

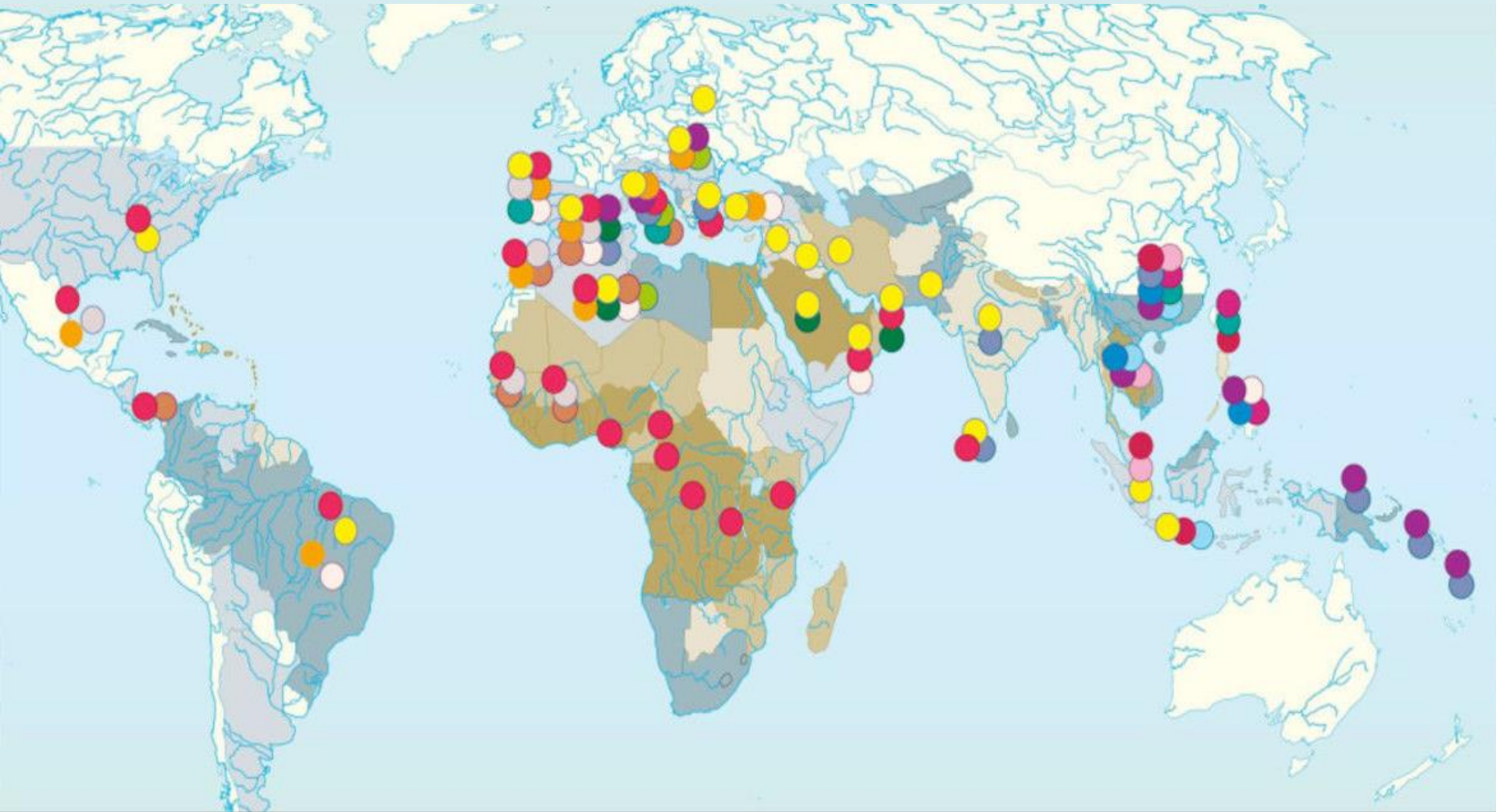
G6PD Deficiency-Induced Hemolysis and Malaria

