



**WISCONSIN**  
UNIVERSITY OF WISCONSIN-MADISON

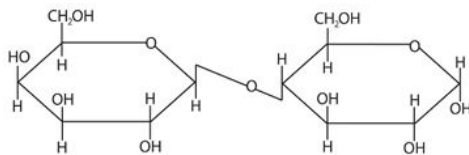
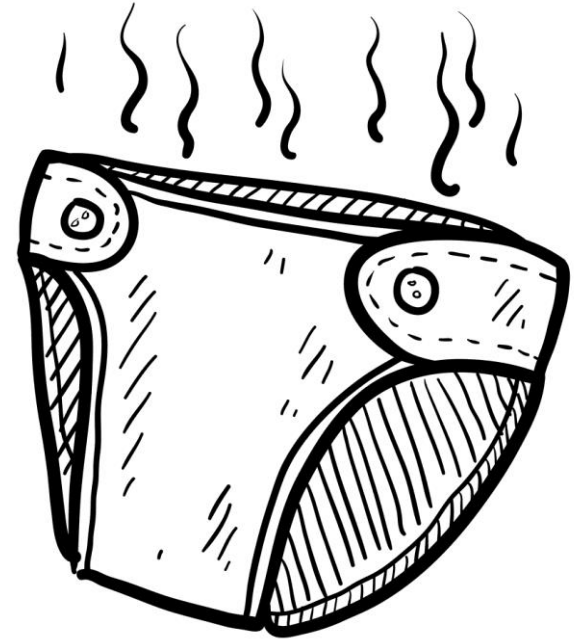


