# Embryology and cardiovascular genomics



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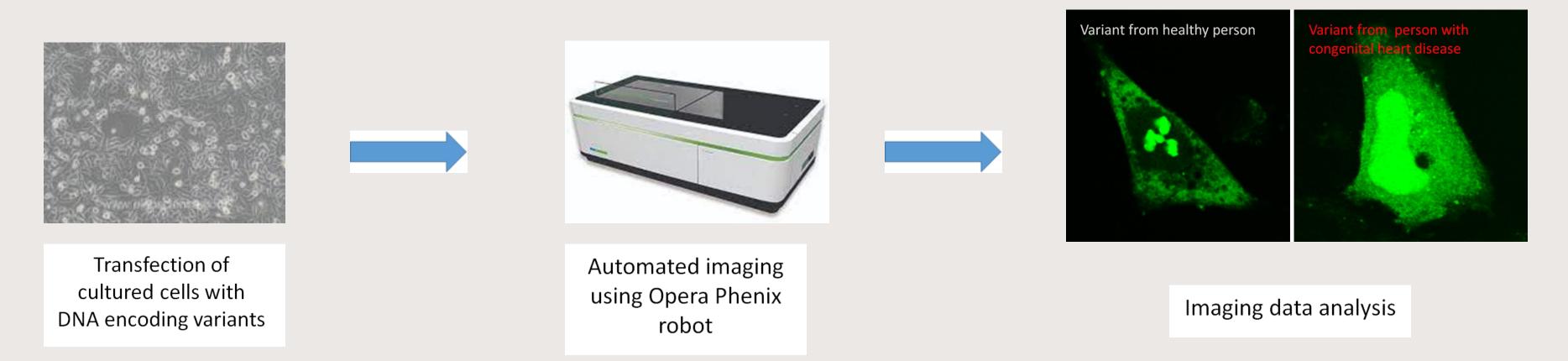
## Why are babies born with heart defects?

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Here at the Embryology laboratory at the Victor Chang Cardiac Research Institute, we are trying to answer this question. 1 in 10 babies are born with a heart defect. Current research tells us that gene mutations or environmental factors like diet, drugs or disease can cause these defects. But so far, we only have an answer for 25% of the families who are looking for answers. So, we've come up with a few ways to change the odds in our favour.

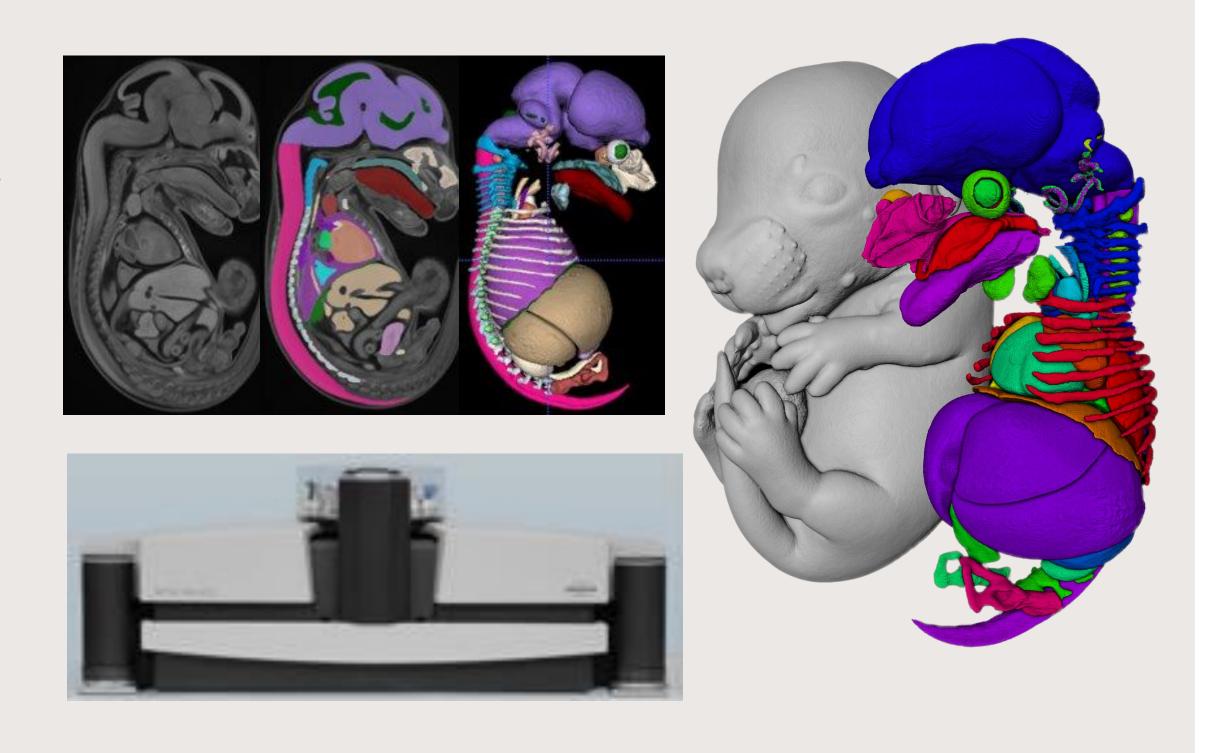
## Whole genome sequencing and functional genomics