Diane Xue

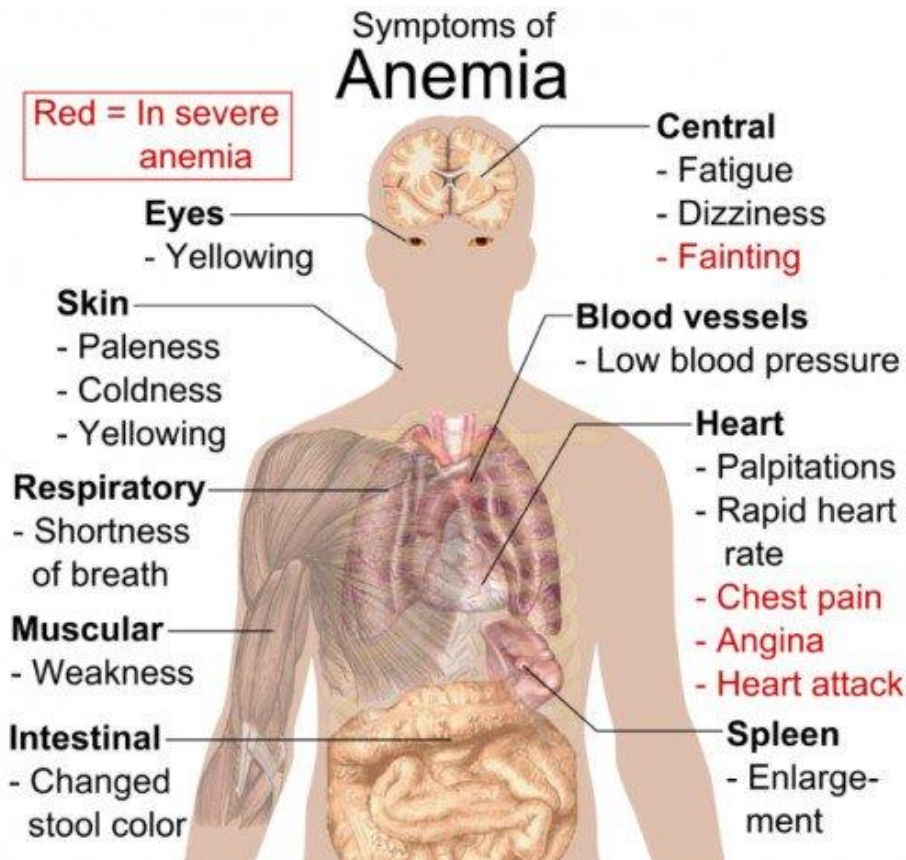


WISCONSIN
UNIVERSITY OF WISCONSIN-MADISON

G6PD Deficiency is the most common Enzymopathy in the World



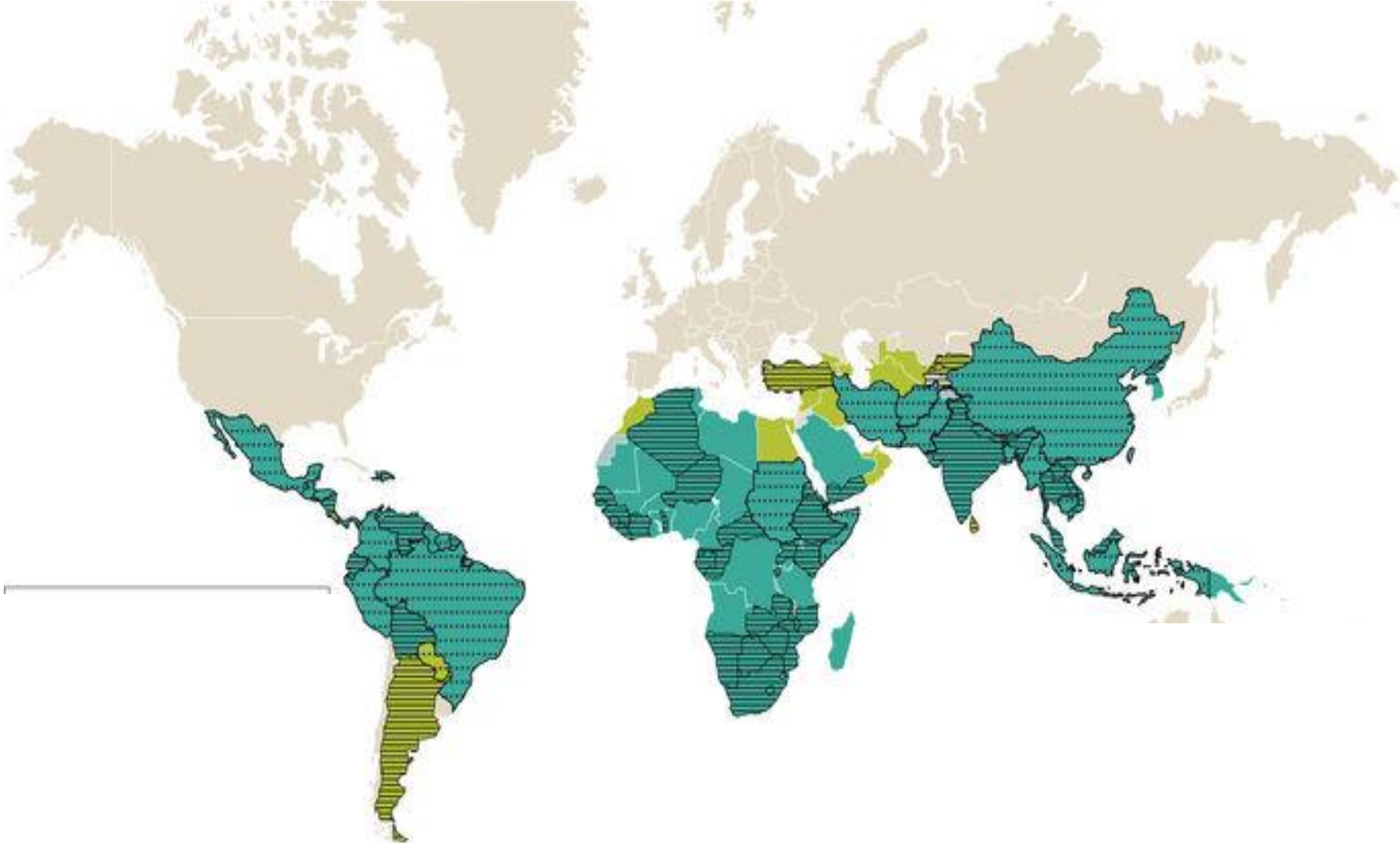
G6PD Deficiency Phenotypes



Hemolytic Anemia
