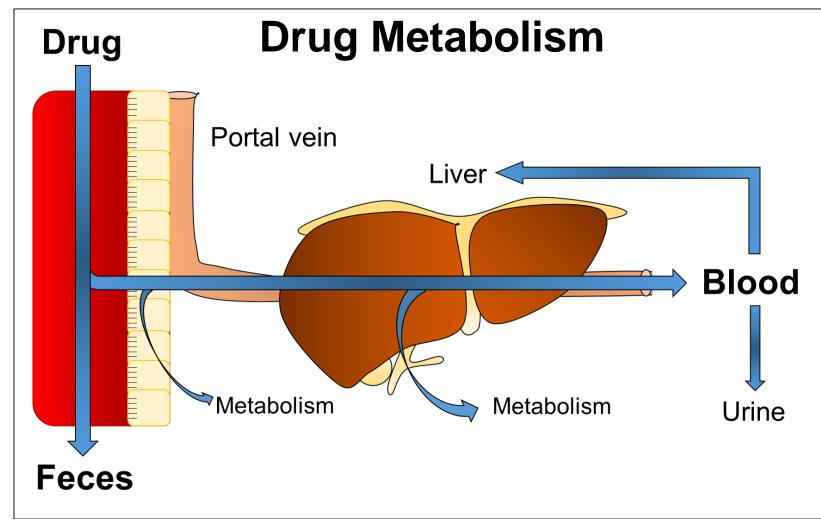
## Elucidating the Pharmacology and Toxicology of Anti-HIV Drug Metabolites

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- Drugs tend to be lipophilic
- Metabolism generates more polar products that can be excreted readily

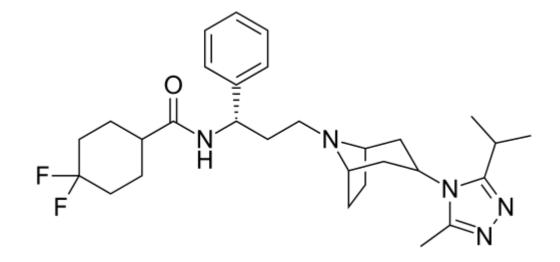
## The Cytochromes P450 (CYP)

- •A superfamily of heme-containing enzymes
- •Metabolize drugs and endogenous molecules
- •The general oxidation reaction
  - $NADPH + H^{+} + O_{2} + R \rightarrow NADP^{+} + H_{2}O + RO$

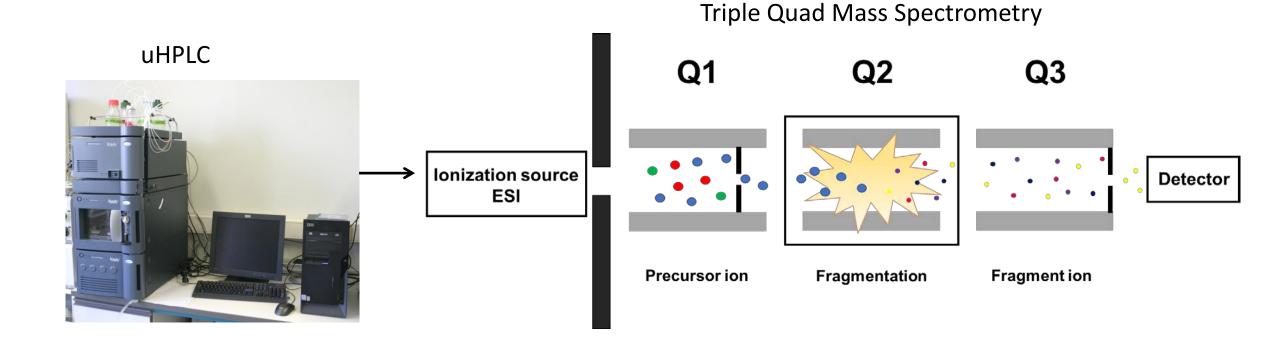
```
•Nomenclature
•Example: CYP3A4
Specific enzyme
Subfamily
```