# Congenital Lactase Deficiency (LCT)



**Hayley Stoneman**

# What is congenital lactase deficiency (CLD)?



**CLD patients cannot digest lactose, resulting in diarrhea**

# LCT is associated with CLD

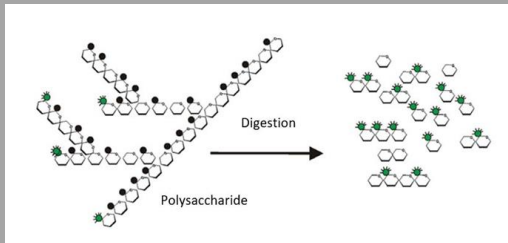
Signal  
sequence

Glycosyl hydrolase family 1

Transmembrane  
region

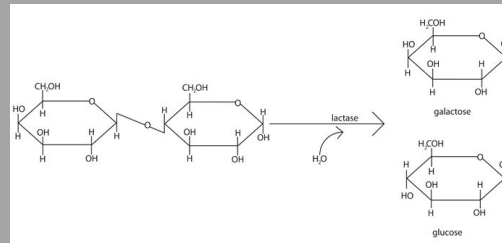


## Biological Process



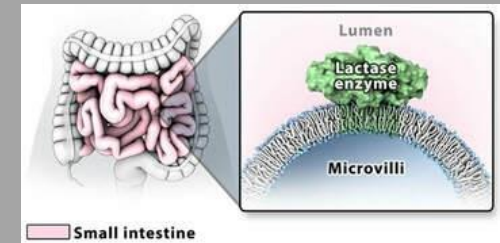
Polysaccharide  
digestion