Whole genome sequencing is a revolutionary technique that lets us identify all variations in a baby's DNA. Currently, we have sequenced the DNA of 156 families affected by congenital heart defects. In these families, we have found mutations genes that could be the cause of the heart defect. In the lab, we are using cell lines to test how these mutations could be affecting heart development.



## Mouse models of congenital heart defects

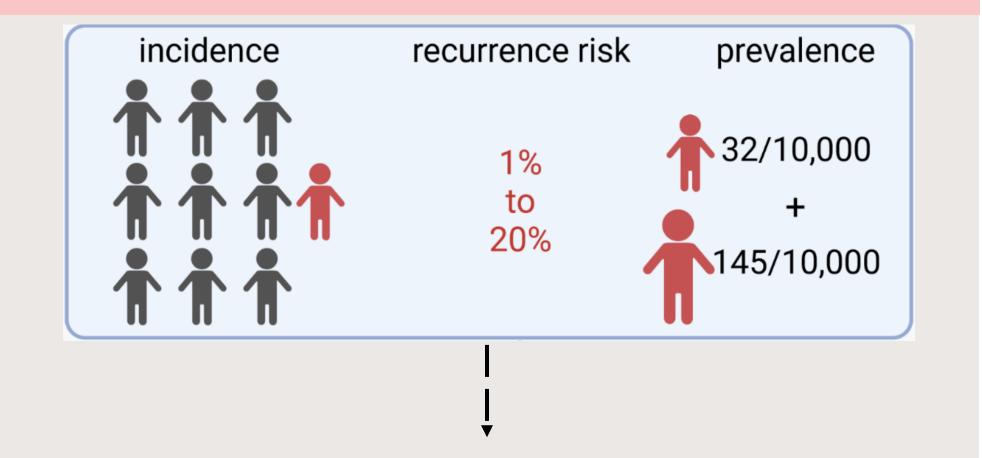
Our lab uses pioneering gene editing techniques to create mouse models with heart defects and analyse them using state of the art imaging techniques (Micro-CT) together with an automated phenotyping pipeline. Studying a human mutation in a mouse model is an essential way to prove that the gene is crucial for development and proper function of the heart.



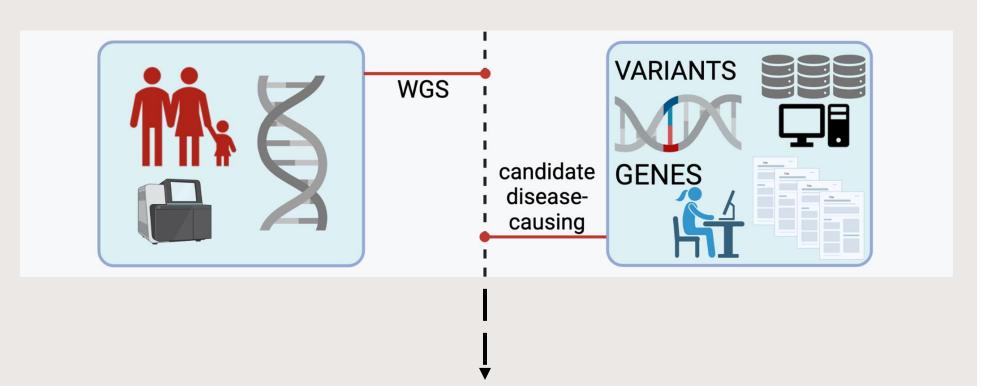
# Where you come in

Our research has lead to the discovery of new heart genes, new pathways needed for heart development, new ways to provide healthcare interventions for at-risk pregnant women. But there is still so much to do and to discover. We need more people to join us. If you're excited about learning about how a single nucleotide change can stop the heart from forming, then we're the lab for you.