## Maraviroc

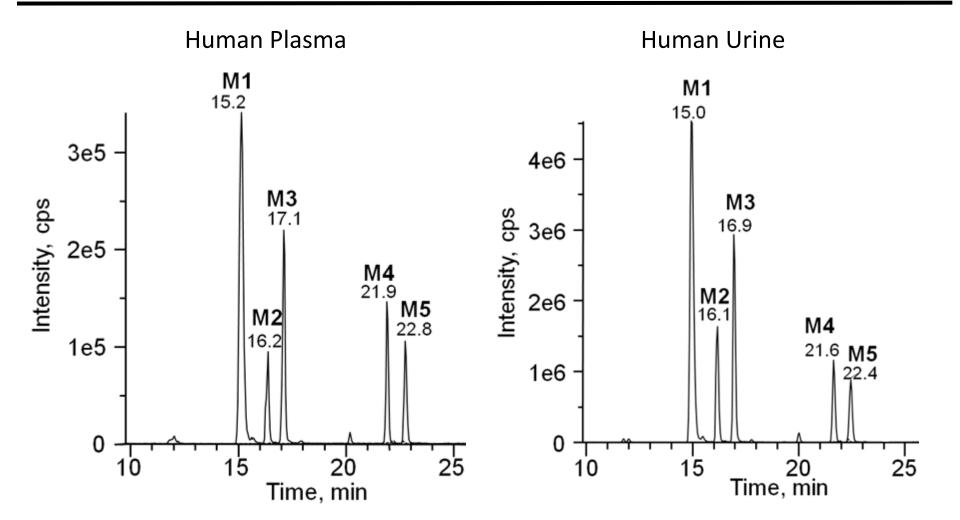
- Used in the treatment of HIV infection
- A chemokine receptor CCR5 antagonist
- A candidate for HIV prevention



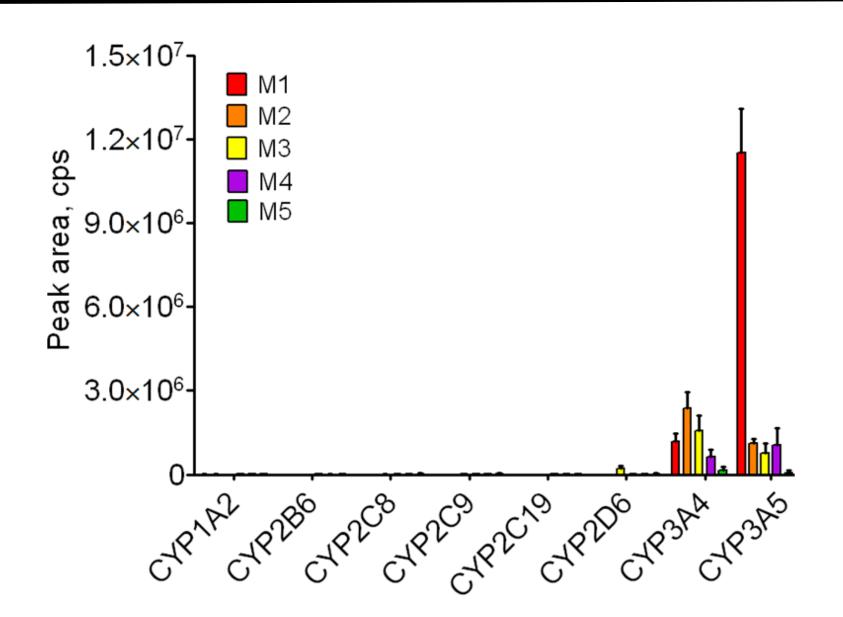
## Separation and Detection of Metabolites: LC-MS/MS



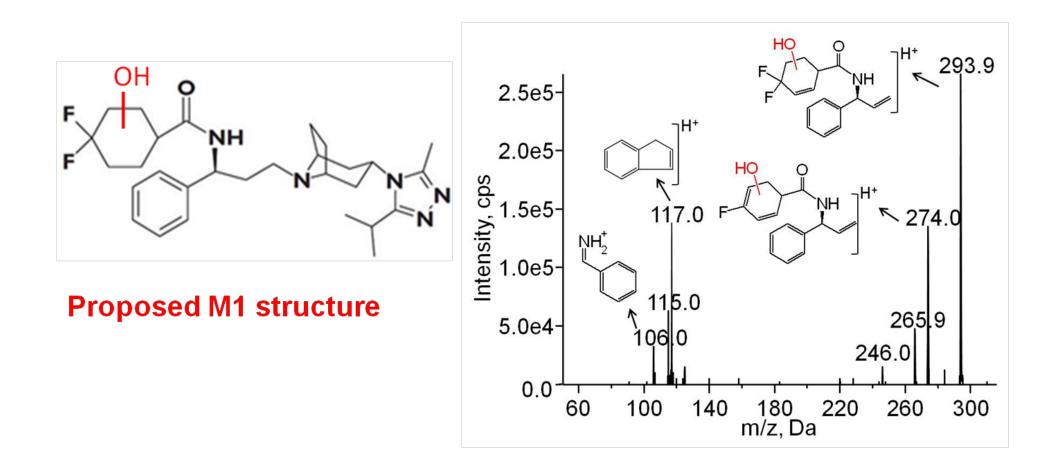
#### Separation of Oxidative Metabolites of Maraviroc