## Molecular Function



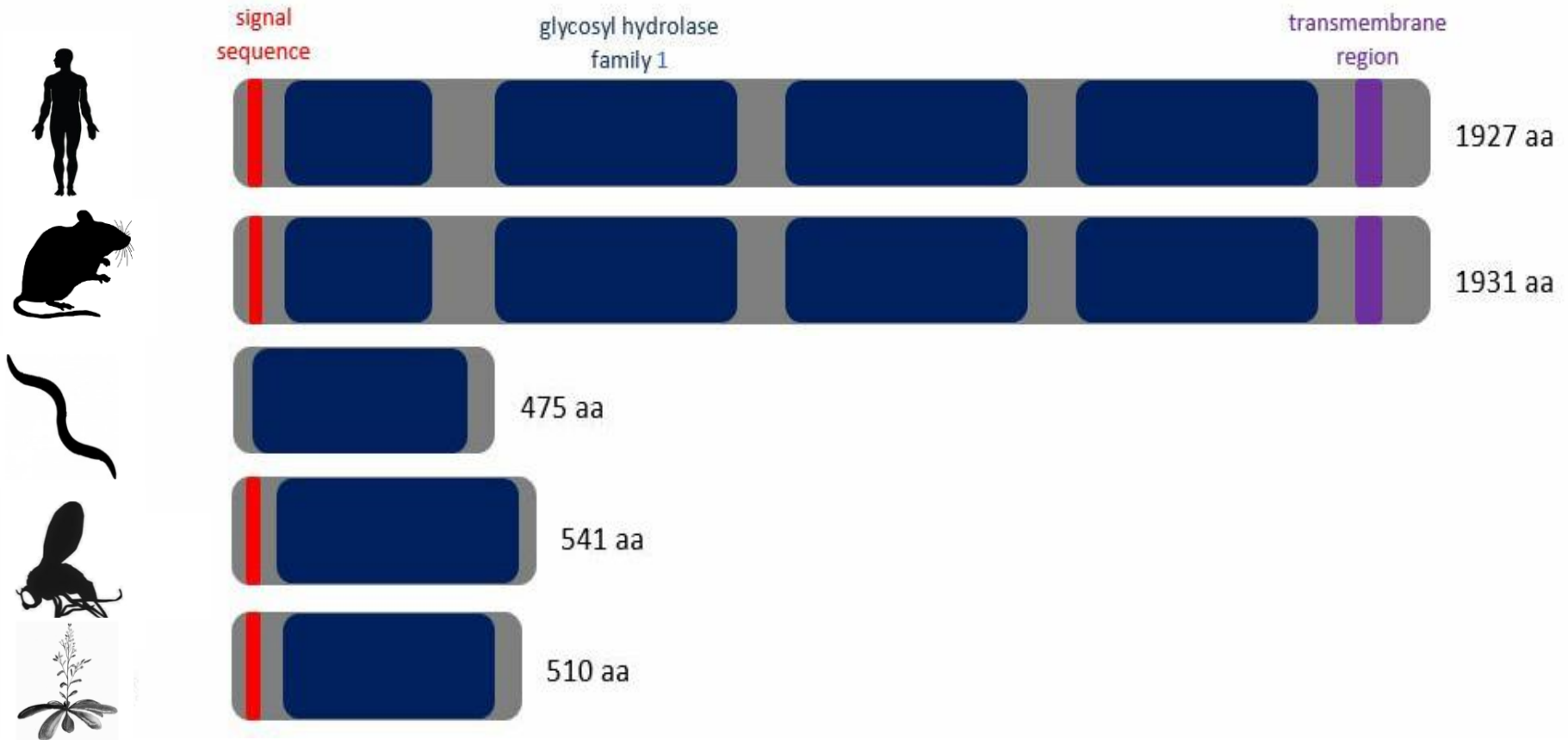
Lactase activity

## Cellular Component



Intestinal tract

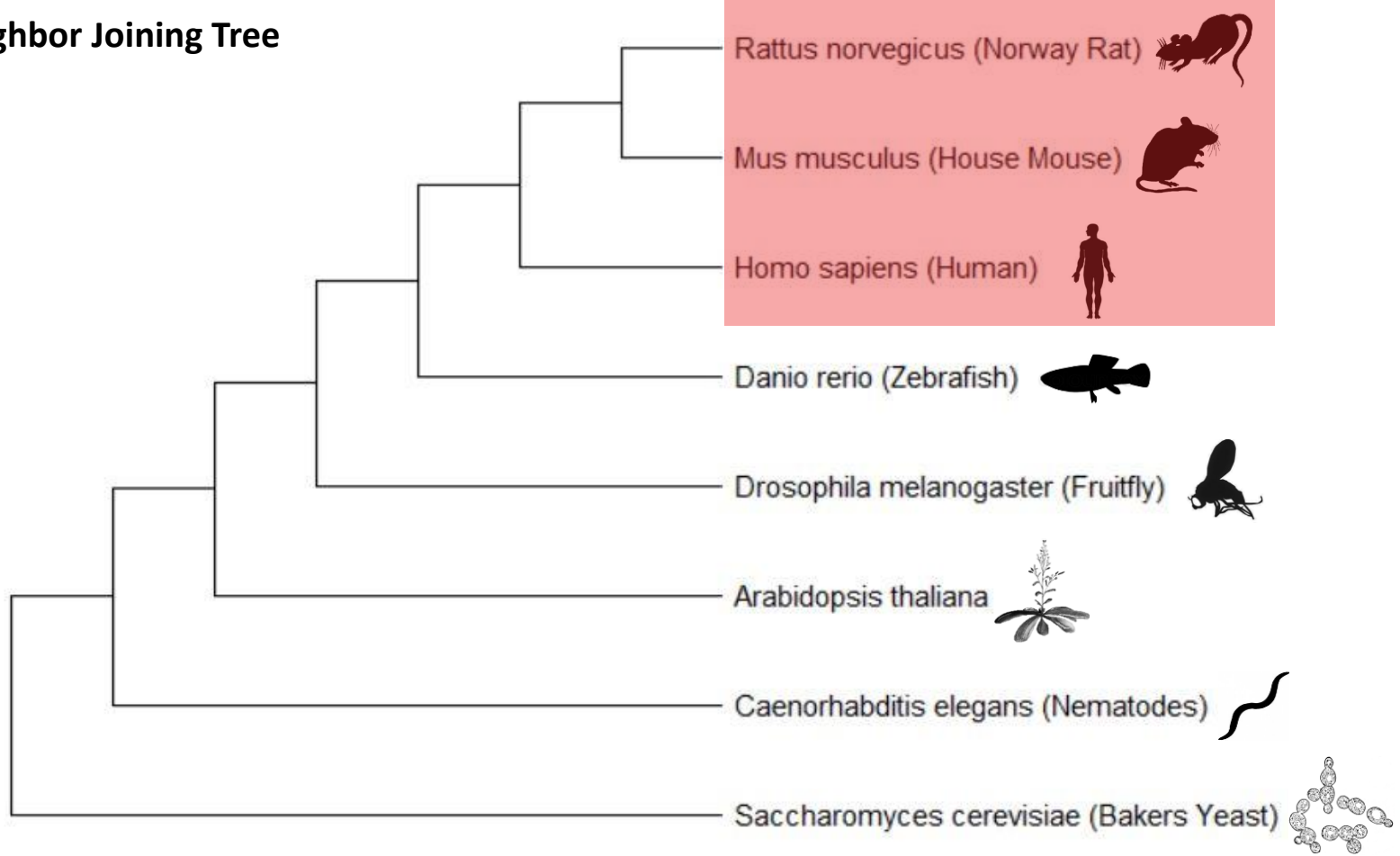
# LCT is well conserved in mammals



All homologs have glycosyl-hydrolase family 1 domains

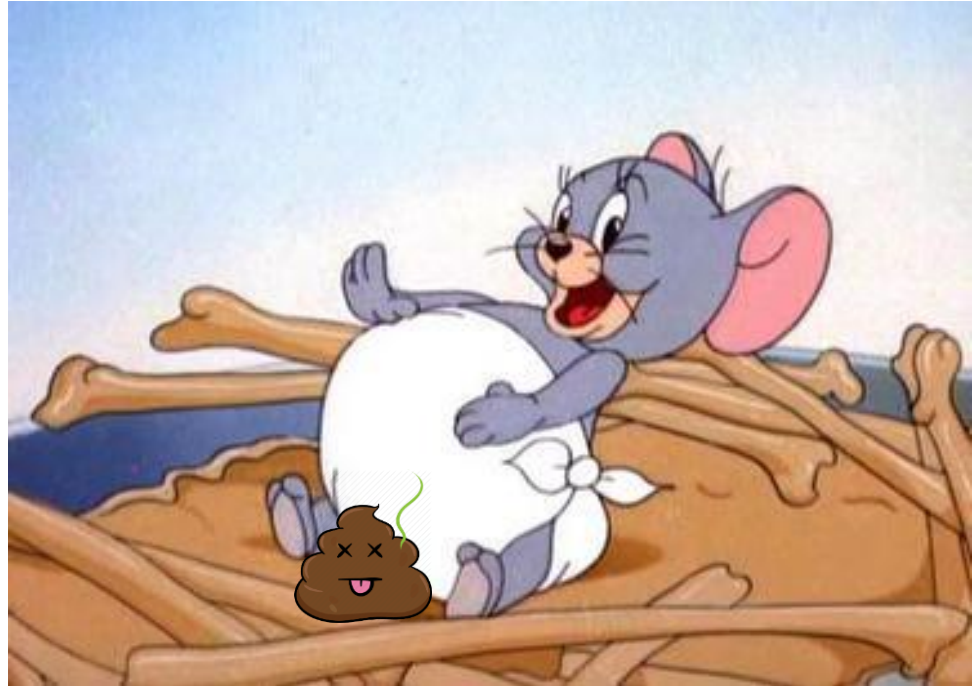
# What is **LCT** phylogeny?

Neighbor Joining Tree



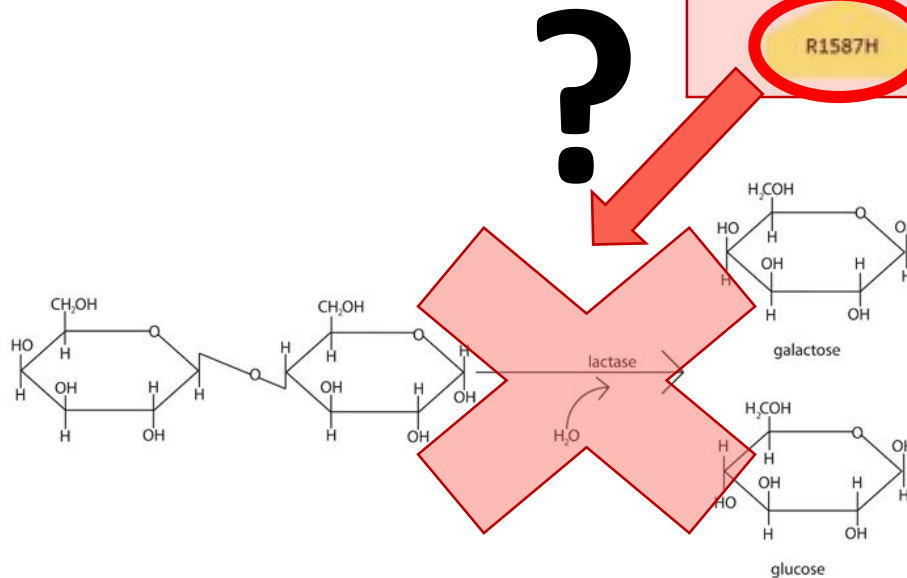
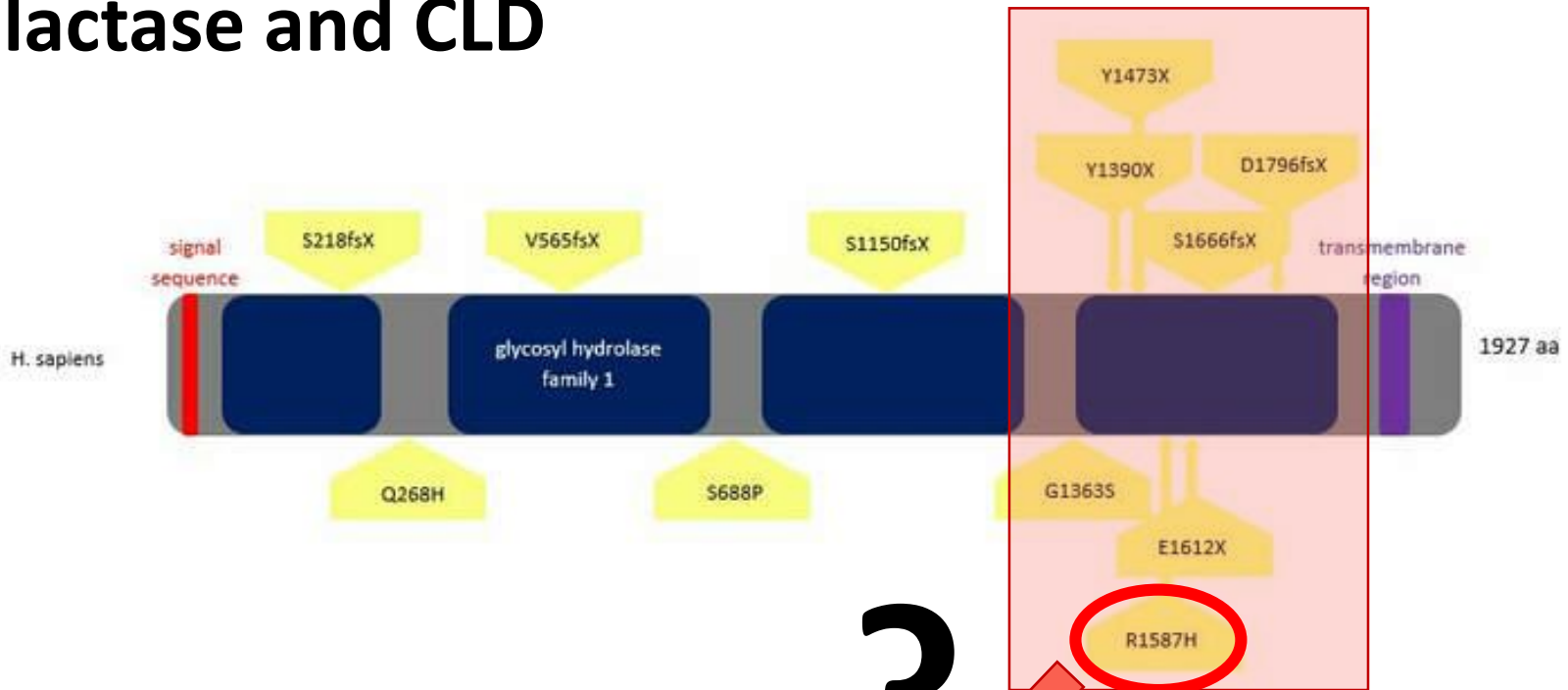
Milk-consuming animals have closest related **LCT** proteins

