

Evaluation of a Fully Automated System for Use in Somatic Mutation Testing in Colorectal Cancer: a Prospective Study with Comparison to Next-Generation Sequencing

M. Rabie Al-Turkmani, Donald C. Green, Alicia A. Finney, Michael A. Suriawinata, Gregory J. Tsongalis

Department of Pathology and Laboratory Medicine, Dartmouth-Hitchcock Medical Center and Geisel School of Medicine at Dartmouth, Lebanon, NH

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BACKGROUND

- Colorectal cancer (CRC) is the third most commonly diagnosed cancer in the United States among both men and women.
- The management of CRC has been aided by the advancement and increased usage of molecular testing.
- Mutational analysis of *KRAS* and *NRAS* is warranted for CRC patients considered for anti-EGFR therapy. *BRAF* codon 600 testing is also recommended in CRC for prognostic stratification.
- The American Society of Clinical Oncology (ASCO) recently expanded their recommendations on molecular testing for CRC patients considered for anti-epidermal growth factor receptor (anti-EGFR) therapy to include *KRAS* and *NRAS* codons 12 and 13 of exon 2, codons 59 and 61 of exon 3, and codons 117 and 146 of exon 4 (so-called "expanded" or "extended" RAS testing).
- The Idylla™ system (Biocartis, Mechelen, Belgium) is a fully automated platform that utilizes a single-use cartridge containing all necessary reagents and steps for sample processing and real-time PCR-based mutation amplification and detection.
- In this prospective study, we evaluated the Idylla™ system against next generation sequencing using colorectal cancer tissue specimens.

DESIGN & METHODS

- Seventy-five CRC tissue specimens which were ordered for next-generation sequencing (NGS) between January 2018 and December 2018 were also prospectively tested on the Idylla™ system using the *KRAS* and *NRAS-BRAF-EGFR S492R* cartridges (Research Use Only), which test for actionable mutations in *KRAS*, *NRAS* and *BRAF*.
- Two 10 µm formalin-fixed paraffin-embedded (FFPE) tissue sections (one for each cartridge) were used for each run on the Idylla™ and all cases met the system's minimum tumor requirement of 10%.
- NGS testing was performed using the Ion AmpliSeq 50-gene Cancer Hotspot Panel v2 (Thermo Fisher Scientific). NGS results and turnaround time of the testing process were compared to those obtained by the Idylla™ system.



