# Detection of BRAF Mutations in Urine and Plasma Cell-Free DNA: Application to the Diagnosis and Management of Histiocytic Disorder Patients

Omar Abdel-Wahab<sup>1,2,</sup> Eli L. Diamond<sup>3</sup>, Minal Patel<sup>2</sup>, Veronica R. Holley<sup>4</sup>, Goran Cabrillo<sup>4</sup>, Raaajit Rampal<sup>2</sup>, Latifa Hassaine<sup>5</sup>, Karena Kosco<sup>5</sup>, Jose Baselga<sup>6</sup>, Razelle Kurzrock<sup>7</sup>, Jason C. Poole<sup>5</sup>, Cecile Rose Vibat<sup>5</sup>, Mark Erlander<sup>5</sup>, Filip Janku<sup>4</sup>, David M. Hyman<sup>6</sup>

RESULTS

<sup>1</sup>Human Oncology and Pathogenesis Program, <sup>2</sup>Leukemia Service, <sup>5</sup>Neurology Service, <sup>6</sup>Developmental Therapeutics, MD Anderson Cancer Center; <sup>5</sup>Trovagene Inc.; <sup>7</sup>Moores Cancer Center, UCSD

--9 patients with indeterminate initial tissue biopsy result (34.6%)

Concordance between urinary cfDNA, plasma cfDNA, and tissue biopsy

--11 patients to be *BRAF*V600E mutant (42.3%)

--6 patients as *BRAF*V600E wildtype (23.1%)

- 40-60% of patients with the systemic histiocytoses (including Langerhans Cell Histiocytosis (LCH) and Erdheim-Chester Disease (ECD)) have a *BRAF*V600E mutation.
- Treatment with RAF inhibitors have dramatic effects on the disease.
- However, contamination of stromal cells, low tumor content, and frequent use of bone biopsy lesions for molecular testing often impede effective detection of BRAFV600E mutations in clinical practice for these patients.
- We therefore sought to assess the utility of urinary and plasma cell-free DNA (cfDNA) analysis to (1) reliably detect *BRAF*V600E mutation and (2) dynamically monitor response to therapy in this unique subset of patients.

# Typical Pathology

Cells die releasing their

genomic DNA into the

fragments enter the

kidneys and are

excreted from the

kidneys into the urine

solated and detected

bloodstream

# **METHODS**

**BACKGROUND** 

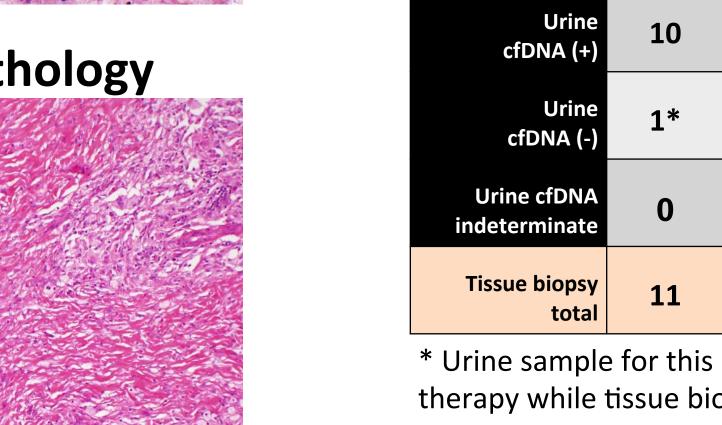
- Between January 2013 and June 2014, 26 consecutive patients with LCH (n=5) and ECD (n=21) were enrolled from Memorial Sloan Kettering Cancer Center and MD Anderson Cancer Center.
- Urinary cfDNA analysis was performed in all patients and plasma cfDNA analysis in 19/26 patients.
- Serial urinary samples in 10 BRAFV600E mutant samples were also obtained to track disease burden with therapy.
- Urine and plasma cfDNA were quantified by a droplet digital PCR (ddPCR; QX-100, BioRad).

### Two-Sten Design for 31 hn RRAF VANNE Assay

Iwo-step besign for 31 bp BRAF volue Assay								
STEP ONE: Pre-amplification with wild-type suppression to decrease amplification of WT patient DNA	Tag A  M  tag B  WT Blocker  WT							
STEP TWO: Amplification with primers complimentary to A,B tags	A M TaqMan Prob	)е 						
(droplet digital PCR)								