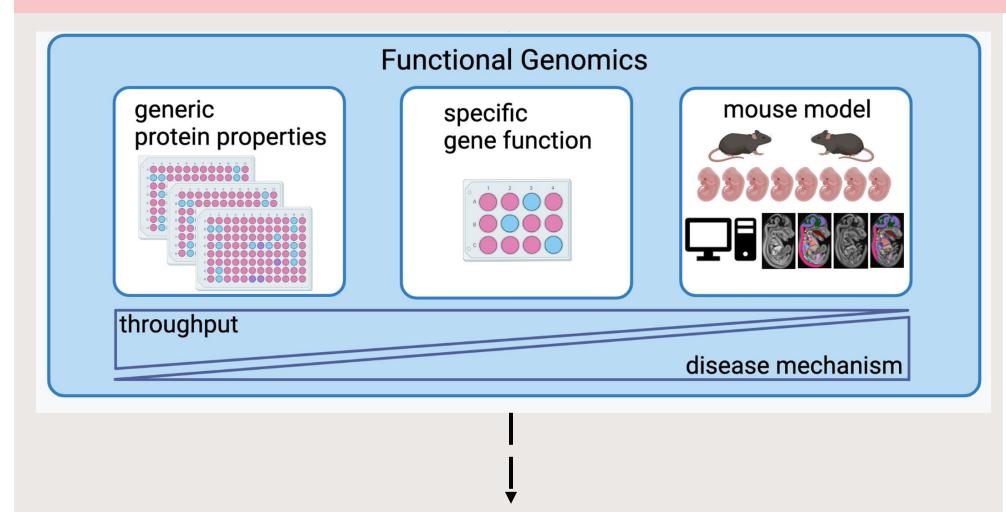
## 1. Recruiting patients and families with congenital heart defects



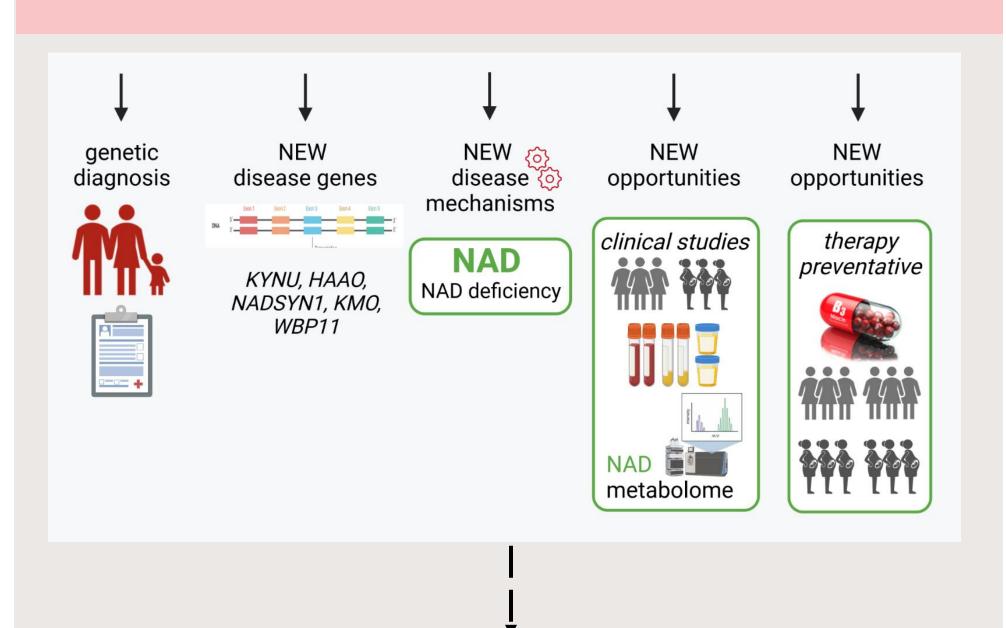
### 2. Genome sequencing analysis and variant identification



#### 3. Functional genomics



#### 4. Outcomes



#### 5. Your honours project!

### What you will walk away with:

Genome sequencing analysis and variant interpretation

Molecular biology techniques (cloning, cell culture, imaging etc)

Micro-CT and automated embryo phenotyping

**Animal handling** 

Understanding of cardiovascular genetics and functional genomics

Laboratory experience in one of the top cardiovascular genetics labs in the world

#### People involved



**Professor Sally Dunwoodie** Deputy director of the VCCRI S.Dunwoodie@victorchang.edu.au

**Click this link** to talk to us!







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