Figure 1. The Idylla™ system

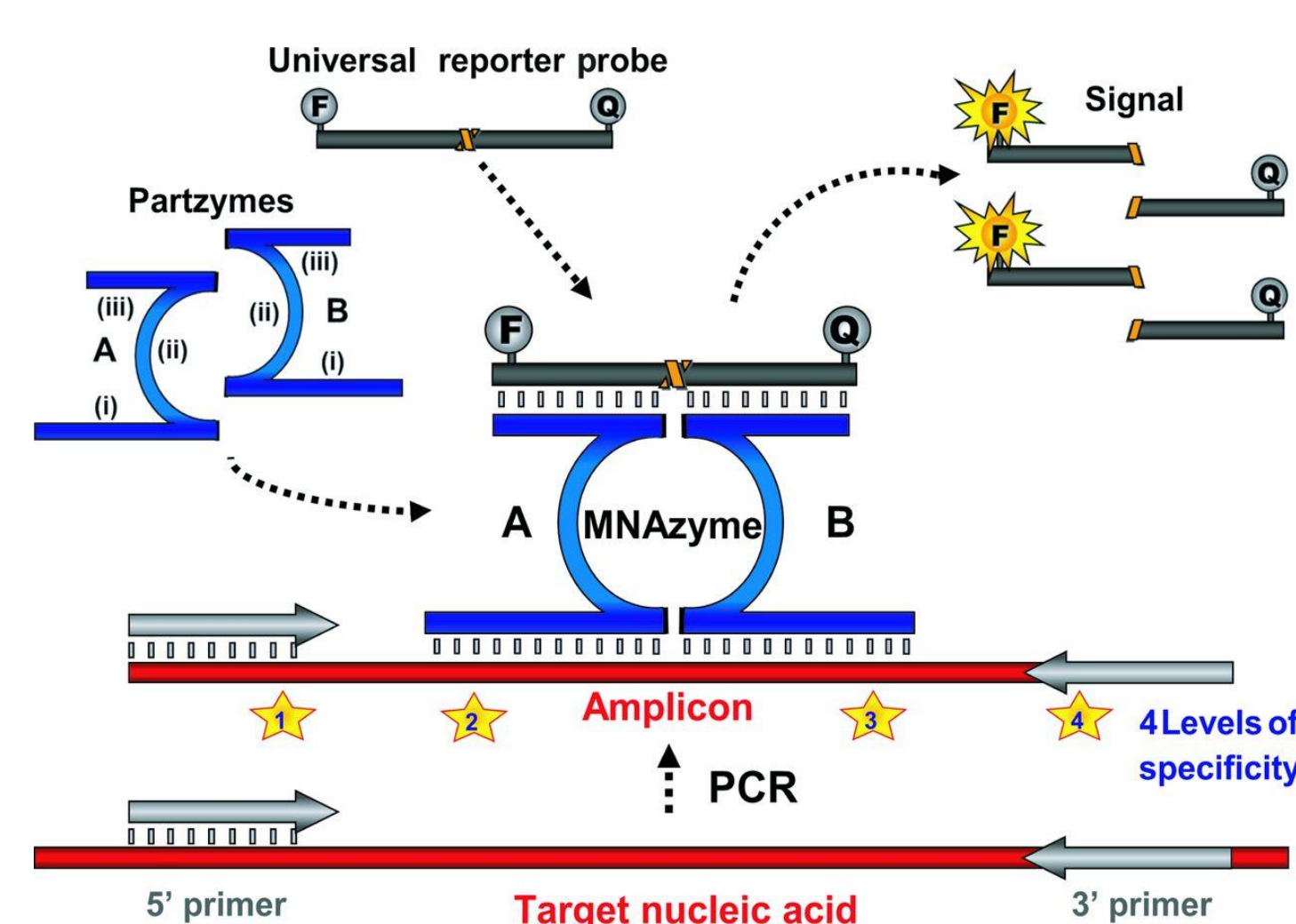


Figure 2. PlexPrime®/PlexZyme® Technology

KRAS Mutation Assay (RUO)		
Exon	Codon	Mutation
2	12	G12C, G12R, G12S, G12A, G12D, G12V
	13	G13D
3	59	A59T, A59E, A59G
	61	Q61K, Q61L, Q61R, Q61H
4	117	K117N
	146	A146P, A146T, A146V

NRAS-BRAF-EGFR S492R Mutation Assay (RUO)			
Gene	Exon	Codon	Mutation
NRAS	2	12	G12C, G12S, G12A, G12D, G12V
		13	G13D, G13R, G13V
	3	59	A59T
		61	Q61K, Q61L, Q61R, Q61H
	4	117	K117N
		146	A146T, A146V
BRAF	15	600	V600E, V600D, V600K, V600R
EGFR	12	492	S492R

Tables 1 & 2. Mutations detectable by the Idylla™ *KRAS* and *NRAS-BRAF-EGFR S492R* Mutation Assays