## **CYP34 and CYP3A5 Metabolize Maraviroc**



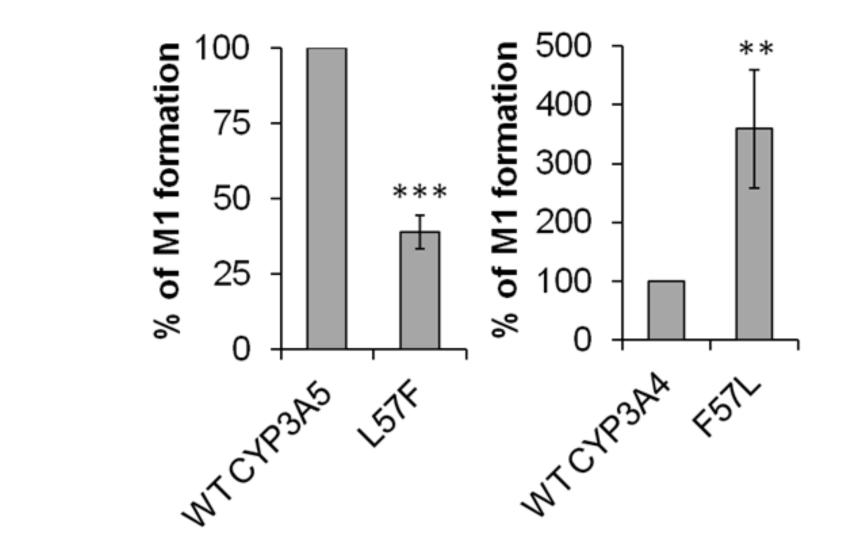
#### M1 Results from Oxygen Insertion on the Difluorocylcohexane Ring of Maraviroc



## **Divergent CYP3A4 and CYP3A5 Residues**

<b>3A4</b> 54 HKGFCMF 60	SRS4	
+G F	3A4 294 ELVAQSI IFIFAGYETTSS 312	
3A5 54 RQGLWKF 60	EL AQSI IFIFAGYETTSS	
Substrate Recognition Site (SRS1)	<b>3A5</b> 294 ELAAQSI IFIFAGYETTSS 312	
3A4 101 VFTNRRPFGPVGFMKSAISI AED123	SRS5	
VFTNRR GPVGFMKSAIS+AED	3A4 368 PI AMRLERV 376	
3A5 101 VFTNRRSLGPVGFMKSAISLAED123	P+A + RLER	
SRS2	<b>3A5</b> 368 PVA   RLERT 376	
3A4 204 VENTKKLLRFDFL 216	SRS6	
VE+TKK L+F FL	3A4 476 KLSLG GLLQP 485	
3A5 204 VESTKKFLKFGFL 216	KL GLLQP	
SRS3	3A5 475 KLDTQGLLQP 484	
<b>3A4</b> 238   CV FPRE 244		
++ FP++	Biochemistry 1998; 37:12536-45	
<b>3A5</b> 238 VS LFPKD244	J Biol Chem 2004; 279:38091-4 Xenobiotica 2006; 36:219-33	
	Curr Drug Metab 2011;12:684-700	