# What model organism should be used to study CLD?



**Mouse diet is easily manipulated and diarrhea is easily observed**

# Unknown how R1587H causes loss of functional lactase and CLD



# What is the primary goal?

To understand how a single amino acid substitution, **R1587H**, affects the structure, function, and interaction of **LCT** to determine how it leads to CLD

## Aim 1

Understand the evolutionary history of LCT and conservation of the C-terminus and R1587

## Aim 2

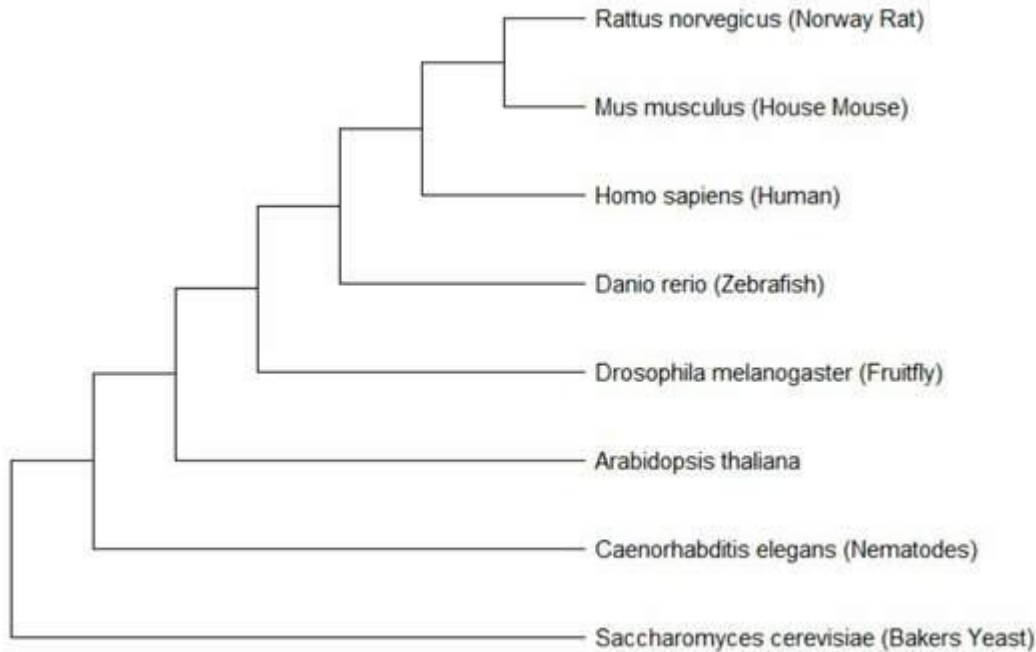
Determine differentially expressed genes and their functions in WT and R1587H mutant mice

## Aim 3

Experimentally determine protein-protein interactions of WT and R1587H mutant mice LCT



