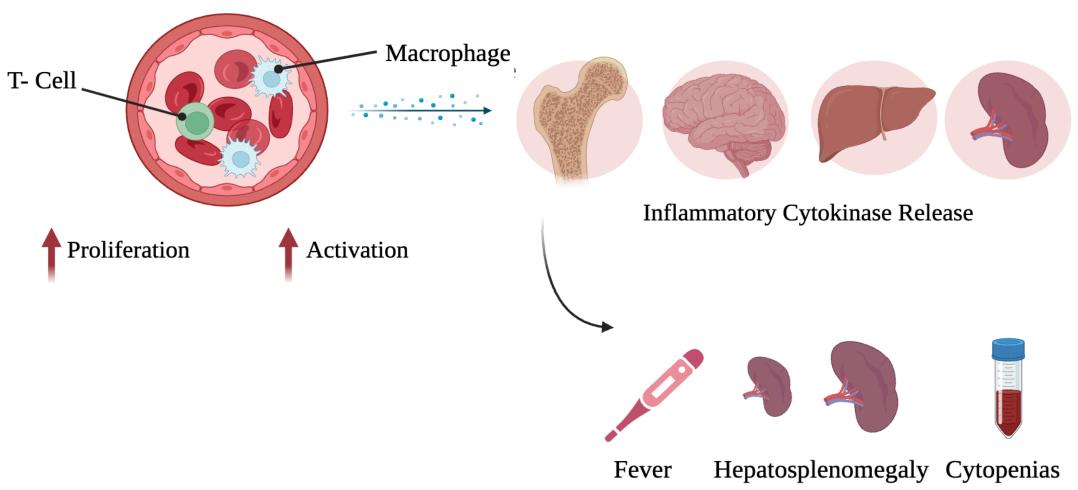
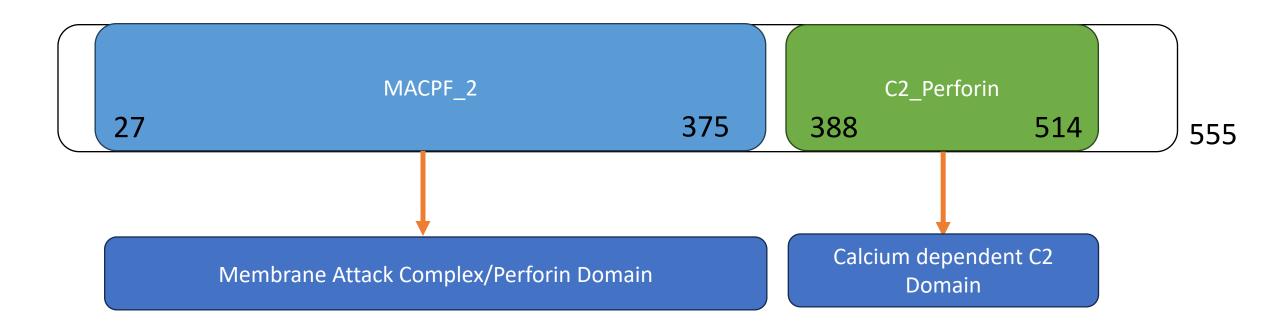


