Discovery of Genetic Variants of the Kinases that Activate Tenofovir in a Compartment-Specific Manner

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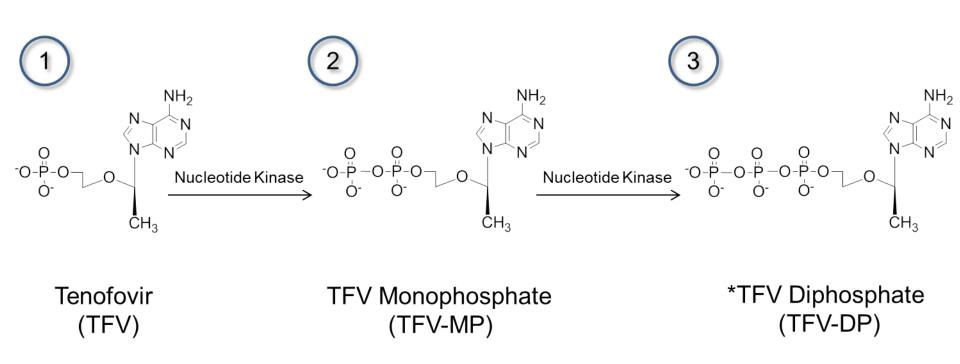
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Tenofovir Requires Activation by Nucleotide Kinases

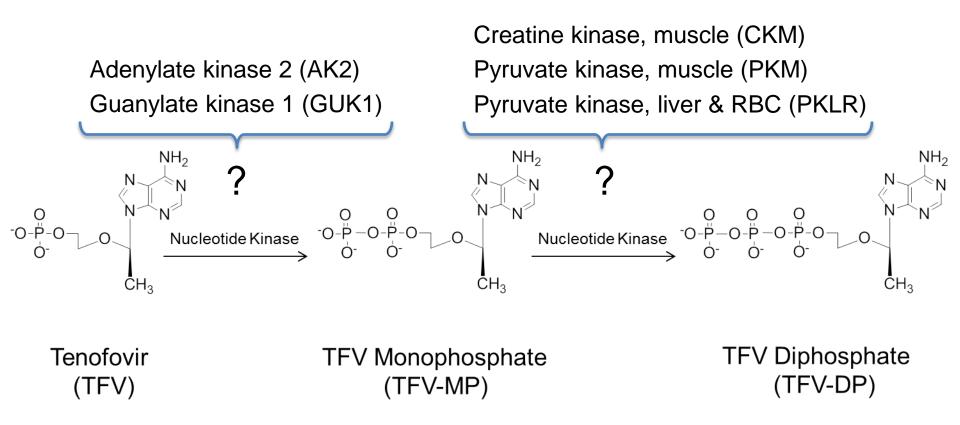


Which nucleotide kinases contribute to tenofovir activation in cells and tissues at risk of HIV infection?

Nucleotide Mono- and Di-Phosphate Kinase Isoforms to be Investigated

- Adenylate Kinases, ATP + AMP \rightarrow 2 ADP
 - 9 isoforms with differential tissue distributions and subcellular localizations
- **Guanylate Kinases**, ATP + GMP \rightarrow ADP + GDP
 - 3 nucleotide phosphorylating isoforms
- Nucleotide Diphosphate Kinases, NTP + NDP ↔ NDP + NTP
 - 4 enzymatically active isoforms
- Creatine Kinases, ATP + Creatine \rightarrow ADP + Phosphocreatine
 - Cytosolic and mitochondrial isoforms
- **Pyruvate Kinases**, Phosphoenolpyruvate + ADP → Pyruvate + ATP
 - 4 isozymes result from differential splicing