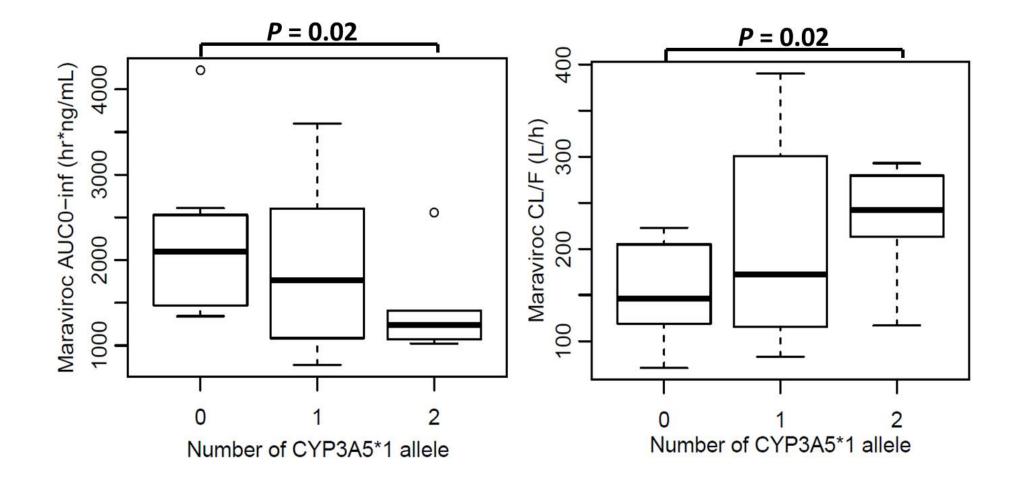
## Residue 57 Differentiates CYP3A4 vs. CYP3A5 M1 Formation



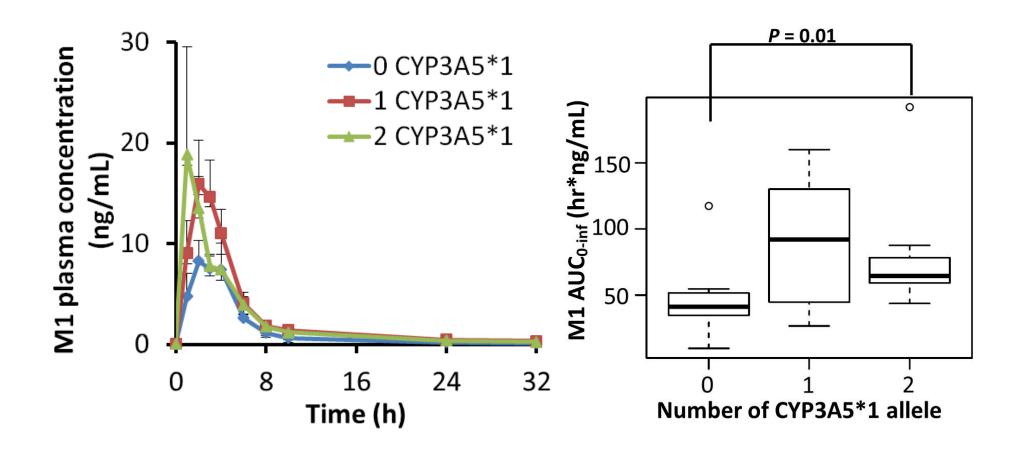
## **Study Design**

- Twenty four healthy volunteers (N=8 in each group)
  - Homozygous \*1 (two \*1 alleles)
  - Heterozygous (one \*1 allele and one variant allele\*3, \*6, or \*7)
  - Dysfunctional (two variant alleles)
- **Treatment**: a single oral dose of 300 mg maraviroc
- Blood collection: predose and 1, 2, 3, 4, 6, 8, 10, 24 and 32 h after the dose
- Concentration measurement
  - LC-MS/MS

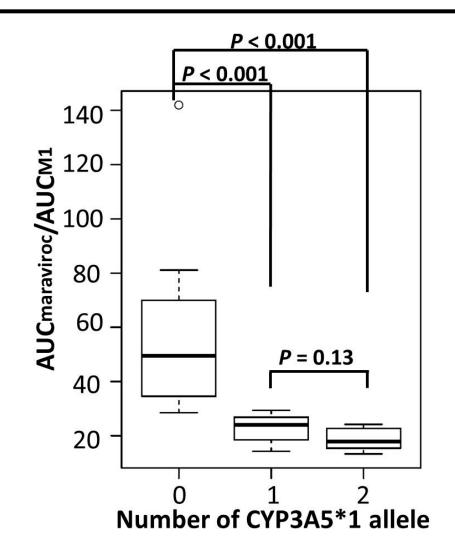
#### Decreased Maraviroc Exposure and Increased Clearance in the CYP3A5 Homozygous \*1 Group