Overactivation of T-cells and macrophages results in symptoms of HLH



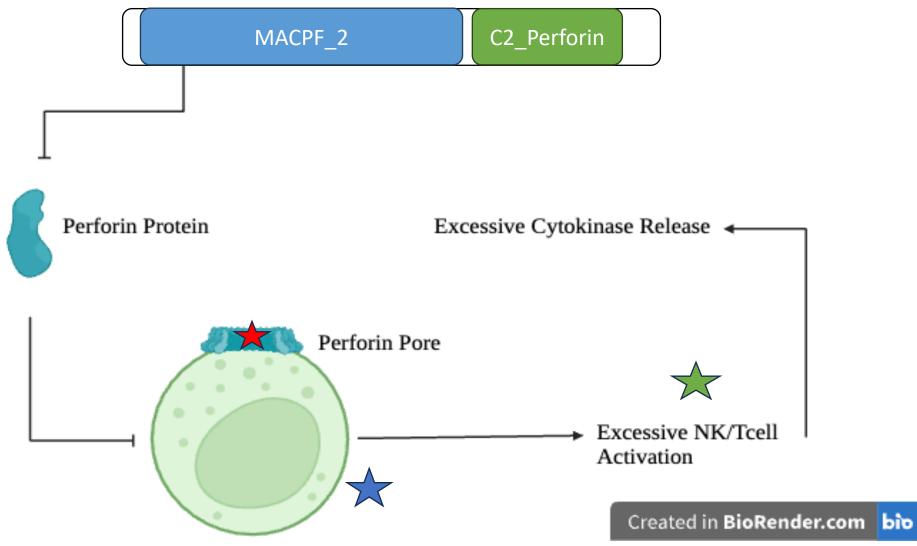
Familial HLH is caused by loss-of-function mutation in PRF1



InterPro: Classification of Protein Families (Paysan-Lafosse et al. 2022)

Loss of PRF1 decreases perforin pore formations and associated cytotoxic responses

Two Pathogenic Alleles Required

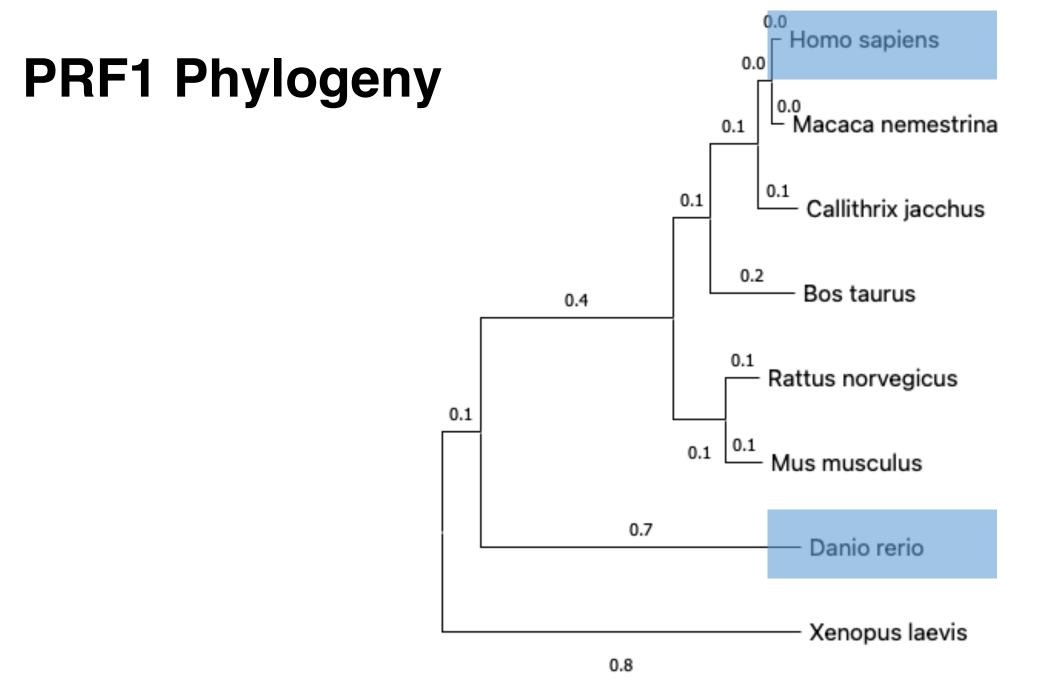


Osinka et al. 2014; InterPro: Classification of Protein Families (Paysan-Lafosse et al. 2022)