Candidate Nucleotide Kinases



siRNA Knockdown of Nucleotide Kinases

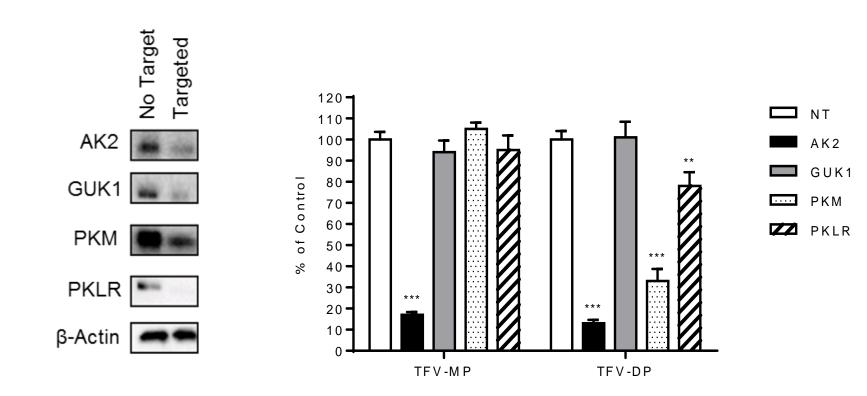
Human Samples

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

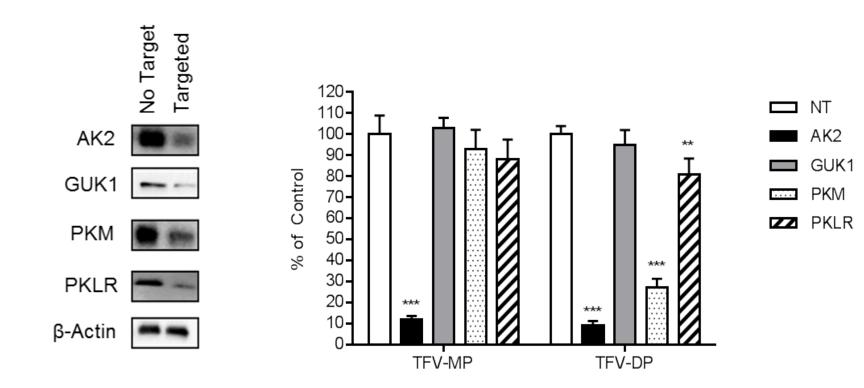


- Delivered siRNA to cells and tissues in culture
- Followed by incubation with TFV
 - Detected TFV metabolites using ultrahigh performance liquid chromatographytandem mass spectrometry

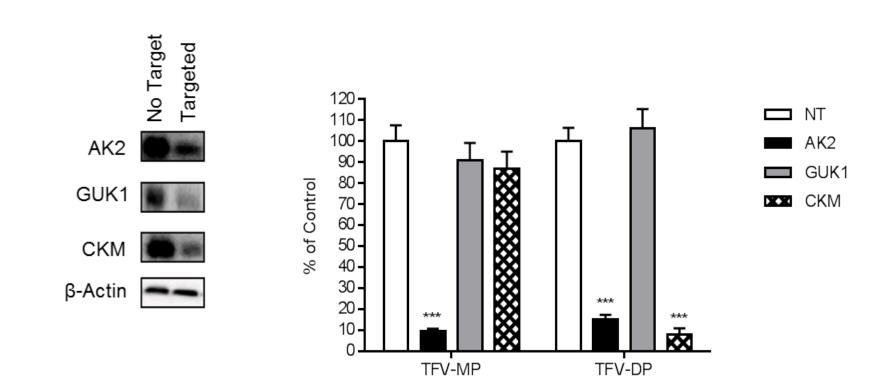
AK2, PKM, and PKLR Contribute to Metabolite Formation in Peripheral Blood Mononuclear Cells (PBMC)



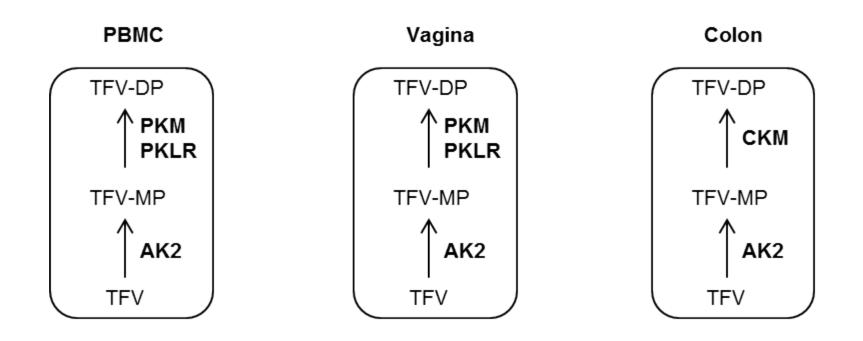
AK2, PKM, and PKLR Contribute to Metabolite Formation in Vaginal Tissue



AK2 and CKM Contribute to Metabolite Formation in Colorectal Tissue



Tenofovir is Activated in a Tissue-Specific Manner



Do human genetic variants exist in the genes encoding these nucleotide kinases?

Next-Generation Sequencing of MTN-001 Samples

18 CKM variants

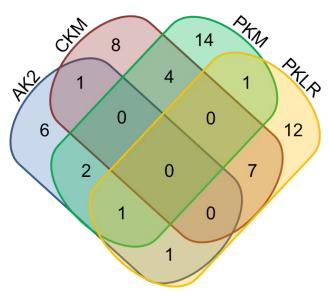
- 17 participants
- 3% frequency of deleterious missense variants – 1 USA, 1 SA, 2 UGA participants

19 PKM variants

- 19 participants
- Unable to predict phenotype of missense variants



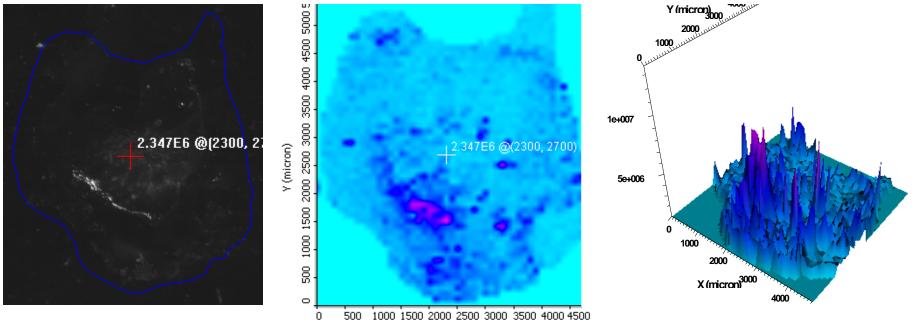
- 11 participants
- 2% frequency of deleterious missense variants – 1 USA, 2 SA participants



22 PKLR variants

- 21 participants
- 3% frequency of deleterious missense variants – 3 USA, 1 SA participants

MALDI-MS Imaging of TFV



X (micron)

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