## VIP-H: Very Important Pharmacogenes in Hmong

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Dates: 2016-2018

## Purpose:

- Measure extent of pharmacogenes known to affect medication doses in the Hmong community
- Evaluate people's responses to their pharmacogenomic results

## Methods:

- Genetic analysis in 198 Hmong adults to identify 24 genetic variations relevant to 20 medicines
- Focus group discussions with 27 participants about their results

## Results

- Significant genetic variations for two medicines (warfarin and clopidogrel), some variations for 8
  medicines and little variations for 10 medicines.
- Community members are interested in results. They perceive that heredity (ncaj cev) influences their composition (roj ntshav), which affects how medicines work for them.
- <u>Poster</u>
- Application: Communicating results at community events and VIP-Hmong website
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