## Of 26 patients, initial tissue BRAFV600E genotyping identified:



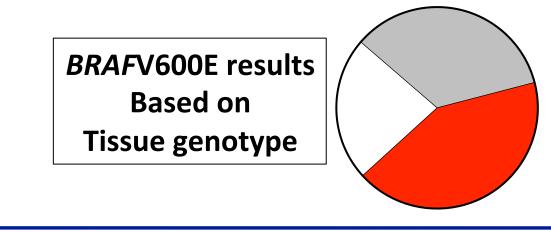


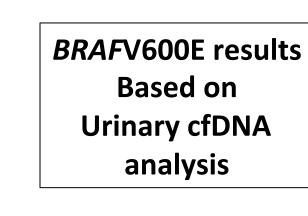
	Tissue biopsy (+)	Tissue biopsy (-)	Tissue biopsy (?)	Urine cfDNA Total	
Urine cfDNA (+)	10	0	4	14	
Urine cfDNA (-)	1*	6	4	11	
Urine cfDNA indeterminate	0	0	1	1	Plas inde
Tissue biopsy					Tis

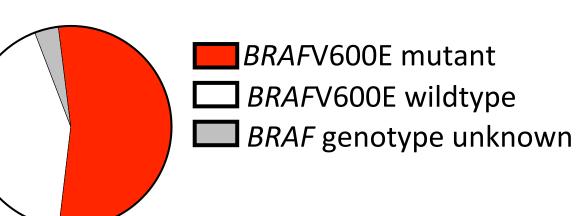
\* Urine sample for this patient was acquired during therapy while tissue biopsy was performed pretreatment

BRAFV600E genotyping results:

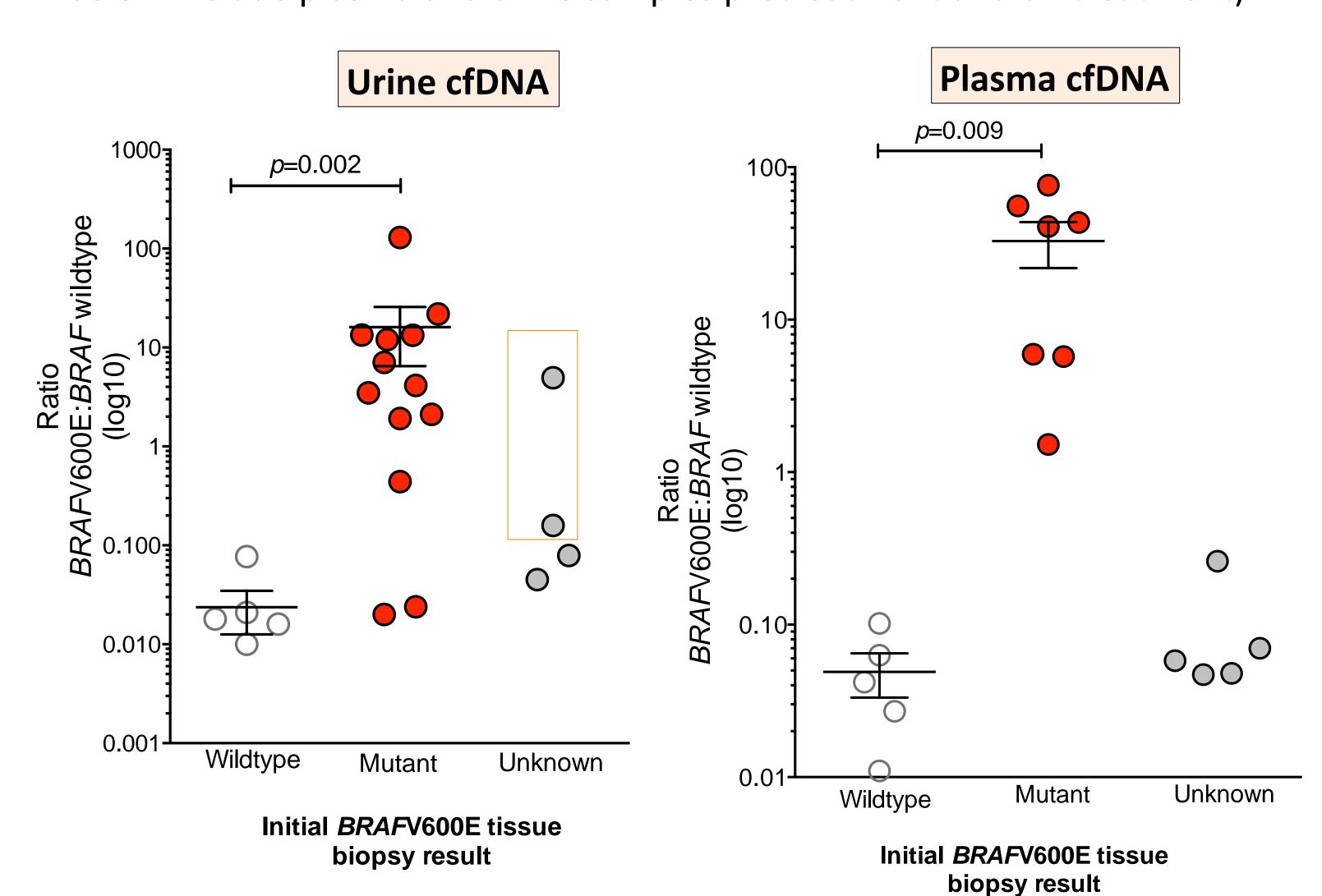
- \*\* These 2 patients subsequently underwent repeat tissue biopsy which confirmed cfDNA test result and allowed the patients to enroll in phase II study of
- cfDNA analysis (both plasma and urine) increased number of patients with known BRAFV600E genotype and identified new additional BRAFV600E mutant patients.