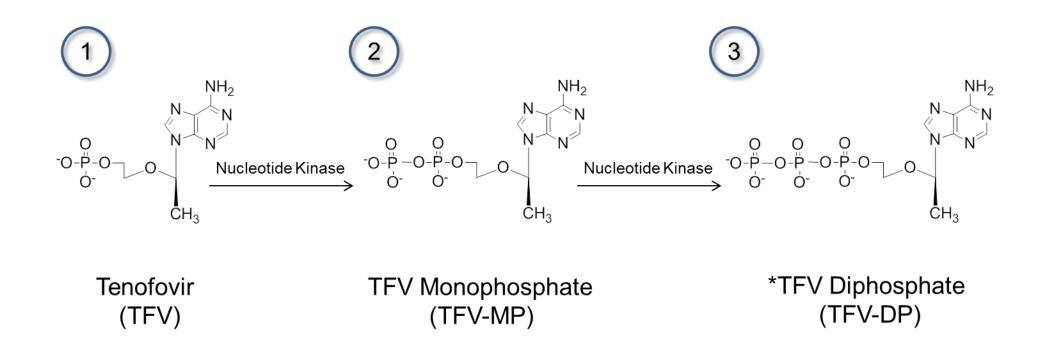
#### Decreased M1 Formation by CYP3A5 Dysfunctional Group



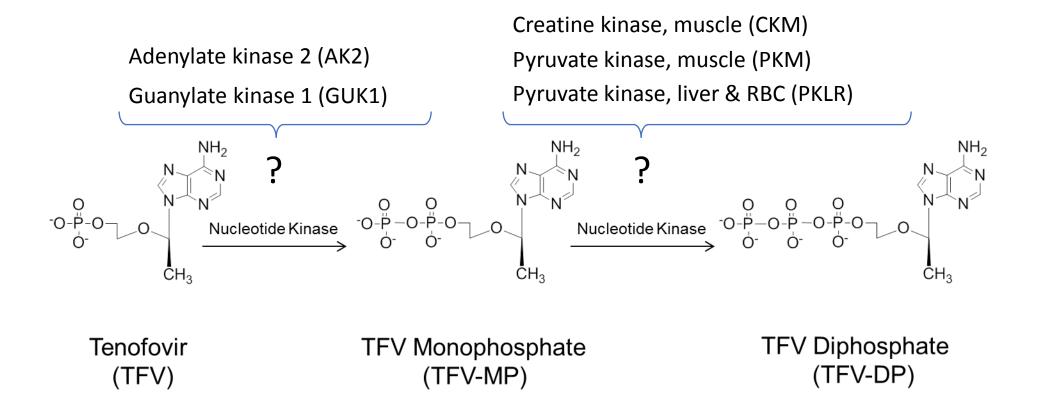
#### Lower AUC Ratio (maraviroc/M1) in CYP3A5\*1 Allele Carriers



#### **Tenofovir is a Nucleotide Reverse Transcriptase Inhibitor**



#### **Candidate Nucleotide Kinases**



#### siRNA Knockdown of Nucleotide Kinases

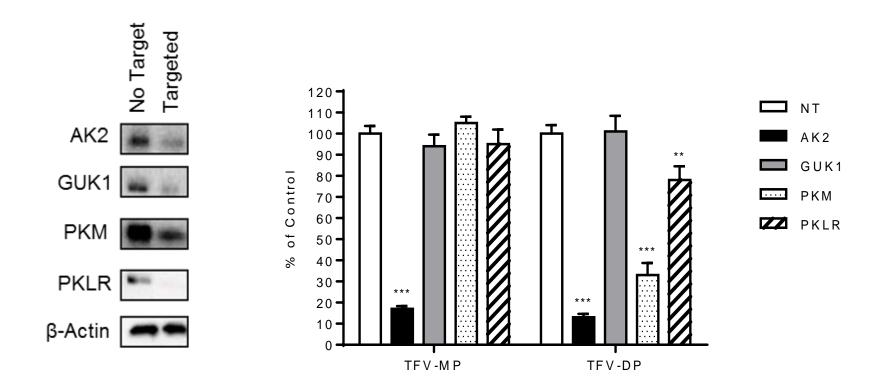


- Cells and tissues:
- 1. Peripheral blood mononuclear cells (PBMC)
- 2. Colorectal tissue
- 3. Vaginal tissue

