						38	
Assay	Codon 59			5					5	
	Codon 61	1			14			5	20	
	Codon 117					4			4	
	Codon 146						18	1	19	
	No mutation	З	2				1	129	135	
	TOTAL	142	40	5	14	4	19	138	362	

Mutations not picked up by Idylla[™] or the reference method (n=8) are not included in the % agreement calculation. Idylla does not pick up the following low prevalent (<1%) mutations: 4xG13C, 1xG13R, 1xG12F (samples excluded from table). Roche Cobas does not pick up mutations in codon 146 (2x).

* Different reference methods were used: cobas[®] KRAS Mutation Test (Roche), Ion Torrent AmpliSeq[™] Colon and Lung Cancer Research Panel (Life Technologies), therascreen[®] KRAS Pyro[®] Kit (Qiagen), therascreen[®] RAS Extension Pyro Kit (Qiagen), HRM screening and pyrosequencing, Sanger sequencing, HRM screening and Sanger sequencing; for the analysis, when Idylla[™] identified a specific mutation in codon 12, 13 or 61, and the cobas[®] KRAS Mutation Test (Roche) reported a "codon 12/13" or "codon 61" result), both results were considered identical.

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Discordant analysis by ddPCR, Idylla™ retest and NGS

98.9% ov	erall concorda	nce	Routine reference methods, including further analysis by ddPCR, Idylla™ retest, NGS							
		Codon 12	Codon 13	Codon 59	Codon 61	Codon 117	Codon 146	No mutation	Total	
Idylla™ KRAS Mutation Assay	Codon 12	142							142	
	Codon 13		38						38	
	Codon 59			5					5	
	Codon 61	1			15			1	17	
	Codon 117					4			4	
	Codon 146						20		20	
	No mutation	1	2					133	136	
	TOTAL	144	40	5	15	4	20	134	362	

IDYLLA™ KRAS POSTERS & PUBLICATIONS

- Maertens G. et al. A solution for same-day extended RAS testing. Poster ESMO 2015
- Vandenbroucke I. et al. A rapid and fully automated multiplex assay for KRAS-BRAF mutations with high mutation sensitivity using novel selective amplification and detection technologies. Poster AACR 2014
- Solassol J. et al. Multi-Center Evaluation of the Fully Automated PCR-Based Idylla™ KRAS Mutation Assay for Rapid KRAS Mutation Status Determination on Formalin-Fixed Paraffin-Embedded Tissue of Human Colorectal Cancer. PLOS ONE 2016
- Weyn C. et al. Clinical performance evaluation of a sensitive, rapid low-throughput test for KRAS mutation analysis using formalin-fixed, paraffin-embedded tissue samples. BMC Cancer 2017

RESEARCH APPLICATIONS

- Dario de Biase. et al. Fully automated PCR detection of KRAS mutations on pancreatic endoscopic ultrasound fine-needle aspirates. J Clin Path 2016.



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