## THE UNIVERSITY OF TEXAS MDAnderson Cancer Center

# Circulating Tumor DNA Assay Performance for Detection and Monitoring of **KRAS Mutations in Urine from Patients with Advanced Cancers**

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## Introduction

- Noninvasive urinary ctDNA-based liquid biopsy approach can be used to detect and track cancer driver mutations for rapid diagnosis and disease monitoring.
- Using a highly sensitive ctDNA mutation detection platform, we examined detection of KRAS G12/13 mutations in urine obtained from patients with advanced cancers, assessed urine sample requirements, and compared the results with matched tumor tissue.

Plasma

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*KRAS*-Positive Advanced Cancers

Urine

Experimental Therapies

Monitor Urine and Plasma KRAS ctDNA for Correlation with Response to Treatment

Urine and plasma samples collected from 41 patients with advanced or metastatic cancers positive for KRAS G12/13 mutations in tissue (colorectal, n=26; lung, n=6; pancreatic, n=4; ovarian, n=2; melanoma, n=1; breast, n=1; other, n=1).

### Study Design

TABLE 1: Patient demographics						
	TOTAL (N=41)	# of patients (%)				
Age, years – Median (range)	56 (38-77)					
Gender	Male	20 (48.8)				
	Female	21 (51.2)				
ECOG Performance status	0	4 (9.8)				
	1	35 (85.4)				
	2	1 (2.4)				
	3	1 (2.4)				
Cancer type	Colorectal cancer	29 (70.7)				
	Non-small cell lung cancer	6 (14.6)				
	Pancreatic cancer	2 (4.9)				
	Ovarian cancer	2 (4.9)				
	Others	2 (4.9)				
KRAS mutation	G12C	7 (17.1)				
	G12D	21 (51.2)				
	G12R	3 (7.3)				
	G12S	2 (4.9)				
	G12V	6 (14.6)				
	G13D	2 (4.9)				

### **KRAS G12/13 Mutation Enrichment NGS Assay**

- Mutant allele enrichment PCR, followed by NGS, utilizes a 31-bp footprint and selectively amplifies mutant DNA fragments while suppressing Wild-Type (WT) sequence amplification.
- Proprietary analysis algorithm allows accurate quantitation of mutant DNA by interpolation to standard curves.
- Mutation enrichment results in approximately 3000-fold increase in ratio of mutant over WT signal for low copy number inputs. For 5 mutant KRAS G12D copies spiked into 18,181 copies of WT DNA (0.0275%), the output sequencing library contains 94% mutant reads (~3418 fold enrichment; **Table 2**).

#### **TABLE 2: Comparison between input ratio of** mutant/WT KRAS copies and output ratio of mutant/WT KRAS sequencing reads

Input MT Copies/WT Copies	Reads/Wild	Mutant Sequ Type Reads <i>KRAS</i> G12V	(% Mutant)
(% Mutant)		_	
5/18,181	13447/858	4269/410	2318/748
(0.0275%)	(94%)	(91.2%)	(75.6%)
15/18,181	34363/1155	9068/423	15726/1053
(0.0825%)	(96.7%)	(95.5%)	(93.7%)
125/18,181	156863/1855	144666/1821	170503/1348
(0.688%)	(98.8%)	(98.8%)	(99.2%)
250/18,181	309123/2307	267933/2452	331498/2216
(1.375%)	(99.3%)	(99.1%)	(99.3%)
500/18,181	508045/1442	472491/2836	585254/1807
(2.750%)	(99.7%)	(99.4%)	(99.7%)

Similar results obtained for other variants (data not shown).

### **Analytical Assay Sensitivity** Lower Limit of Detection (LLoD) = 0.002%

- LLoD was defined as the lowest number of copies for which frequency distribution of the copy number events, upon repeated measurements, falls within the 95% confidence interval of expected frequency distribution determined by Poisson statistics.
- KRAS G12/13 assay LLoD:
- 1 mutant copy in a background of 18,181 copies of Wild-Type DNA (0.006%; **Table 3**).
- 2 mutant copies in a background of 100,000 copies of Wild-Type DNA (0.002%; data not shown).