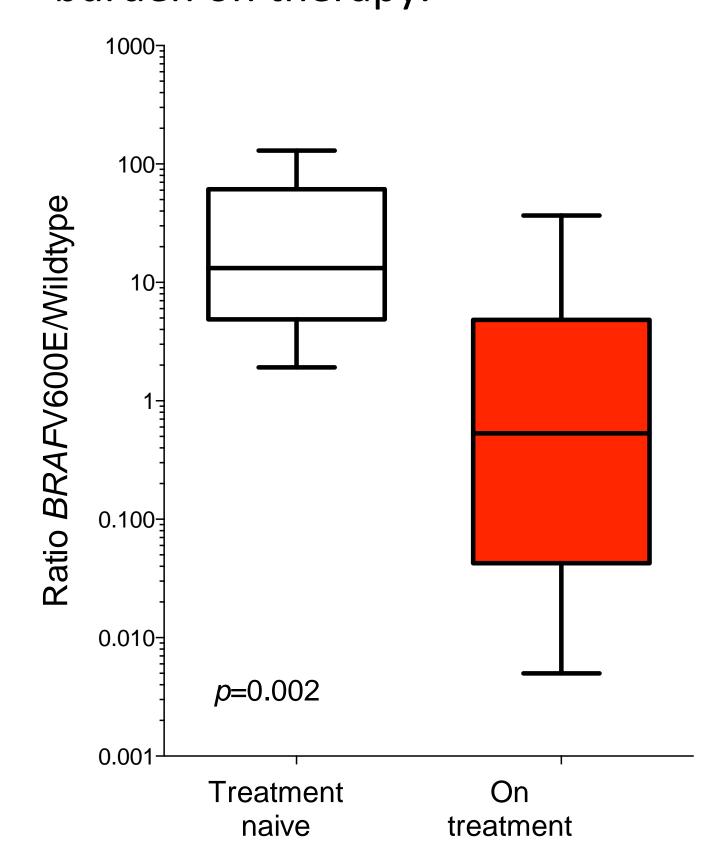
Quantitative analyses of BRAFV600E mutation in urinary and plasma cfDNA reliably detects BRAFV600E mutation based on tissue genotype (analyses below include plasma and urine samples pretreatment and on treatment).



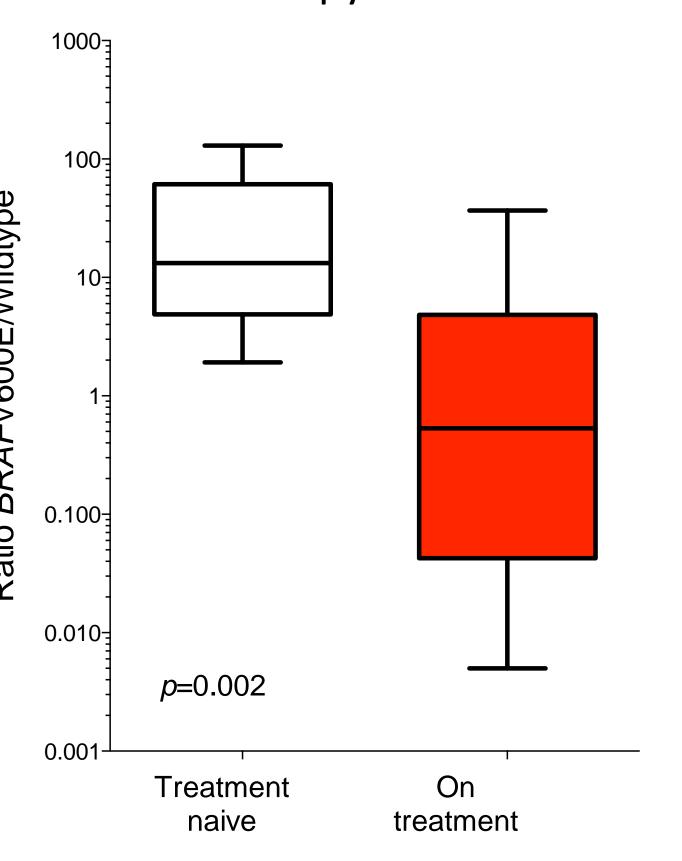
 Two patients with an indeterminate biopsy result but with clearly positive Urinary cfDNA BRAFV600E mutation subsequently were found to have BRAFV600E mutation in tissue with a repeat biopsy (orange box above).