RESULTS

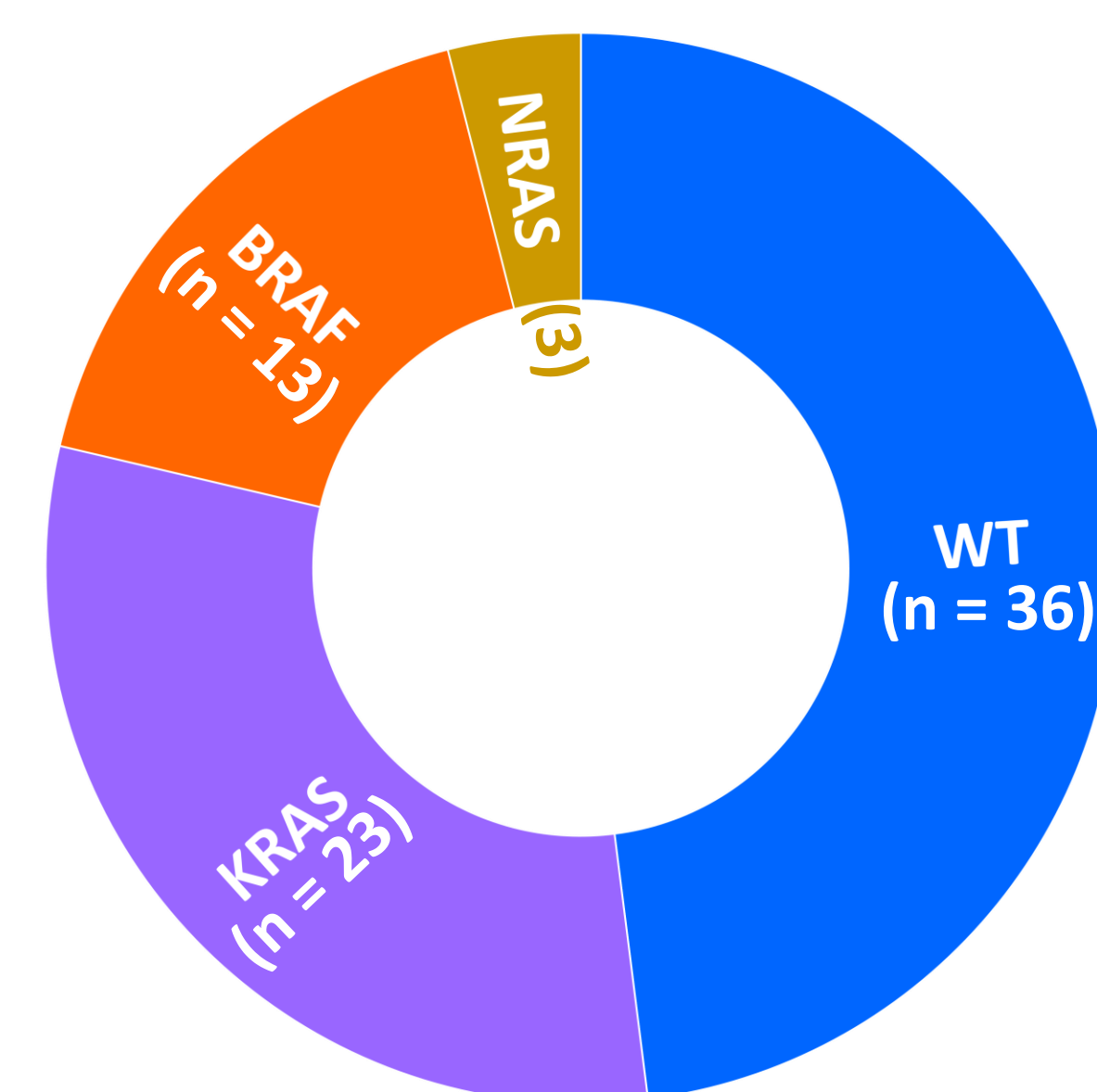


Figure 3. Distribution of *KRAS*, *BRAF*, and *NRAS* mutations among study samples