# Aim 1: Understand evolutionary history of LCT and conservation of R1587 and the C-terminus



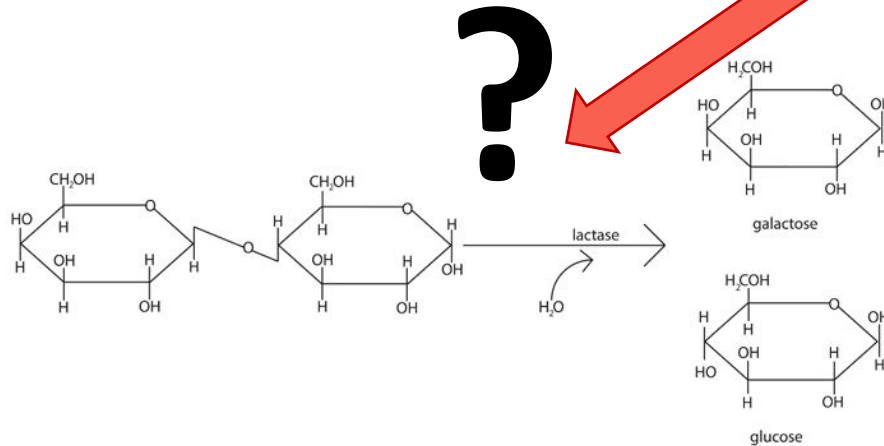
**Approach:** Use amino acid sequence to build phylogenetic trees and observe conserved regions

# Aim 1: Understand evolutionary history of **LCT** and conservation of R1587 and the C-terminus

What

Why

Hypothesis



**Rationale:** Evolutionary history can elucidate the importance of R1587 in functional lactase among milk consuming organisms

# Aim 1: Understand evolutionary history of **LCT** and conservation of R1587 and the C-terminus

What

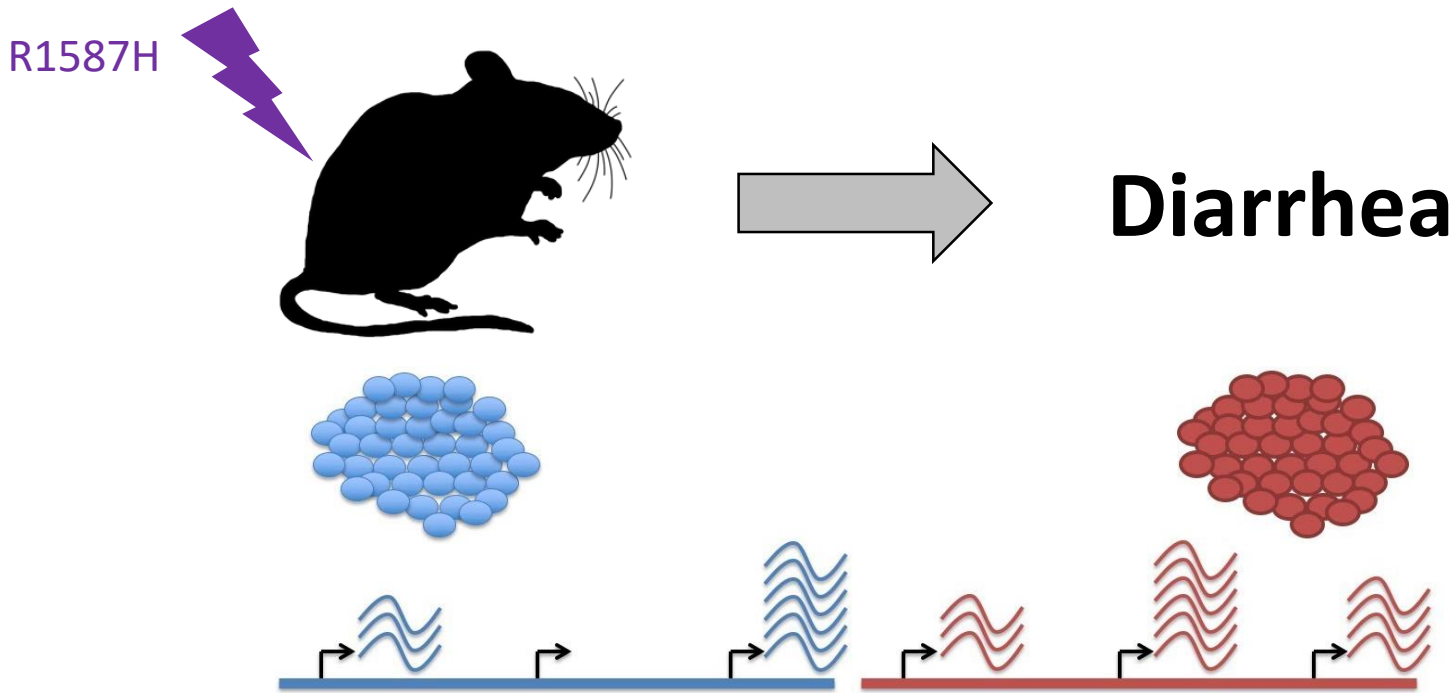
Why

Hypothesis



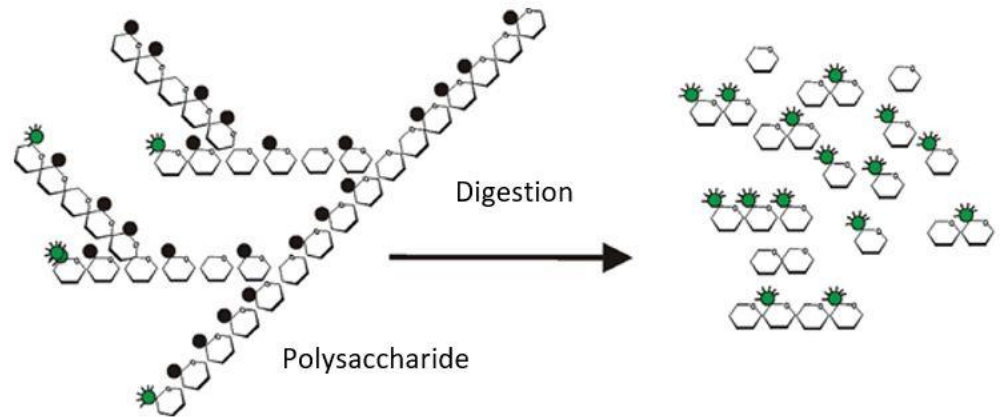
**Hypothesis:** R1587 will be conserved among milk consuming organisms