11/100 deaths

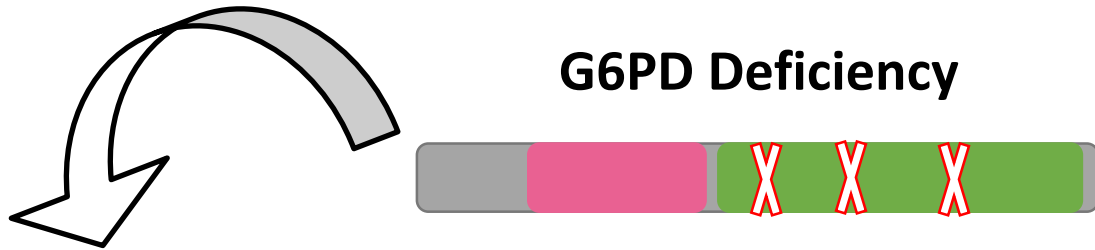
Neonatal Jaundice
7.3/1000

G6PD Deficiency's Indirect Consequence



Malaria = 720,000 deaths per year

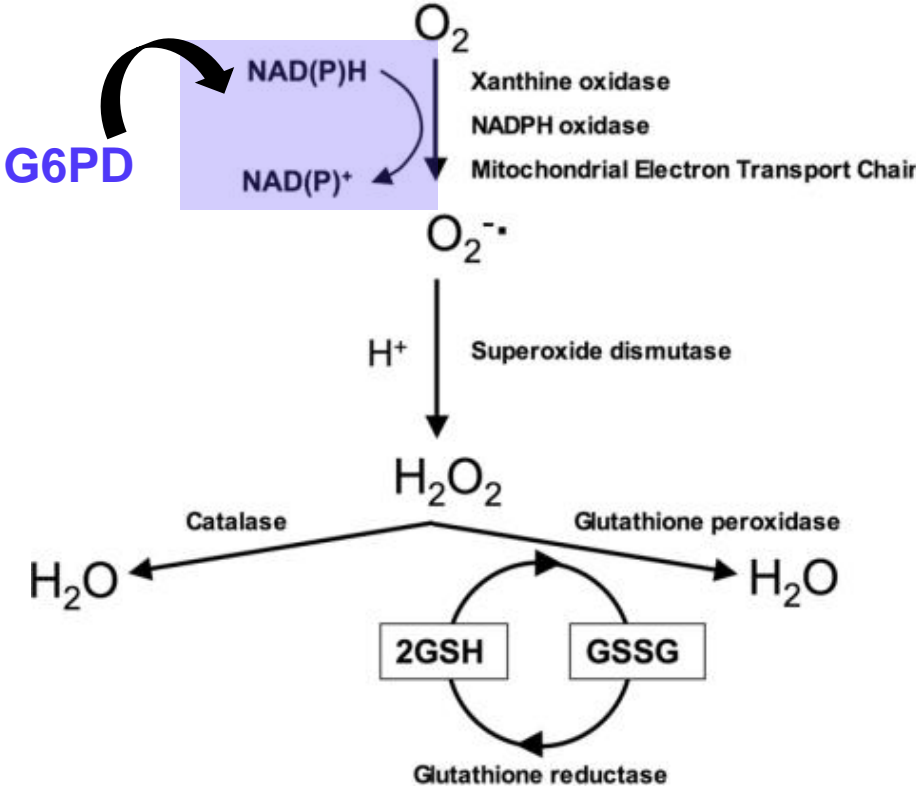
How are G6PD deficiency, Hemolysis, and Malaria related?