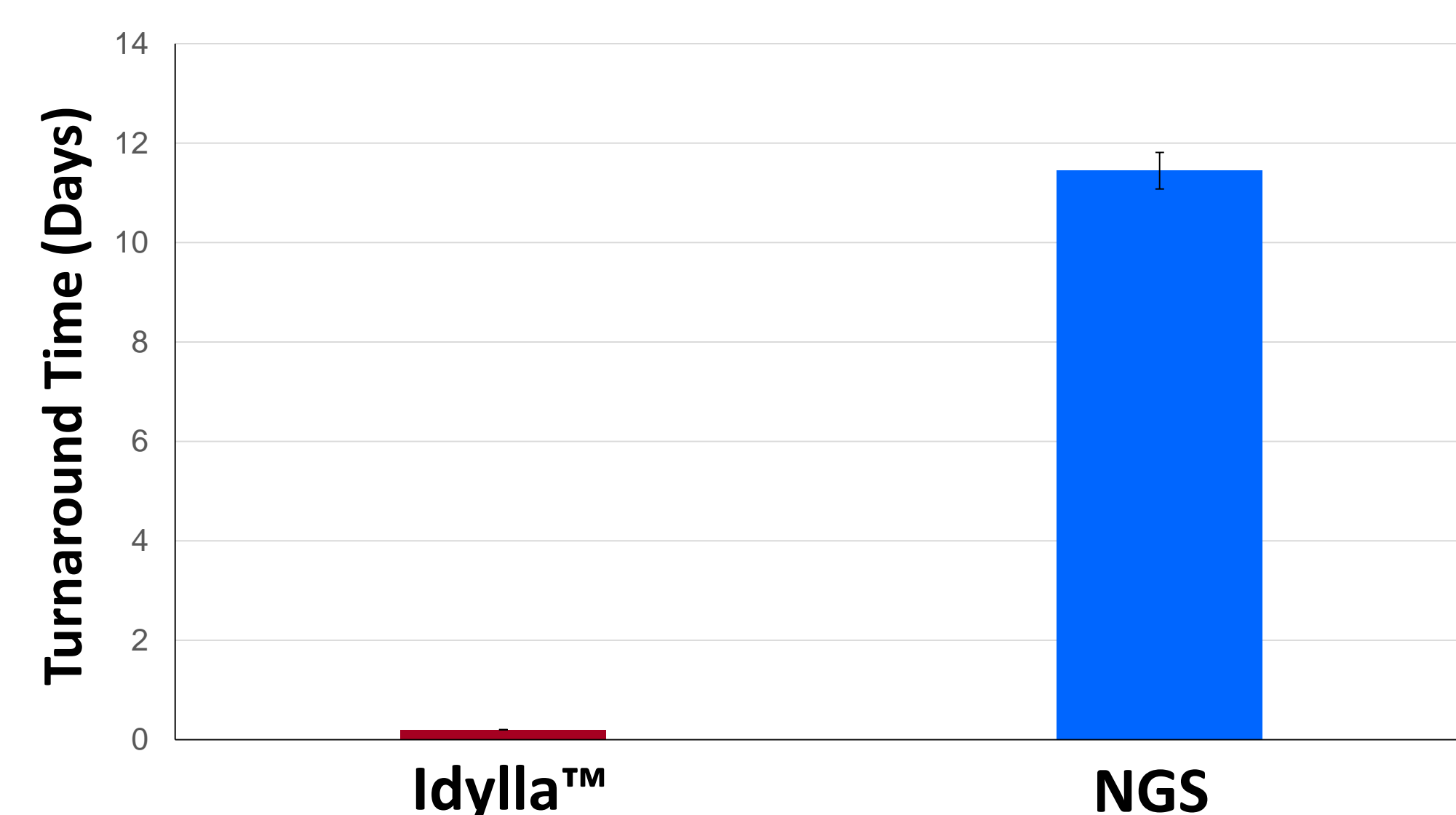


Figure 4. Comparison of Idylla™ and NGS average testing turnaround time

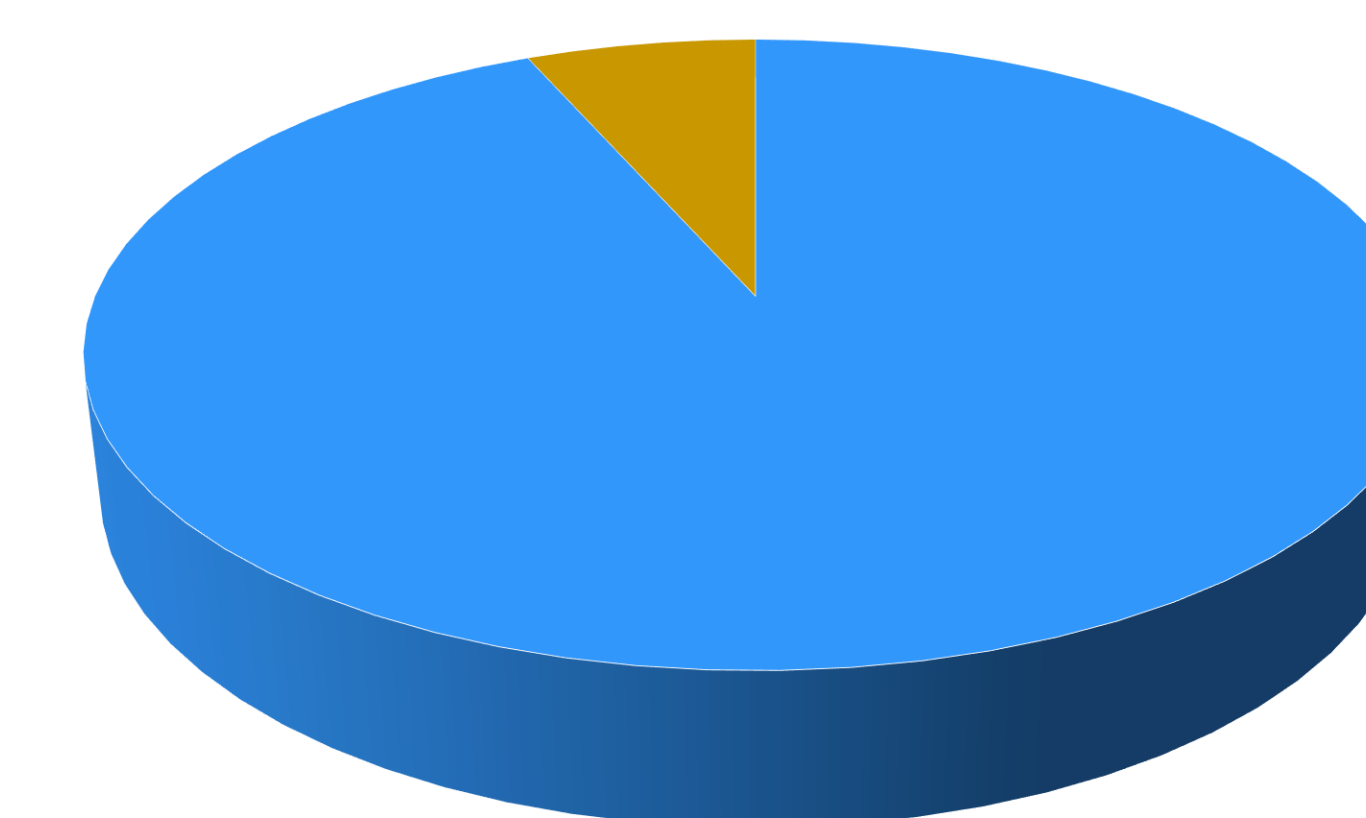
Gene Mutated	Number of samples	Idylla™	NGS
BRAF	10	<i>BRAF</i> V600E/D	<i>BRAF</i> V600E
	1	<i>BRAF</i> V600E/D	<i>BRAF</i> V600E, <i>KRAS</i> T20M*
	1	No mutation detected	<i>BRAF</i> D594N*
	1	No mutation detected	<i>BRAF</i> K601E*
KRAS	3	<i>KRAS</i> G12A	<i>KRAS</i> G12A
	1	<i>KRAS</i> G12C	<i>KRAS</i> G12C
	7	<i>KRAS</i> G12D	<i>KRAS</i> G12D
	1	<i>KRAS</i> G12D, <i>NRAS</i> G12A/V [†]	<i>KRAS</i> G12D
	1	<i>KRAS</i> G12S	<i>KRAS</i> G12S
	1	<i>KRAS</i> G12S, <i>NRAS</i> G12A/V [†]	<i>KRAS</i> G12S
	4	<i>KRAS</i> G12V	<i>KRAS</i> G12V
	3	<i>KRAS</i> G13D	<i>KRAS</i> G13D
	1	<i>KRAS</i> Q61K	<i>KRAS</i> Q61K
	1	<i>KRAS</i> A146P/T/V	<i>KRAS</i> A146T
NRAS	1	<i>NRAS</i> G12D	<i>NRAS</i> G12D
	1	<i>NRAS</i> Q61K	<i>NRAS</i> Q61K
	1	<i>NRAS</i> Q61R	<i>NRAS</i> Q61R
None (WT)	36	No mutation detected in <i>BRAF</i> , <i>KRAS</i> , or <i>NRAS</i>	No mutation detected in <i>BRAF</i> , <i>KRAS</i> , or <i>NRAS</i>