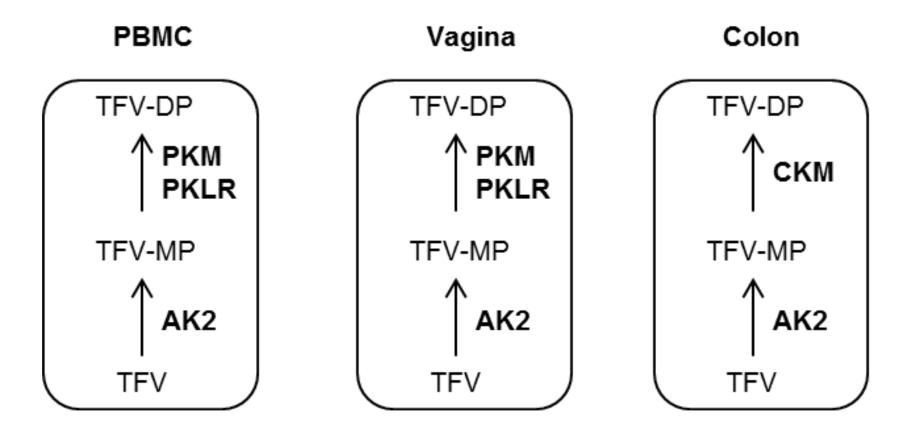


- Delivered siRNA to cells and tissues in culture
- Followed by incubation with TFV
  - Detected TFV metabolites using ultra-high performance liquid chromatography-tandem mass spectrometry

#### AK2, PKM, and PKLR Contribute to Metabolite Formation in Peripheral Blood Mononuclear Cells (PBMC)

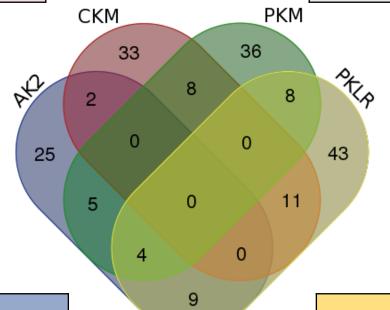


## **Tenofovir is Activated in a Tissue-Specific Manner**



Creatine Kinase, Muscle (CKM)			
Country	Individuals Sequenced	Individuals with one or more variant	% of individuals with one or more variant
United States	582	23	4%
South Africa	231	17	7%
Uganda	24	5	21%
Thailand	171	9	5%
Total	1,008	54	5%

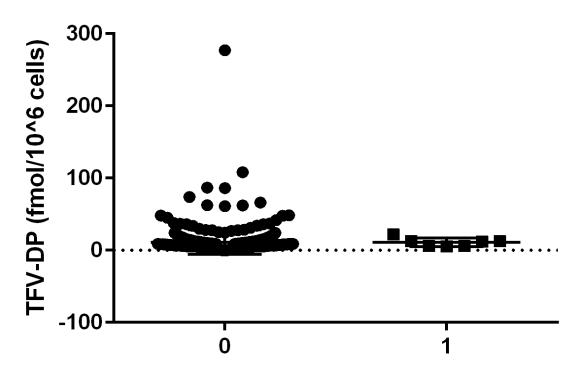
Pyruvate Kinase, Muscle (PKM)			
Country	Individuals Sequenced	Individuals with one or more genetic variant	% of individuals with one or more variant
United States	582	30	5%
South Africa	231	15	6%
Uganda	24	5	21%
Thailand	171	11	6%
Total	1,008	61	6%



Pyruvate Kinase, Liver and Red Blood Cell (PKLR)			
Country	Individuals Sequenced	Individuals with one or more genetic variant	% of individuals with one or more variant
United States	582	46	8%
South Africa	231	16	7%
Uganda	24	3	12.5%
Thailand	171	10	6%
Total	1,008	75	7%

Adenylate Kinase, 2 (AK2)			
Country	Individuals Sequenced	Individuals with one or more genetic variant	% of individuals with one or more variant
United States	582	33	6%
South Africa	231	7	3%
Uganda	24	3	12.5%
Thailand	171	2	1%
Total	1,008	45	4%