Hemolysis



How does G6PD cause Susceptibility for Hemolysis?



No Build-up of Reactive Oxygen Species

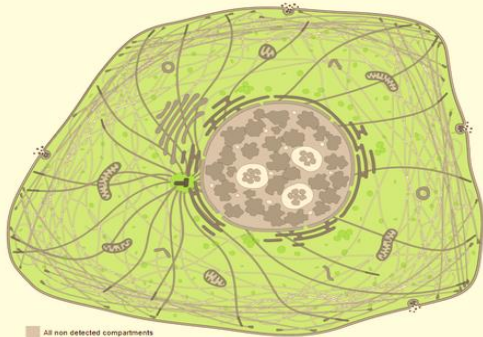
What is the function of normal G6PD?



NAD binding domain

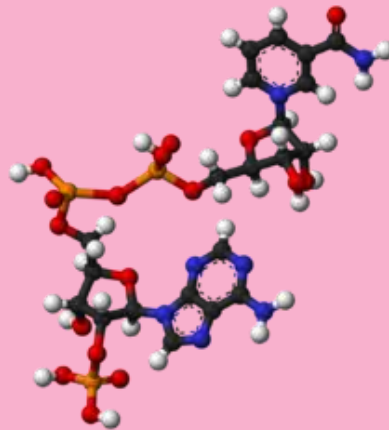
C-terminal domain

Cellular Component



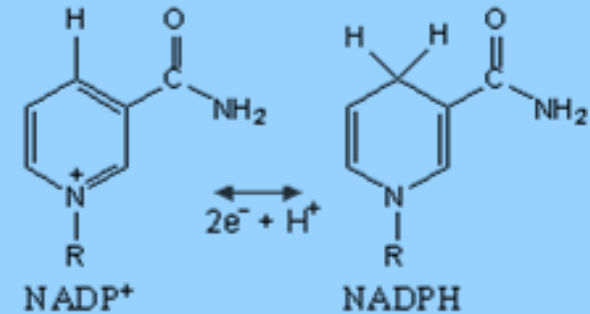
Cytoplasm

Molecular Functions



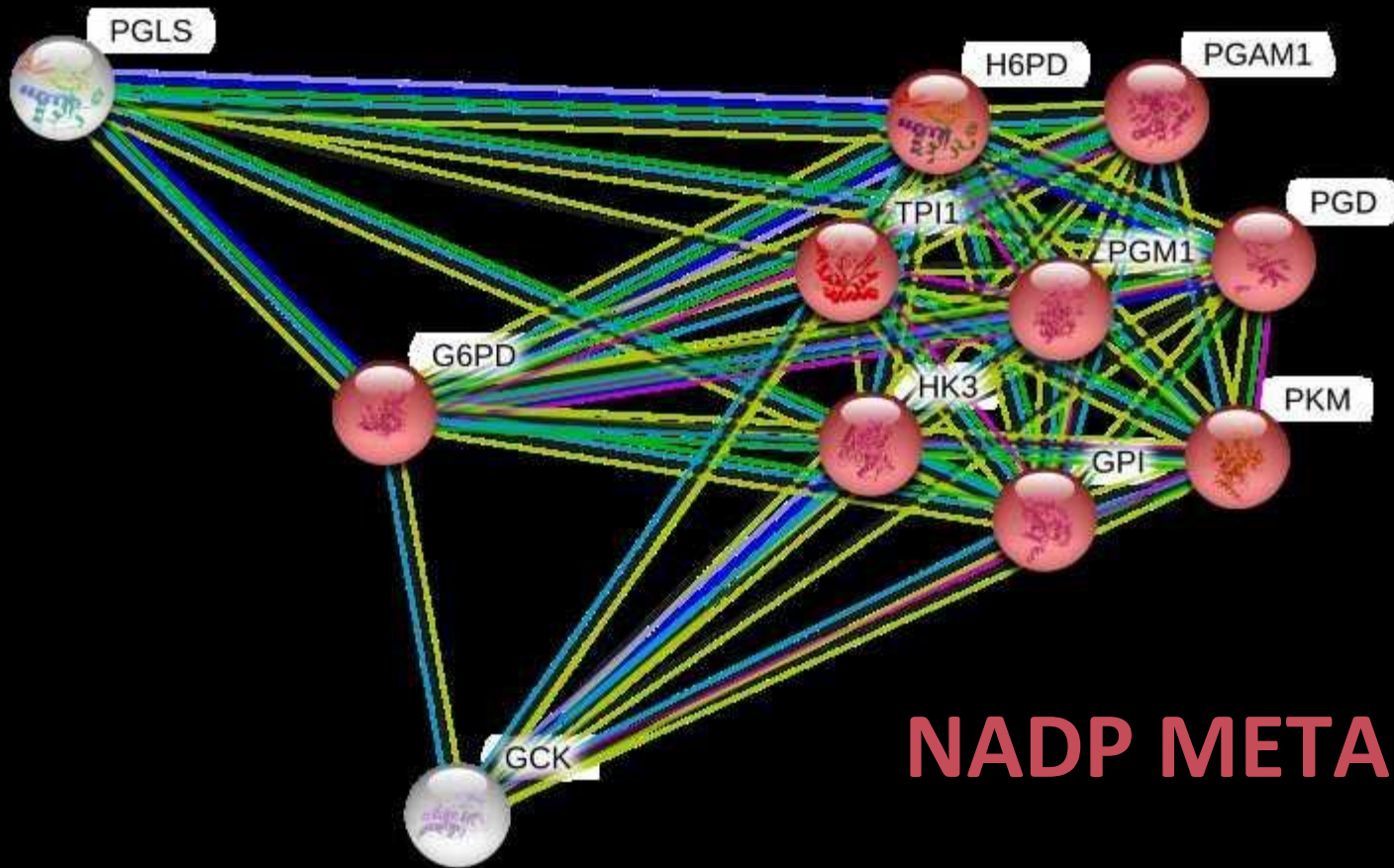
NADP Binding

Biological Process



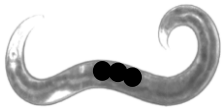
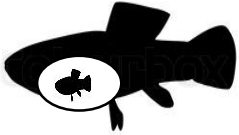
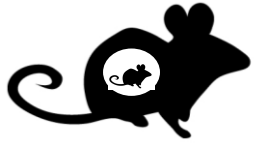
NADP Metabolism

What Proteins Interact with G6PD?



NADP METABOLISM

How well conserved is *G6PD*?



Can G6PD be upregulated?

Proc. Natl. Acad. Sci. USA
Vol. 82, pp. 1465–1469, March 1985
Genetics

Tissue-specific levels of human glucose-6-phosphate dehydrogenase correlate with methylation of specific sites at the 3' end of the gene

(DNA methylation/housekeeping genes/transcriptional regulation)

GIORGIO BATTISTUZZI*[†], MICHELE D'URSO*[‡], DANIELA TONIOLO[‡], G. M. PERSICO[‡], AND LUCIO LUZZATTO*

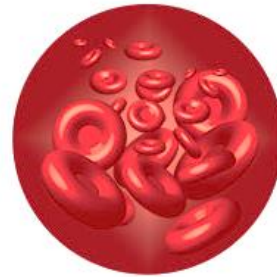
*Department of Haematology, Royal Postgraduate Medical School, Ducane Road, London W12 OHS, England; and [‡]International Institute of Genetics and Biophysics, Consiglio Nazionale delle Ricerche, Via Marconi 10, 80125 Naples, Italy

Communicated by Paul A. Marks, October 24, 1984

**** THIS STUDY WAS ONLY DONE ON MALES**



Goal: How does *G6PD* Methylation affect *G6PD* levels in Pregnant Females?



