











Contact information: cdzhang@cmmt.ubc.ca

# Gene Regulatory Region Variations Potentially Modulating the Drug Metabolism Response

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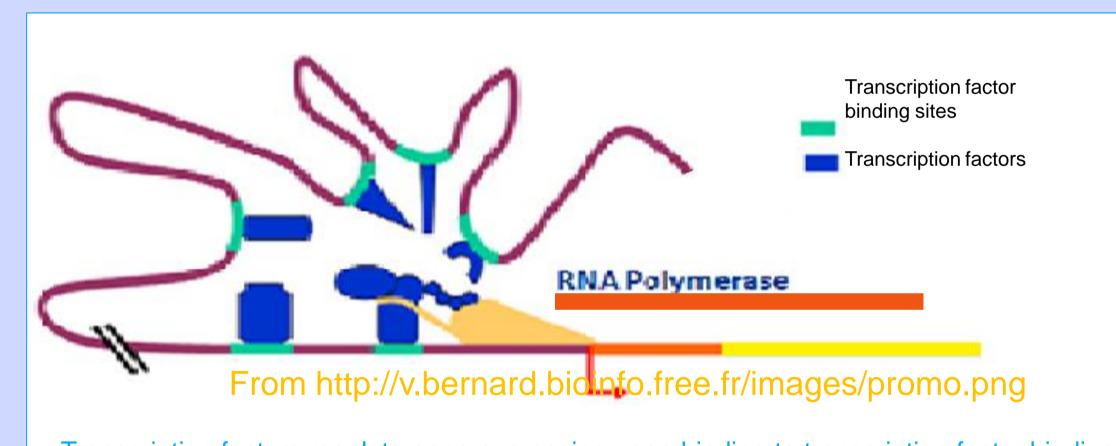
## Motivation

Adverse drug reactions (ADRs) are among the most common causes of hospitalization in paediatric patients. Genetic factors account for 20% to 95% of ADRs.

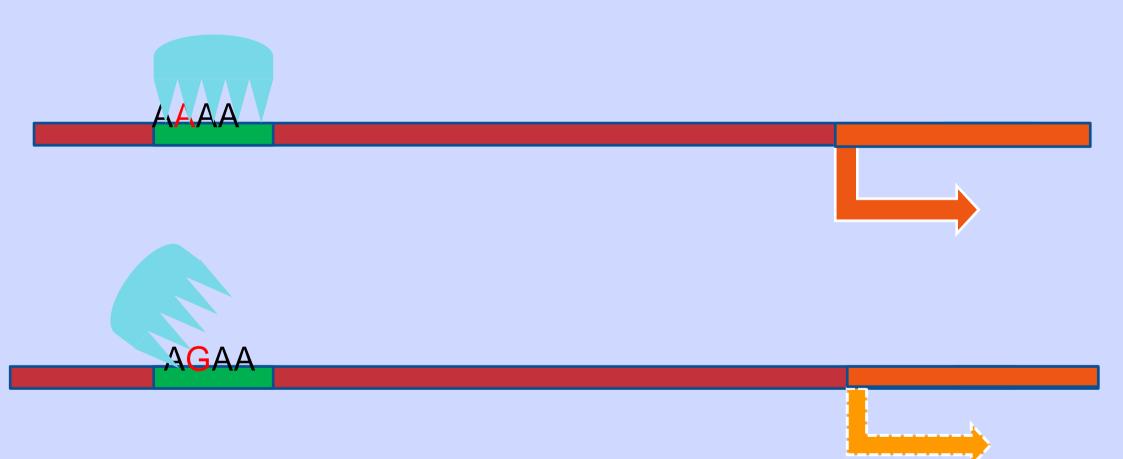


From www.surryothai.acth/node/1238

Individual differences in gene regulatory regions can lead to drug metabolism differences



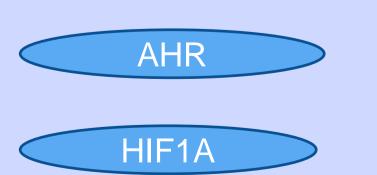
Transcription factors regulate gene expression upon binding to transcription factor binding sites



Genetic differences in TFBSs may block TF binding and alter gene expression.

### Focus

3 transcription factors (TFs) are known to have regulated the expression of a whole range of drug metabolizing genes.



NFE2L2

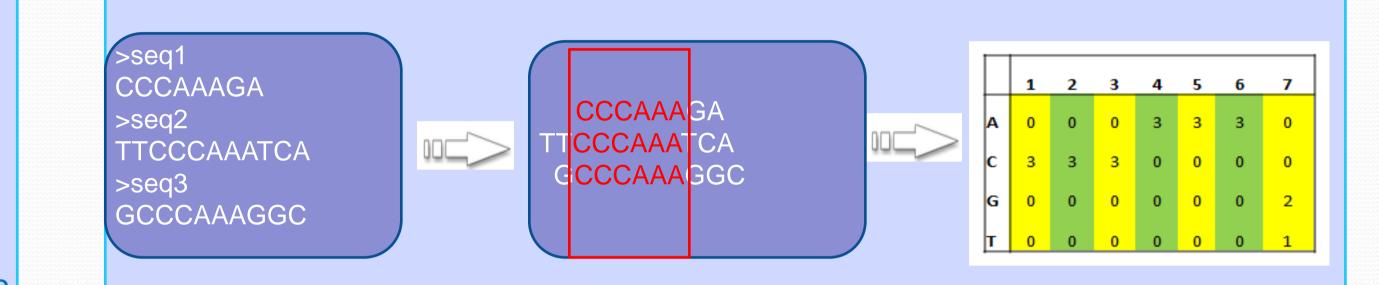
Responses to aryl hydrocarbon compounds

Responses to hypoxia

Responses to antioxidant and xenobiotics

#### Methods & Results

Building a transcription factor binding site (TFBS) profile for each of the 3 transcription factors



