### Wolff-Parkinson-White Syndrome and PRKAG2

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http://www.beatmap.net/portfolio-detail/human-cardiovascular-system-3d-renderings/

### What causes Wolff-Parkinson-White?



Mayo Clinic, Wolff-Parkinson-White Syndrome 2011

### Current Disease Treatment

Medications = Adenosine, and antiarrhythmic drugs

Electric cardioversion therapy

Radiofrequency catheter ablation



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### **Missense Mutations in PRKAG2**

PRKAG2 = Protein Kinase AMP-Activated Gamma 2

Mutation alters activity of 5'-AMP-activated protein kinase subunit gamma 2 protein (**AAKG2**)



### AAKG2 Protein

#### **Heterotrimeric (AMPK)**

- Catalytic subunit = alpha
- Regulatory subunits = beta and gamma

#### Gamma Subunit

Contains four CBS domains



### PRKAG2 Gene Ontology

#### **Biological Processes**

**ATP Biosynthesis Process Carnitine Shuttle** Cell Cycle Arrest **Cellular Lipid Metabolic Process** Fatty Acid Biosynthetic Process **Glycogen Metabolic Process Insulin Receptor Signaling** Pathway Intracellular Protein Kinase Cascade **Regulation of Protein Kinase** Activity **Regulation of Fatty Acid** Oxidation **Regulation of Glucose Import Regulation of Glycolysis** 

#### **Molecular Function**

ADP Binding ATP Binding CAMP-Dependent Protein Kinase Inhibitor Activity CAMP-Dependent Protein Kinase Regulator Activity Phosphorylase Kinase Regulator Activity Protein Kinase Activator Activity Protein Kinase Binding

#### **Cellular Component**

AMP-Activated Protein Kinase Complex Cytosol Nucleoplasm

### PRKAG2 Highly Conserved





### **CBS** Domains



#### Highly evolutionarily conserved

Often come in tandem pairs

Restore cellular **ATP** balance during metabolic stress

### Heart vs. No Heart



## Where are ATP-binding sites in CBS domains?



## Are these ATP-binding sites important in **PRKAG2 missense mutations**?

Human Mutation	Reduce AMP/ATP Binding	Structurally Close (Interacting)	
R302Q	YES	YES (YES)	
H383R	YES	YES (YES)	
T400N	YES	YES (YES)	
N488I	NO	NO (NO)	
R531Q	YES	YES (YES)	
R531G	YES	YES (YES)	
S548P	No Data	YES (YES)	

Xiao, B., Heath, R., Saiu, P., et al. (2007) Structural basis for AMP binding to mammalian AMP-activated protein kinase. *Nature* 449, 496- 500

## Will these missense mutations have the same effect in a model organism without a heart?



### Without Heart???

## Amino acids causing ATP- binding problems are conserved in *C. elegans*



**Hypothesis**: Dauer formation would be observed if human missense mutation introduced into heartless *C. elegans*.





Wallingford Lab (http://www.bio.utexas.edu/faculty/wallingford/Pages/Xenopus%20Links.html) http://triviascience.blogspot.com/2012/10/why-human-babies-are-so-cute.html

### What is STK11/LKB1?

**Protein Kinase** 

Serine/Threonine 11

Role in G1 cell cycle arrest

Tumor suppressor

Mutation causes Peutz-Jeghers syndrome

Activates AMPK-related proteins by **phosphorylation** 

# What is the function of STK11 in ATP regulation?



### STK11 activates AMPK



#### **Hypothesis**: Mutations in STK11 would inhibit ATP binding and alter development of the heart



## How would PRKAG2 be expressed if STK11 were mutated?

PRKAG2 Expression					
Tissue	Cardiac	Skeletal	Brain	Liver	
WT					
PRKAG2 Mutant					
STK11 Mutant					



**Underexpressed** 

### Conclusions

PRKAG2 and Wolff-Parkinson-White have been well studied/characterized

### CBS domains are found in all three domains of life = **highly conserved**

Mutations in PRKAG2 homologs may still have significant implications in heartless organisms

### **Future Directions**

Highly conserved--where did these CBS domains stem from?

Looking into the AMPK protein in order to potentially develop treatments for STK11 related diseases

Are there any other proteins affected by mutation to PRKAG2 gene?

## Questions?

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