Normal Amount of
Red Blood Cells



Anemic Amount of
Red Blood Cells



How does *G6PD* Methylation affect G6PD levels (NADP Production) in Pregnant Females?

Aim 1:

Knockout conserved methylation sites of G6PD to identify methylation sites associated with normal levels of NADPH production in pregnant females.

Bisulfite DNA-Seq

ClustalOmega

CRISPR

Fluorescent Spot Test

Aim 2:

Identify a small molecule that affects methylation of G6PD and can be taken with Primaquine in pregnant females without hemolysis

Bisulfite DNA-Seq

Chemical Screen

Fluorescent Spot Test

Bisulfite DNA-Seq

Aim 3

Determine differences in protein interaction complexes required for NADP metabolism between methylated and unmethylated G6PD deficient pregnant zebrafish

CRISPR

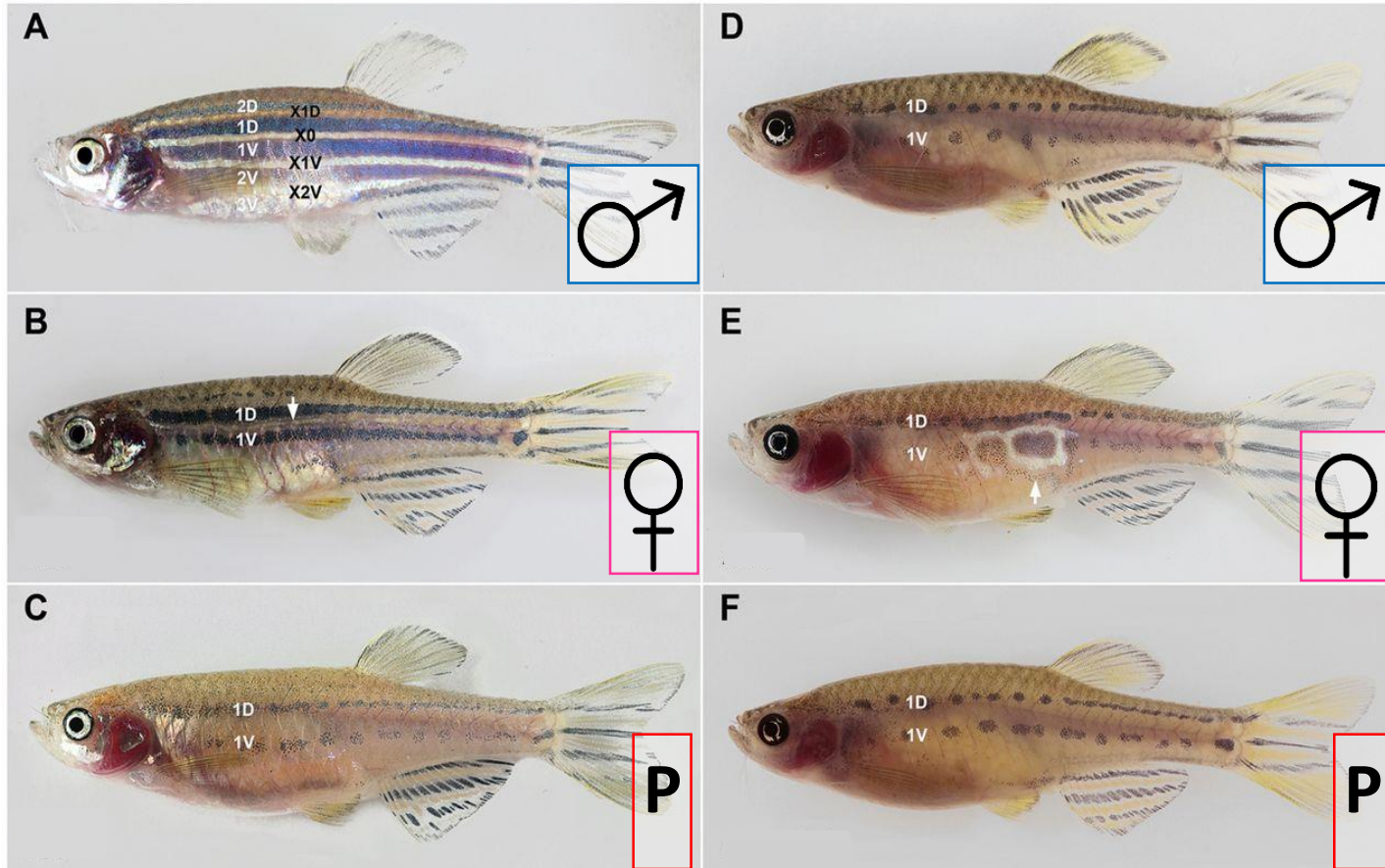
TAP-MS

Fluorescent Spot Test

What Model Organism Will I Use?