Table 3. Idylla™ results compared with NGS

*Mutations not included in the list of mutations detectable by the Idylla™

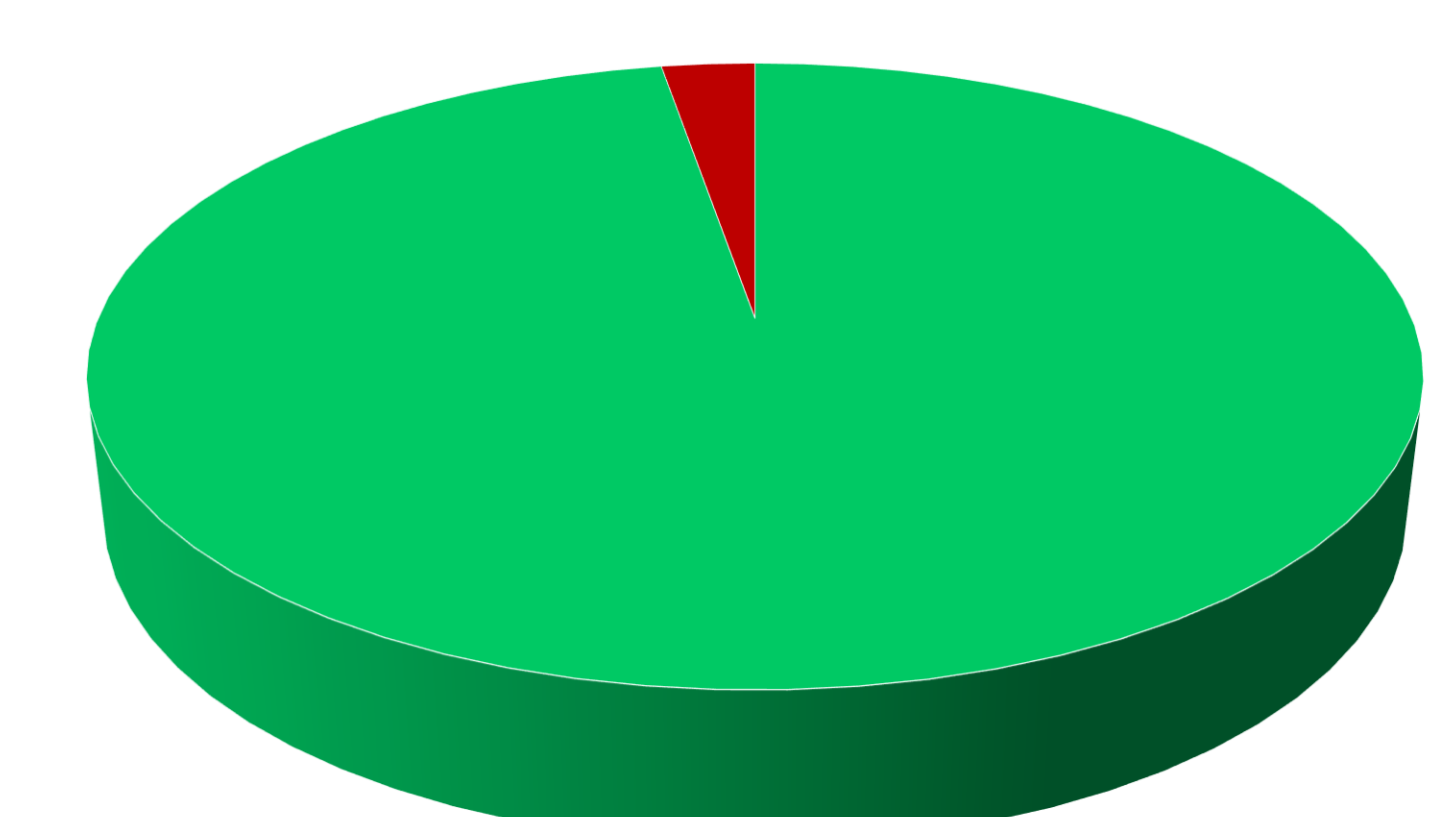
[†]Additional *NRAS* mutation detected by the Idylla™ but not by NGS

A. Variant Concordance



■ Concordant ■ Discordant

B. Clinical Concordance



■ Concordant ■ Discordant

Figure 5. Variant (A: 93%, 70/75) and clinical (B: 97%, 73/75) concordance between Idylla™ and NGS results

CONCLUSIONS

- The Idylla™ system offers rapid and reliable testing of clinically actionable mutations in colorectal cancer specimens directly from FFPE tissue sections.
- This prospective study demonstrates high concordance between the Idylla™ system and NGS.
- The Idylla™ system is ideal for centers that lack highly trained molecular staff or infrastructure and can complement NGS in larger centers by providing rapid turnaround time.