# Aim 2: Determine differentially expressed genes and their functions in R1587H and WT mice



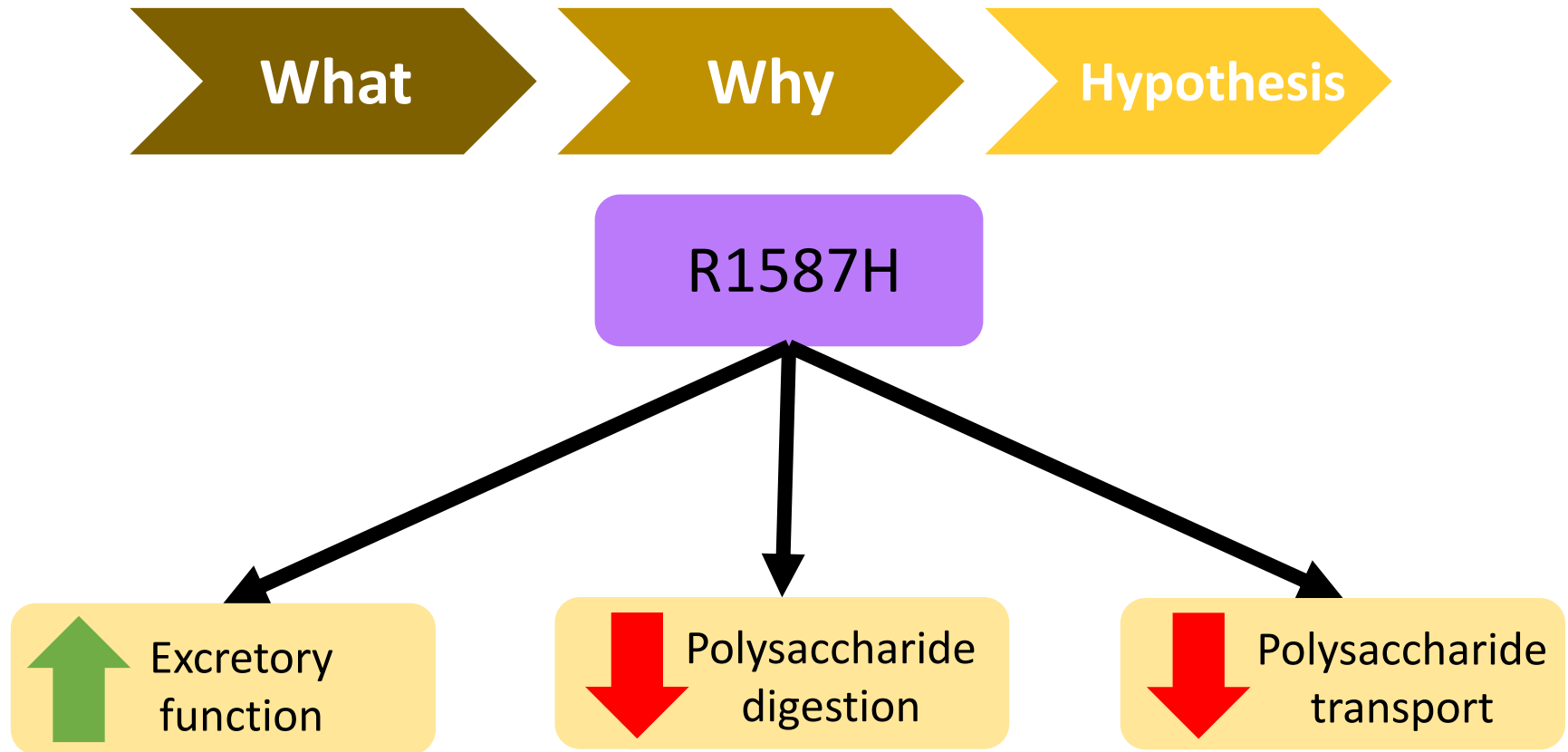
**Approach:** Create mutant mouse line and confirm CLD phenotype, then perform RNA-seq to identify gene expression