Protein Domains Across Species

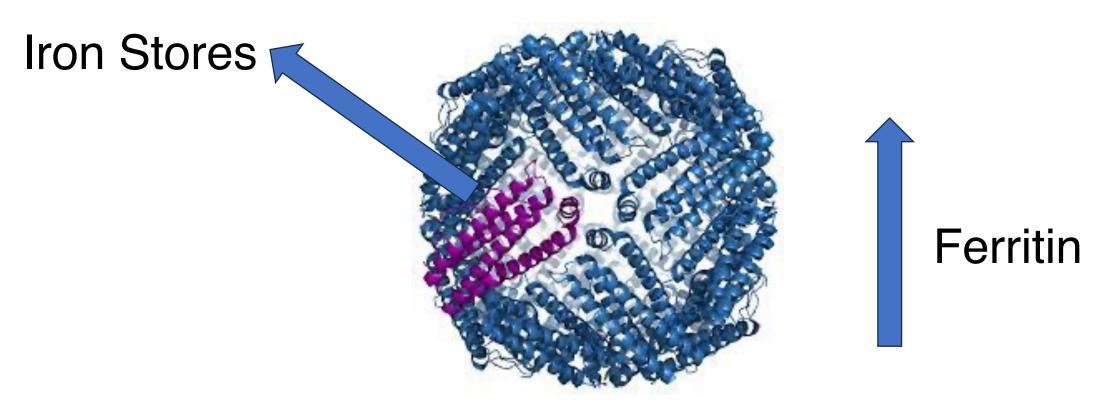
Homo Sapiens

27 MACPF_2 375 388 C2_Perforin 514



MEGA11: Molecular Evolutionary Genetics Analysis version 11 (Tamura, Stecher, and Kumar 2021)

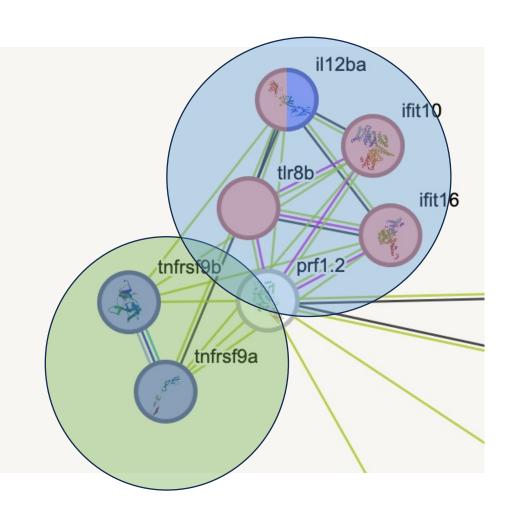
The relationship between hyperferritinemia and PRF1 mutation is unclear

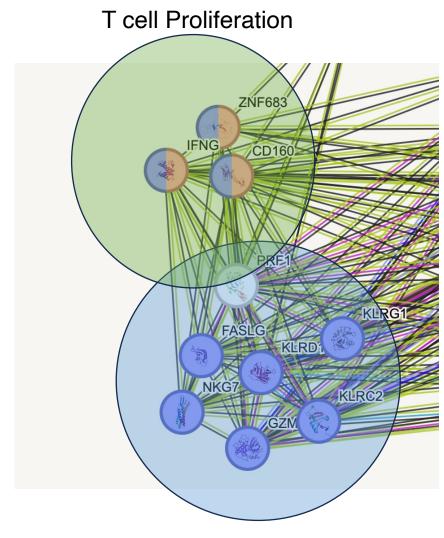


Hypothesis: iron homeostasis in fHLH will result from varying expression of ferritin encoding genes [FTL FTH], and subsequent protein concentrations