The observed frequency distribution of copy number events falls within the expected distribution for the LLoD of 1 mutant copy in a background of 18,181 copies of WT DNA.

### Clinical Sensitivity of KRAS G12/13 ctDNA Assay

Of 41 patients with advanced cancers, 23 had pre-treatment baseline urine samples with urine volumes 40-110 mL available and 29 had pre-treatment plasma (1-4 mL) samples available.



- Urinary DNA yields ranged from 15 to 23059 ng (median, 1059 ng).
- Plasma DNA yields ranged from 12 to 1846 ng (median, 55 ng).
- (**Table 4**):

### **Colorectal Cancer**

- 100% (4/4) for urine with recommended volume of at least 90 mL.
- 78% (14/18) for urine with all volumes (40-110 mL). Multiple cancers
- 80% (4/5) for urine with recommended volume of at least 90 mL.
- 70% (16/23) for urine with all volumes (40-110 mL).
- **Colorectal Cancer** 
  - 100% (16/16) Multiple cancers

### - 83% (24/29)

### Results

 
 TABLE 3: Verification of KRAS G12/13 assay
LLoD

lumber of Mutant Copies	0 (Not-detected)	1+ (Detected)
Expected (95% CI) [1 copy/rep]	7 (2-14)	13 (6-20)
Observed G12A	2	8
Observed G12C	5	15
Observed G12D	3	17
Observed G12R	10	10
Observed G12S	6	14
Observed G12V	4	16
Observed G13D	3	17

#### TABLE 4: Urine/tissue concordance for KRAS G12/13 detection

Colorectal Cancer	Urine (90-110 mL)	100% (4/4)
	Urine (40-110 mL)	78% (14/18)
	Plasma (1-4 mL)	100% (16/16)
Multiple Cancers	Urine (90-110 mL)	80% (4/5)
	Urine (40-110 mL)	70% (16/23)
	Plasma (1-4 mL)	83% (24/29)

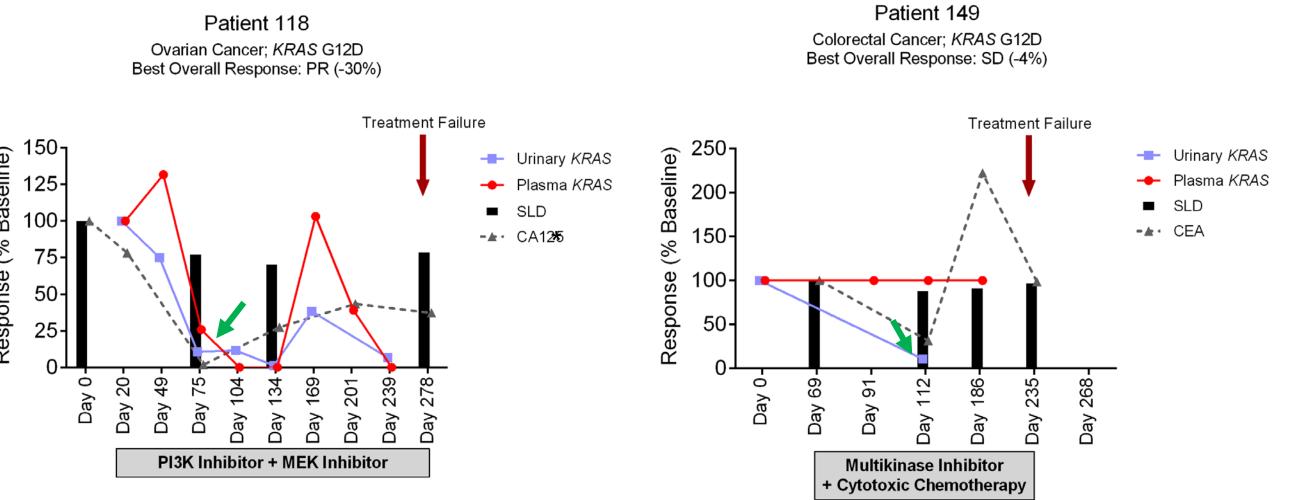
Positive percent agreement (PPA) for urinary KRAS G12/13 with tumor tissue as reference