# Aim 2: Determine differentially expressed genes and their functions in R1587H and WT mice



**Rationale:** Determining differentially expressed genes will elucidate interrupted biological processes

## Aim 2: Determine differentially expressed genes and their functions in R1587H and WT mice



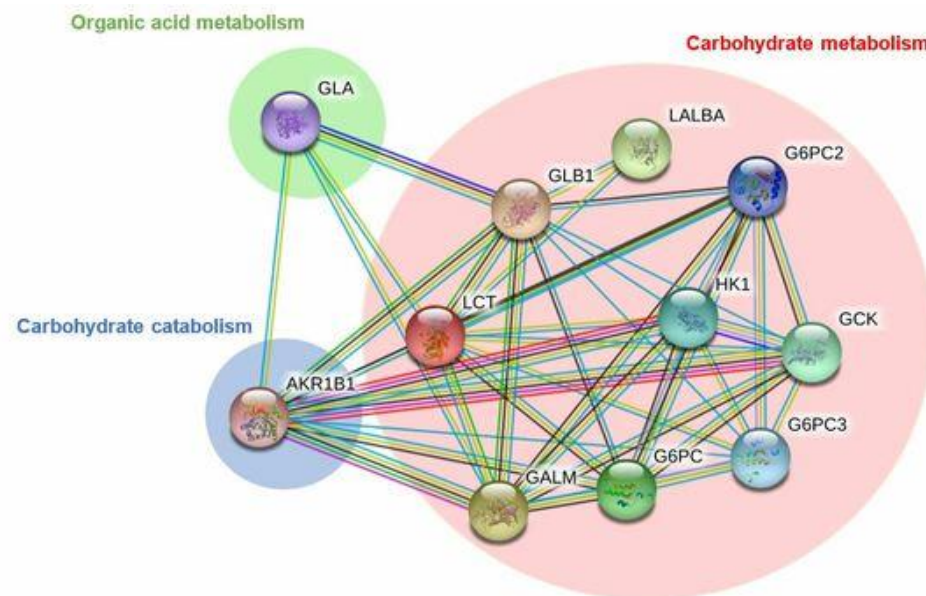
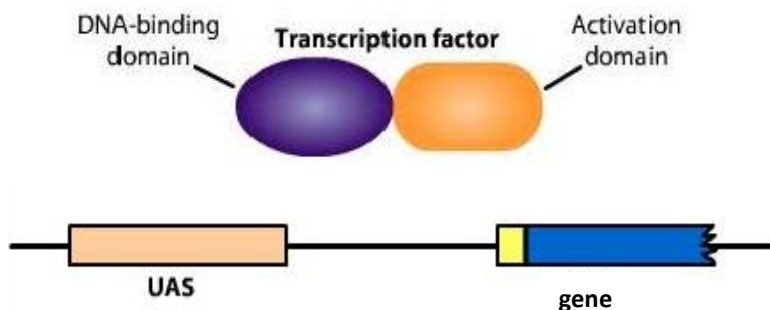
**Hypothesis:** R1587H mice will have upregulated genes related to excretory function downregulated genes in polysaccharide digestion/transport

# Aim 3: Experimentally determine protein-protein interactions of WT and R1587H LCT

What

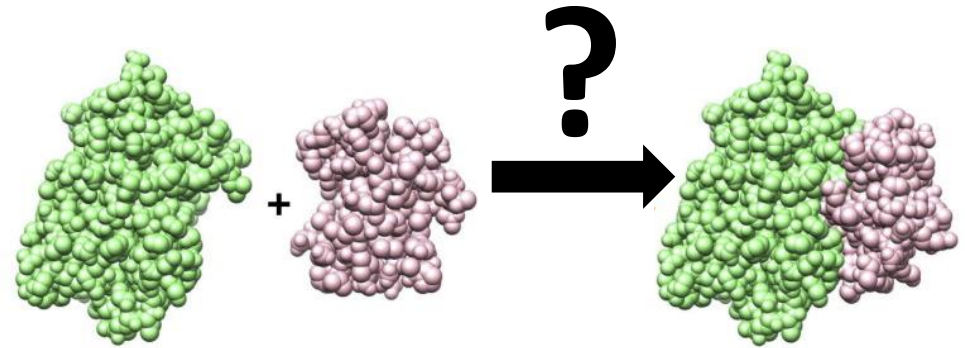
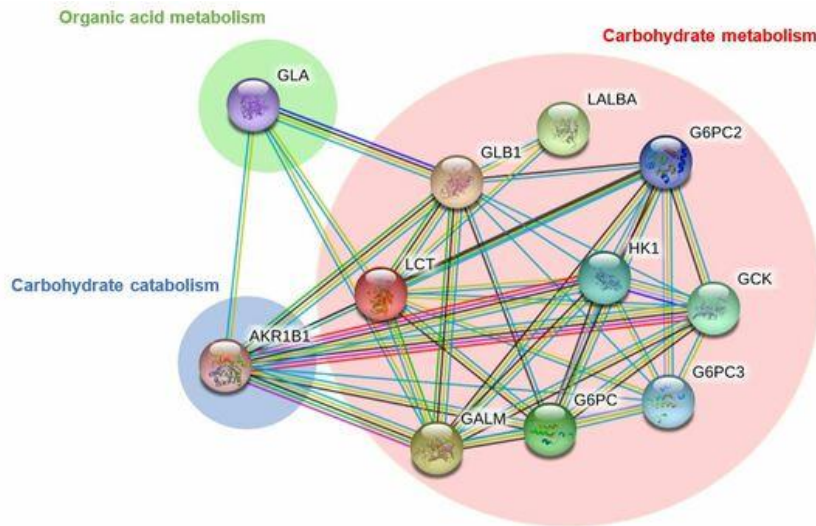
Why

Hypothesis



**Approach:** Use mammalian two-hybrid system to identify protein-protein interactions and sort according to GO terms

# Aim 3: Experimentally determine protein-protein interactions of WT and R1587H LCT



**Rationale:** Create first experimentally determined interaction network for LCT and determine which protein-protein interactions inhibited by R1587H