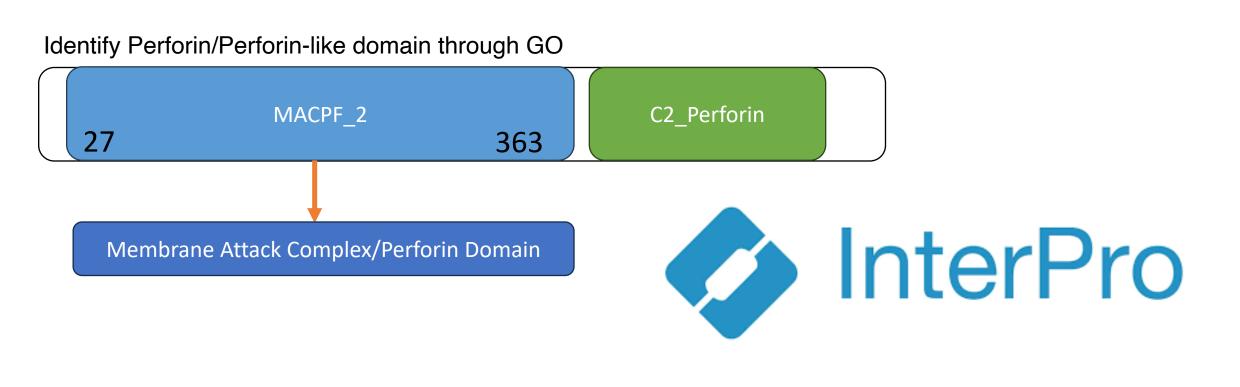
PRF1 Interactions





Innate Immune Response

Aim 1 Identifying Protein Domain Essential for Pore Formation in Danio Rerio



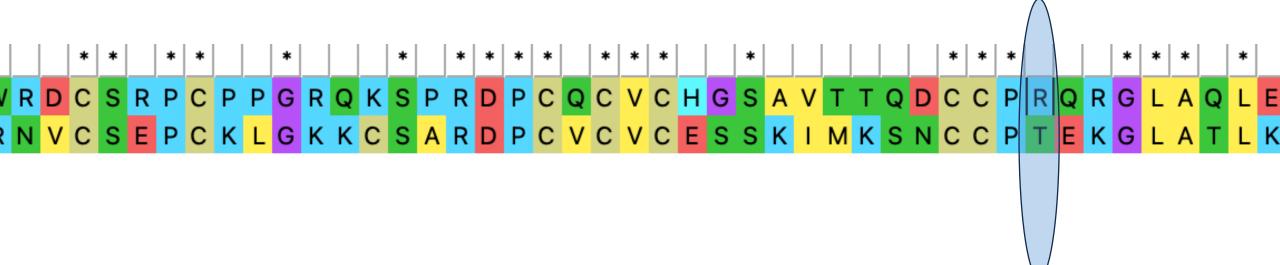
NCBI Protein Blast

GO Analysis

Confirmation By Screening

InterPro: Classification of Protein Families (Paysan-Lafosse et al. 2022)

Aim 1 Identifying Protein Domain Essential for Pore Formation in Danio Rerio



Multiple Sequence Alignment

GO Analysis

Aim 1 Identifying Protein Domain Essential for Pore Formation in Danio Rerio

MACPF_2

C2_Perforin

NCBI Protein Blast

GO Analysis

Confirmation By Screening

(Murtha et al., 2003); Created in Biorender

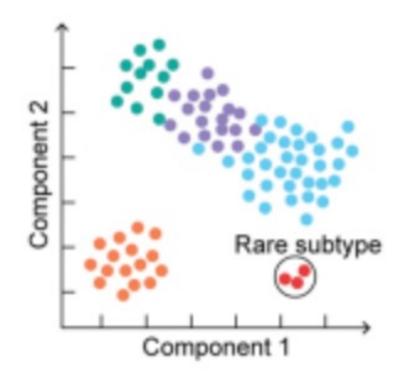
Aim 2 Identifying Gene Expression in PRF1 and Iron Homeostasis Genes