 Quantitative cfDNA analysis of BRAFV600E burden in urine in treatment naïve versus on treatment samples detects decreased mutational burden on therapy.

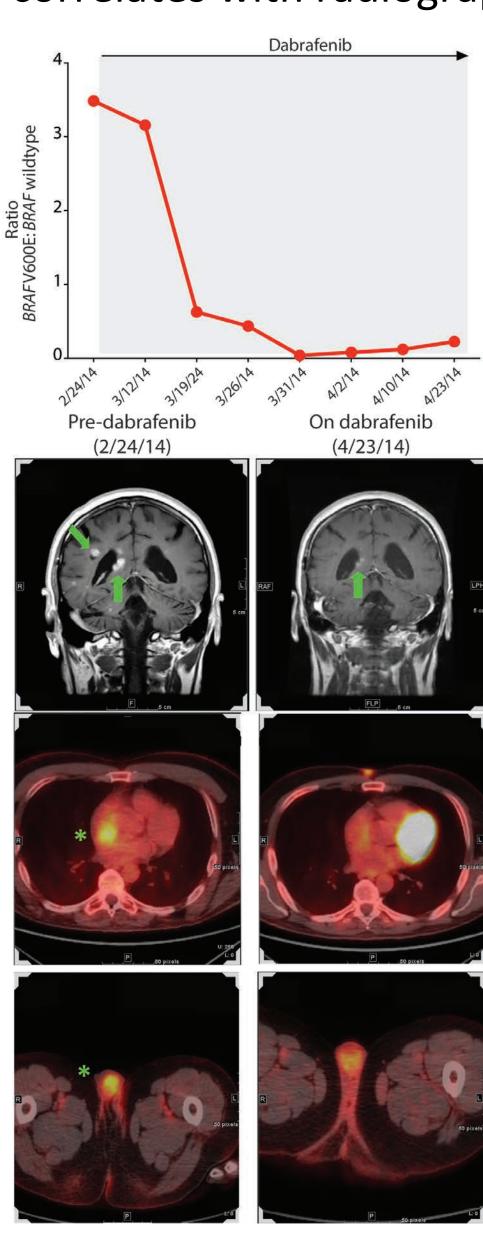
RESULTS

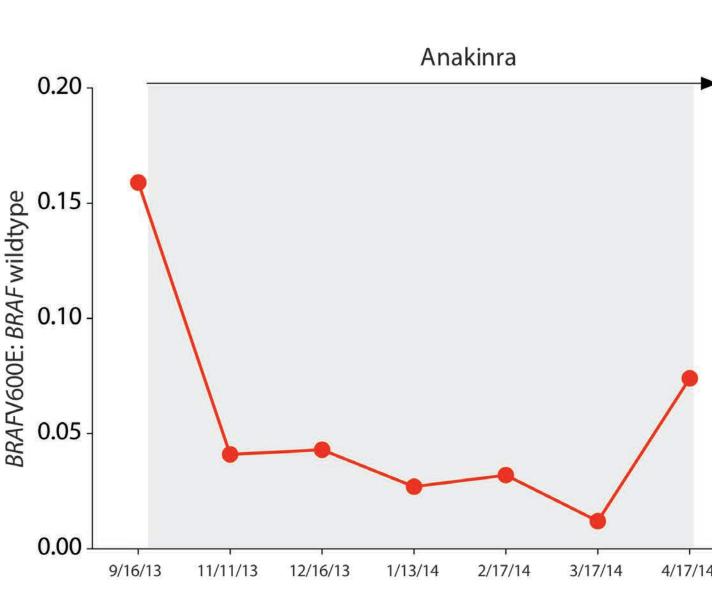


 Serial analysis of BRAFV600E mutation in urinary cfDNA of patients treated with vemurafenib reveals progressive decrement of BRAF mutant allele burden with therapy.



- Patient 4 Patient 5 Patient 6
- BRAFV600E burden in urine in correlates with radiographic response.
- BRAFV600E allele burden in urine changes dynamically with therapy.





### CONCLUSION

- BRAFV600E mutation detection in urine and plasma cfDNA provides a reliable means of detecting BRAFV600E mutations in histiocytosis patients.
- Serial monitoring of cfDNA BRAFV600E mutant allele burden from urine provides a convenient means of dynamically monitoring this unique disease.