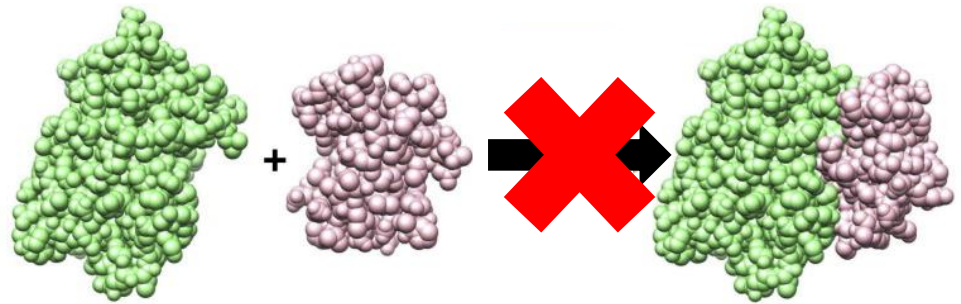
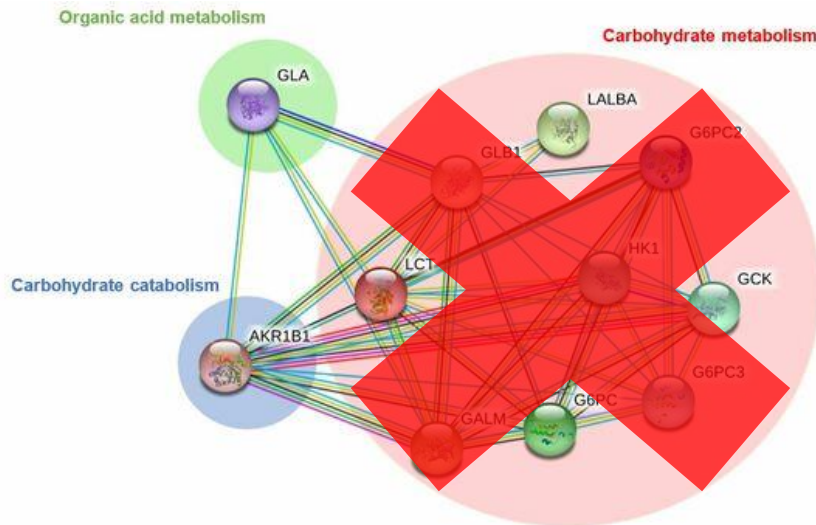


# Aim 3: Experimentally determine protein-protein interactions of WT and R1587H LCT

What

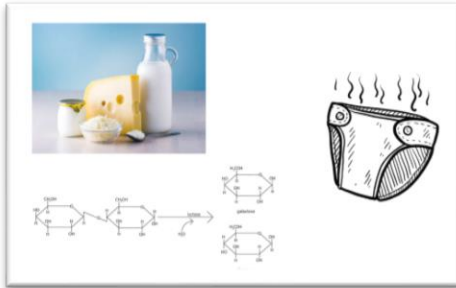
Why

Hypothesis

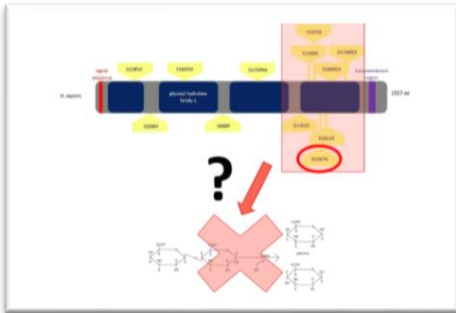


**Hypothesis:** Interaction network will involve carbohydrate metabolism proteins and R1587H will interrupt many of these interactions

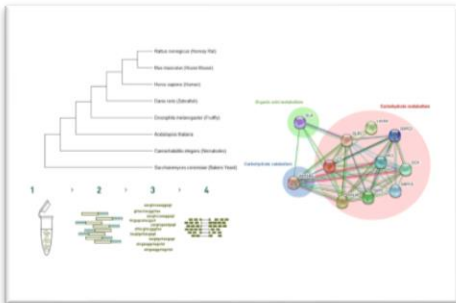
# Summary



CLD is a disease caused by lack of lactase (**LCT**) functional enzyme resulting in inability to digest lactose leading to diarrhea



It is unknown how missense mutations in the C-terminal glycosyl-hydrolase family domain lead to CLD



Phylogenomic, transcriptomic, and proteomic techniques will be used to study the functional contribution of **R1587H** to **LCT**

# Future Directions



**Develop  
treatment for CLD**

# Questions?



# References

Genetics Home Reference (2019). Lactose Intolerance. U.S. National Library of Medicine. Retrieved from <https://ghr.nlm.nih.gov/condition/lactose-intolerance#genes>

Diekmann, L et al. (2015, March). Congenital lactose intolerance is triggered by severe mutations on both alleles of the lactase gene. BMC Gastroenterology. Retrieved from [bmcgastroenterol.biomedcentral.com/articles/10.1186/s12876-015-0261-y](http://bmcgastroenterol.biomedcentral.com/articles/10.1186/s12876-015-0261-y)

Kuokkanen, M. et. al. (2006, February). Mutations in the Translated Region of the Lactase Gene (LCT) Underlie Congenital Lactase Deficiency. American Journal on Human Genetics. Retrieved from <https://www.sciencedirect.com/science/article/pii/S0002929707623647>

Behrendt, M. et al. (2009, June). Impaired Trafficking and Subcellular Localization of a Mutant Lactase Associated with Congenital Lactase Deficiency. Gastroenterology. Retrieved from <https://www.sciencedirect.com/science/article/pii/S0016508509001425>

Luo, Y. et al. (1997, February). Mammalian two-hybrid system: a complementary approach to the yeast two-hybrid system. Biotechniques. Retrieved from <https://www.ncbi.nlm.nih.gov/pubmed/9043710>

**\*all images hyperlinked**