In vitro and in vivo functional testing of SNPs in the 3'UTR of CYP2B6

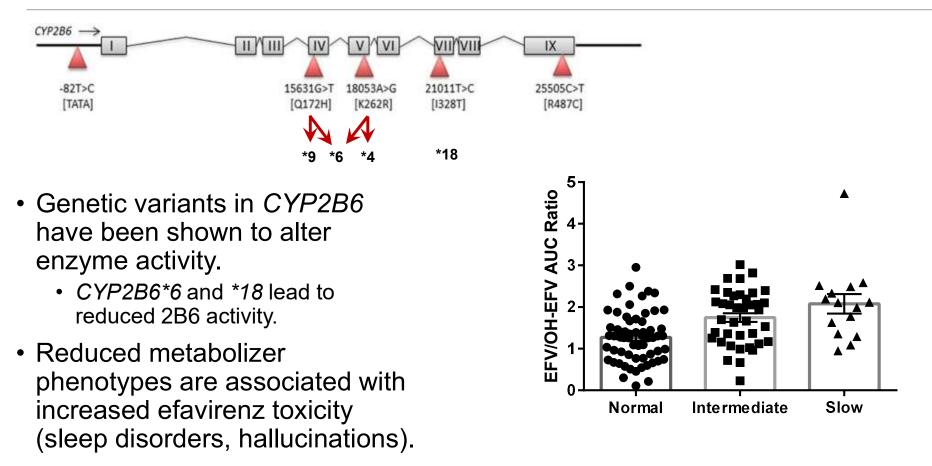
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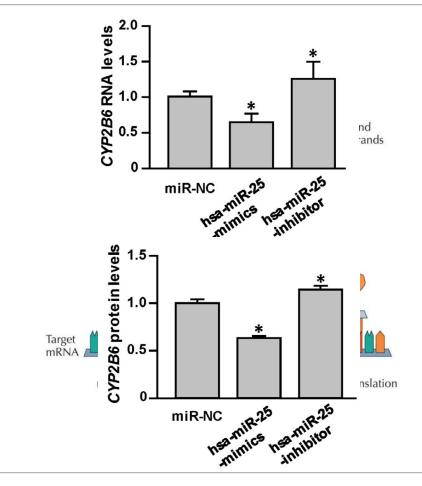
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Background - CYP2B6



Background - microRNA



- microRNA Length: 17-22 nucleotides
- Mechanism of action: bind to the 3'UTR of target mRNA
 - Seed sequence critical for miRNA targeting
- Scope: 2588 mature miRNAs identified in humans.
- miRNAs have been predicted and experimentally validated to target many genes, including CYPs.

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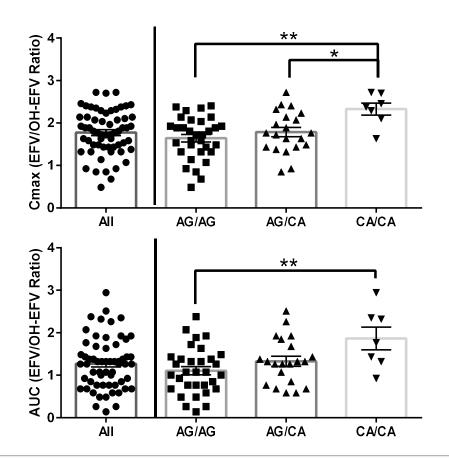
Hypothesis

• Variability in efavirenz pharmacokinetics are associated with genetic variants that alter miRNA regulation of *CYP2B6.*

3'UTR variants are associated with CYP2B6 activity in vivo

- Retrospectively sequenced the *CYP2B6* 3'UTR of 200 healthy human volunteers administered a singe dose of efavirenz (100/600 mg).
- CYP2B6 activity for 114 volunteers: Cmax and AUC_{0-48hr} (EFV/8OH-EFV)
- Two variants, rs12979270 and rs12979898 variants were in perfect LD in our population.
 - rs70950385

Among normal CYP2B6 metabolizers, the rs70950385 variant is associated with decreased CYP2B6 activity *in vivo*.

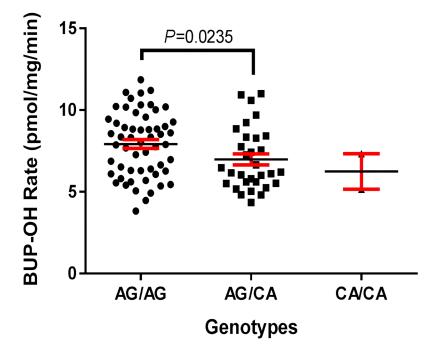


⁽Higher ratio=less metabolism) 5

rs70950385 variant is associated with CYP2B6 activity in vitro

 Sequenced the 3'UTR of 90 liver tissue samples; CYP2B6 activity was determined in microsomal preparations using bupropion as a probe.

Among all human liver samples tested, rs70950385 variant is associated with decreased CYP2B6 activity.



Proposed Mechanism

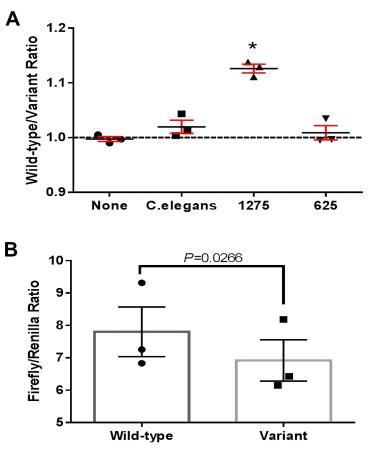
 rs70950385 (AG→CA) variant predicted to create a miRNA binding site for miR-625-5p and miR-1275.

Α Wild-type 5' A C A C A C U G U A G U C U U C C C C A G 3' CYP2B6 mRNA 3' C C U G A U A U C U U G A A A G G G G G A 5' hsa-miR-625-5p Variant 5' A C A C A C U G U A G U C U U C C C C C A 3' CYP2B6 mRNA 3' C C U G A U A U C U U G A A A G G G G G A 5' hsa-miR-625-5p Wild-type Β 5' A C A C A C U G U A G U C U U C C C C A G 3' CYP2B6 mRNA I I I I I I3' C U G U C G G A G A G G G G G U G hsa-miR-1275 Variant 5'ACACACUGUAGUCUUCCCCCA3' CYP2B6 mRNA I I I I I I3' C U G U C G G A G A G G G G G U G hsa-miR-1275

rs70950385 (CA allele) creates a miRNA binding site

- Created firefly luciferase plasmids containing either wildtype or variant miRNA binding sites.
- Transfected ± predicted miRNA or control miRNA in HepG2 cells.
- Firefly luciferase activity normalized to Renilla luciferase control.





Conclusions

- The rs70950385 variant decreases CYP2B6 activity in vitro and in vivo.
- Genetic variants in the 3'UTR have the ability to alter enzyme activity by interfering with miRNA binding.
- Genetic variants in the 3'UTR may explain variation in metabolism and drug response.

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