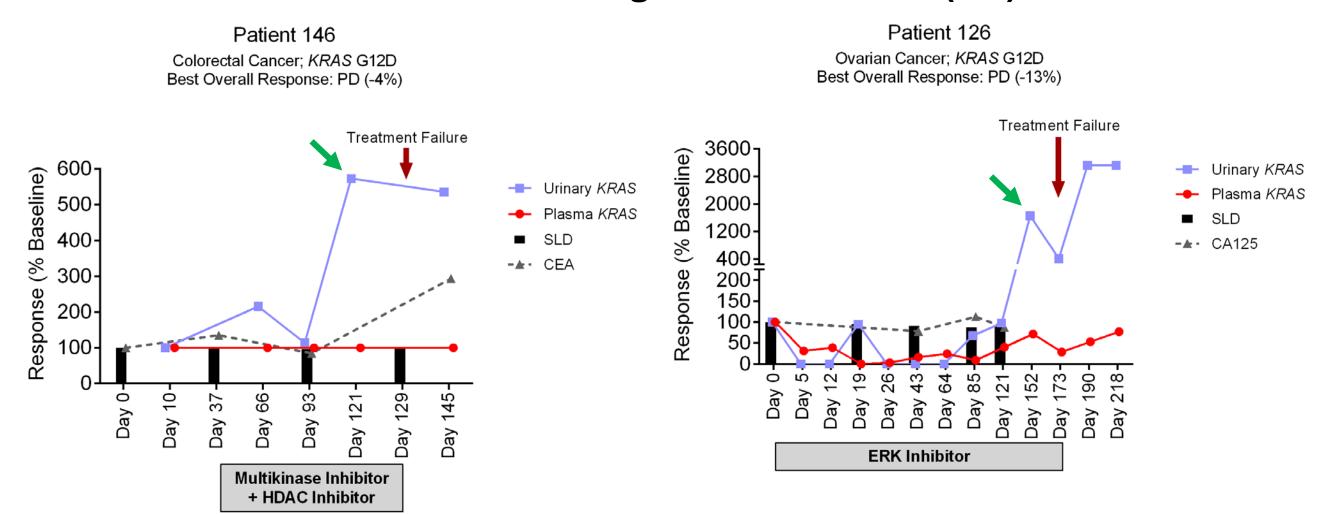
Positive percent agreement (PPA) for plasma KRAS G12/13 with tumor tissue as reference:

### Longitudinal Monitoring by ctDNA KRAS G12/13

Representative cases showing dynamics of urine and plasma ctDNA KRAS G12/13 in patients with advanced or metastatic cancers on experimental therapies.





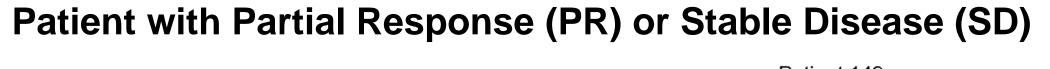


response ().

\*SLD, sum of the longest diameters of index lesions.

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Early decreases from baseline (by 89-96%) are observed for urine KRAS G12/13 signal in patients with PR or SD as best overall response using RECIST 1.1 criteria ( $\downarrow$ ).

### Patients with Progressive Disease (PD)

Increases in urine KRAS G12/13 precede clinical progression in patients with PD as best overall

### Conclusions

 Mutation enrichment NGS assay for KRAS G12/13 mutation detection in ctDNA has a single copy analytical sensitivity (0.002%-0.006%).

 In a blinded study of 41 patients with advanced or metastatic cancers, including colorectal, pancreatic, ovarian, lung, melanoma, and breast cancers, KRAS G12/13 mutation concordant with tissue was detected in 80% of urine samples with recommended volume of at least 90 mL and 83% of plasma samples with volumes 1-4 mL. In a colorectal cancer cohort, KRAS G12/13 mutation detection sensitivity was 100% for both urine and plasma.

Kinetics of KRAS G12/13 mutation signal in urine ctDNA corresponds to treatment outcomes.

 Analysis of urine and plasma may be a viable approach for diagnostic detection of KRAS mutations and therapeutic monitoring of patients with advanced cancers.