Characterizing Pharmacogenomic Drug Response: A Pilot Study

Samantha Adams, Megan Reineck, France Moosa
The University of Findlay

Introduction

- •Pharmacogenomic testing is a growing topic of interest especially in ambulatory pharmacy practice which may allow pharmacists to make more precise and safer medication recommendations for patients.
- •Pharmacogenomic testing will allow the profession to offer more individualized medication therapy for our patients.
- •Cytochrome P450 enzymes are essential for the metabolism of many medications. Although this class has more than 50 enzymes, six of them metabolize 90% of drugs, with two most significant enzymes being CYP3A4, and CYP2D6²
- •Each medication may take a different pathway in metabolism such as degradation or prodrug models and this may affect whether or not a medication is helpful, harmful, or ineffective for a specific patient

Objectives

- •Determine the feasibility of performing genetic testing for medication therapy based on genetic markers, specifically CYP2D6
- •Determine patients that are rapid, normal, or poor metabolizers in reference to CYP2D6

Methods

Collection

- Determine qualifying patients (>18 years old, on > 2 prescription medication, patient under one of our contracting physicians) and obtain consent from patients
- Perform buccal swab and de-identify using numbers and letters

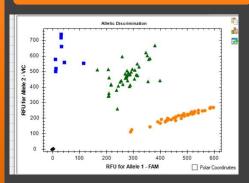
Isolation

 Remove the DNA from the buccal swabs using PBS and protease solutions from QIAGEN followed by heat shocking to elute DNA and clear contaminants

RT-PCR

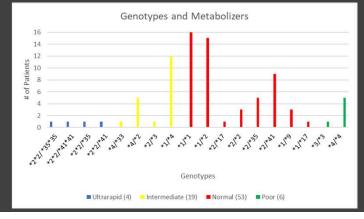
- Plating of samples included using MAD mix with the specific allele primer, followed by the pipetting of each sample into an individual well of the plate. 1 plate was completed for each specific allele
- There were 90 patients whose DNA was run in the PCR machine. 6 non-template control wells were used to ensure that the mix being added to the DNA was pure.
- Completed plates were then run through the BIO-RAD CFX96-Real time system (RT-PCR machine) to amplify DNA for each patient for each specific allele.

Results



2D6 Allele Variant	Variant Count (N=196)	Variant Frequency
*1	63	0.35
*2	59	0.3278
*3	3	0.0167
*4	31	0.167
*9	6	0.0333
*17	3	0.0167
*33	1	0.0056
*35	14	0.0778
*41	16	0.0889

- The picture is the result of CYP2D6*2 from the RT-PCR
- The table shows the different allele variants and the frequencies among different alleles
- The graph shows the different genotypes of the patients along with their classification



Discussion

- Study weaknesses included: low sample size, multiple researchers isolating samples (increased human error), genotyping costs a great deal of time and money
- Study Strengths: multi-tiered student learning approach, most common alleles were tested for, as well as a few that were unique
- The current processes used are still being optimized and these results are preliminary

Conclusions

- Majority of patients were normal metabolizers, followed by intermediate, poor, then ultrarapid metabolizers as seen in the charts
- In the future, chart review can occur for the patients to determine if they are taking medications that are conducive to their genome.
- Implementation of pharmacogenomic testing such as this into an ambulatory care clinic can help pharmacists and physicians deliver individualized care to patients.

References

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