RNA Sequencing

Cluster Analysis

Aim 2 Identifying Gene Expression in PRF1 in Iron Homeostasis Genes

Identify Gene Clusters relating to Cytotoxicity and Iron Homeostasis



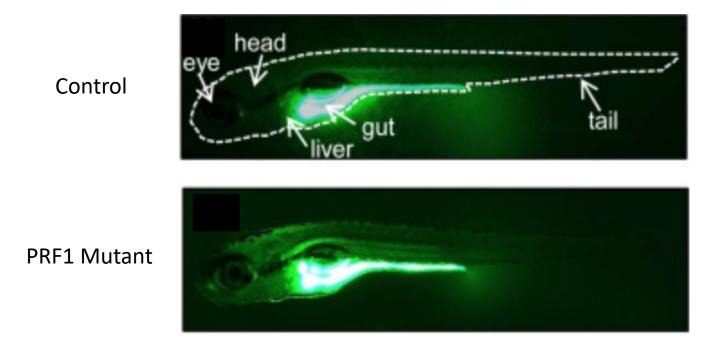
(Hwang et al., 2018)

RNA Sequencing

Cluster Analysis

Aim 2

Identifying Gene Interactions in PRF1 and Iron Homeostasis Genes



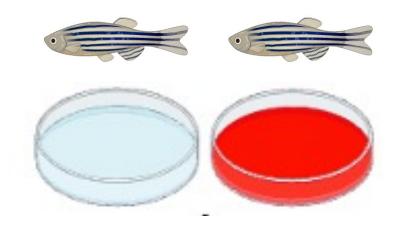
Screen for Ferritin in vivo with fluorescence assay

(Ayaat et al., 2020)

RNA Sequencing

Cluster Analysis

Aim 3 Analysis of proteome via metabolic labeling of control/KO tissues



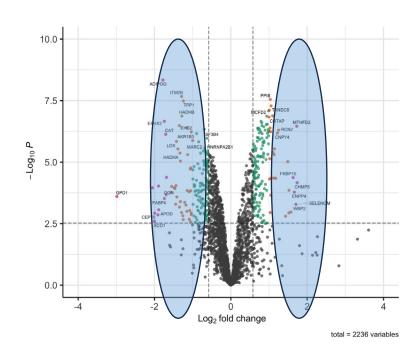
(Hwang et al., 2018)

Isotopic Labeling

Analysis

Aim 3 Analysis of proteome via metabolic labeling of KO tissues

Identify Ferritin in increased expression regions

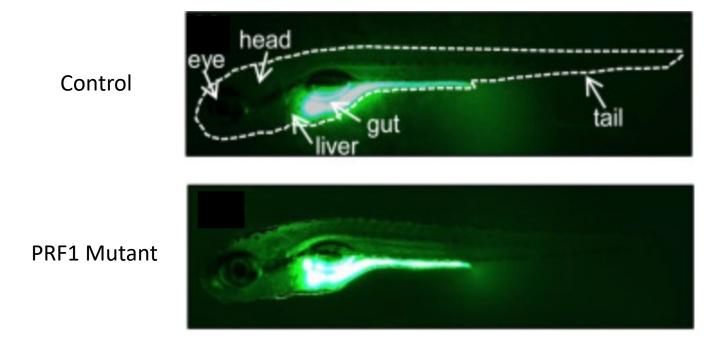


(Hwang et al., 2018)

Isotopic Labeling

Analysis

Aim 3 Analysis of proteome via metabolic labeling of KO tissues



Screen for Ferritin in vivo with fluorescence assay

(Ayaat et al., 2020)

Isotopic Labeling

Cluster Analysis

Summary

- Today, we identified what fHLH is and how PRF1 mutation causes damage to the immune response. The relationship between this condition and high iron homeostasis is unknown.
- We explored Danio Rerio, a model organism for PRF1 mutation, and how it can be used to elucidate this relationship
- We proposed multiple genomics/proteomics approaches to exploring the relationship between these pathways.

Future Directions





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