Gene Set Size Position Frequency Matrices Transcription factors NFE2L2 627 sequences 2077 sequences HIF1A1

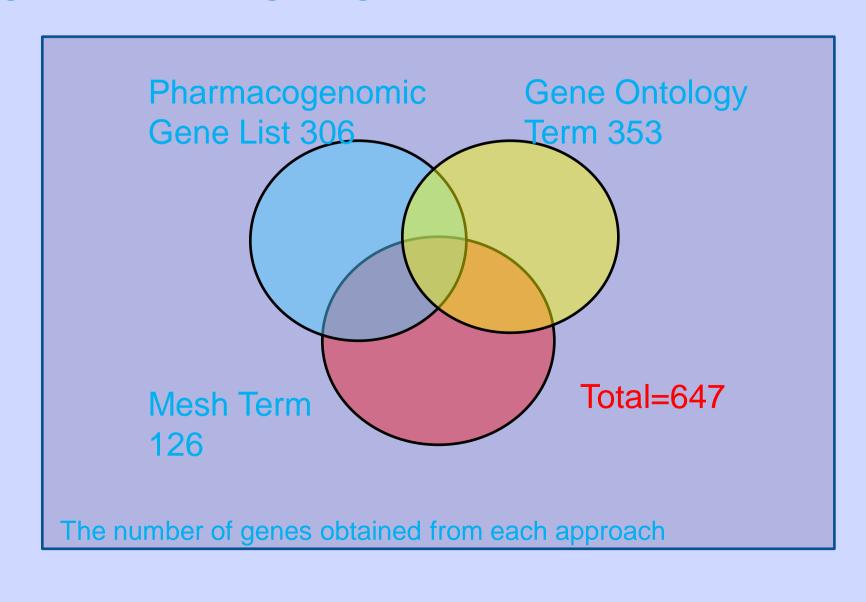
AHR<sup>2</sup> ? sequences

Prediction of over-represented transcription factor binding sites based on position frequency matrices(

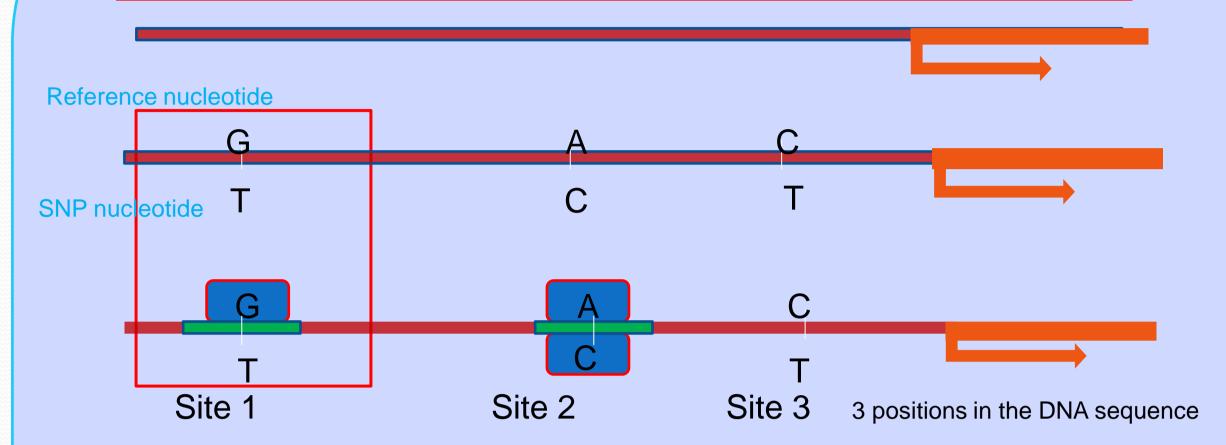
#### Compiling a list of drug-metabolizing genes

- ➤ Pharmacogenomic Gene List obtained from Dr. Colin Ross, Hayden Lab, CMMT
- ➤ Gene Ontology term search using BioMart http://www.ensembl.org/biomart
- Finding genes relating to drug metabolism using Mesh term search tool developed by Warren Cheung, Wasserman Lab

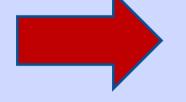
http://dgene.oicr.on.ca/digenei/gene\_browse.html#



Finding the transcription factor binding sites overlapping Nucleotide Polymorphisms (SNPs)



Search for SNP sites or reference sites that match the TFBS exclusively The unique form of nucleotide at the specific position which may be able to account for drug metabolism differences.



Produced a prioritized list of regulatory region variations that may be responsible for individual

differences in drug metabolism

# **Clinical Implications**

http://i.ehow.com/images/a05/d1/bg/safe-tylenol-infant-drops-acetaminophen\_-200X200.jpg

- Highlight potential regulatory sequences in the drug metabolizing genes for personalized genome analysis
- Predict and therein prevent adverse drug reactions based on individual genome scan
- "Ideal" drug design tailored to individual drug metabolising system.

#### Reference

Kung A. and Xia X. Preferential Binding of HIF-1 to Transcriptionally Active Loci Determines Cell-Type Specific Response to Hypoxia. Genome Biology RII3-RII3.12(2009);

Pansoy A. et.al. 3-Methylcholoanthrene Induces Differential Recruitment of Aryl Hydrocarbon Receptor to Human Promoters. Toxicological Sciences ,1-32 (2010);

Portales-Casamar E. et. al. PAZAR: a Framework for Collection and Dissemination of Cis-regulatory sequence Annotation. Genome Biol (2007)