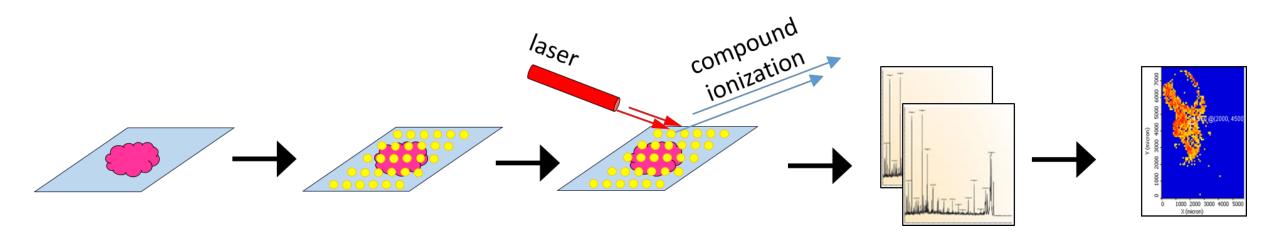
## Tenofovir Diphosphate Levels In Genotyped Individuals



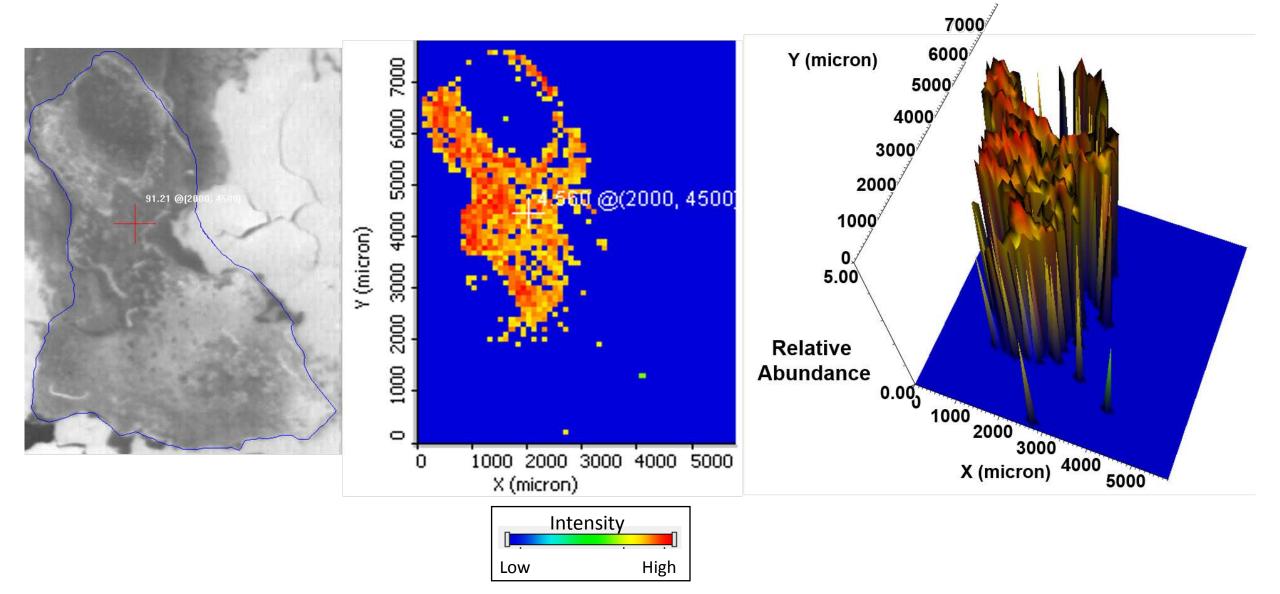
Individuals with no	
Predicted AK2 Deleterious Variants Detected	
Individuals with Predicted AK2 Deleterious Variants Detected	

Number of AK2 Deleterious Variants Detected

## Matrix-assisted laser desorption/ionization coupled to mass spectrometry (MALDI-MS) for understanding spatial distribution



# MALDI-MS/MS Reveals Heterogeneity in TFV distribution within Cervical Tissue



## Acknowledgements

Past/Present Lab Members: Lindsay Avery Philip Cox Dominique Figueroa Carley Heck Elisabeth Hersman Julie Lade Yanhui Lu Erin Madeen Joseph Tillotson Elaine To

<u>Collaborator:</u> Craig Hendrix Senior Colleagues: Zeruesenay Desta Charlie Flexner Evan Kharasch Deanna Kroetz Ken Thummel

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