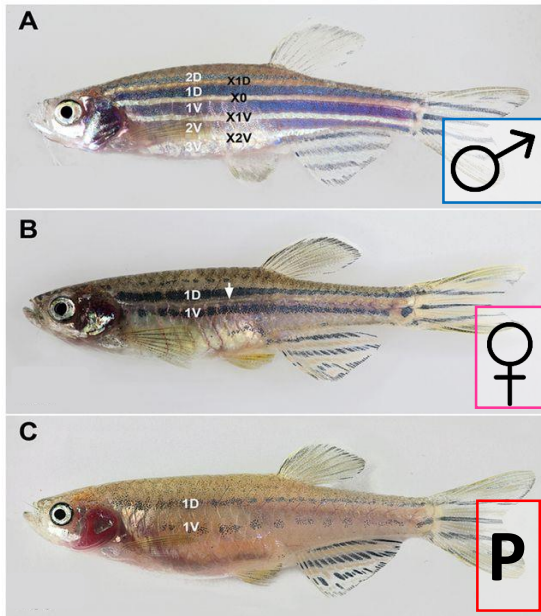
WT (Normal NADPH)

G6PDd (Low NADPH)

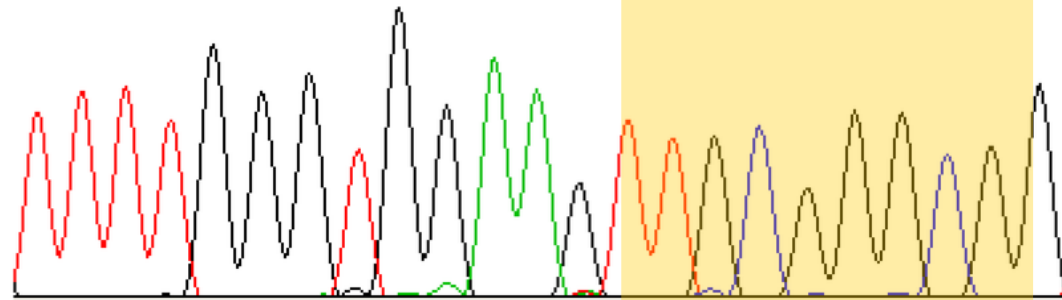


Aim 1: Identify conserved methylation sites in G6PD normal pregnant zebrafish using Bisulfite Sequencing

WT (Normal NADPH)



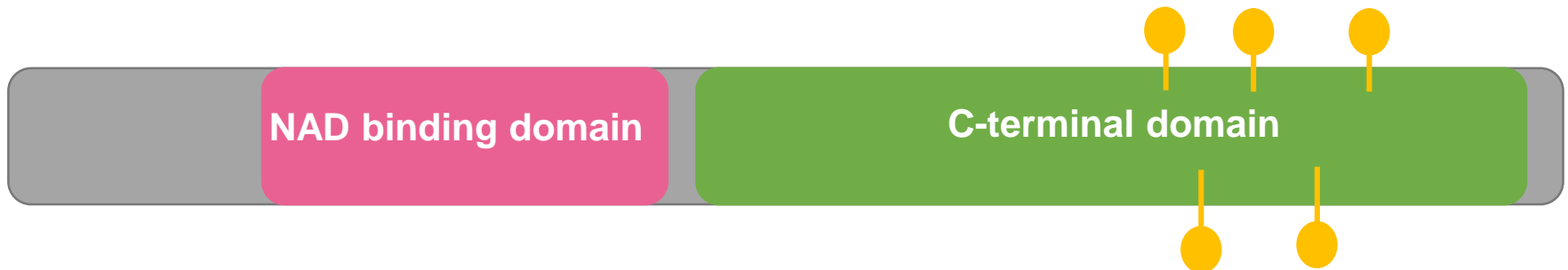
C C C C G G G C G G A A G C T G C G G G C G G
T T T T G G G T G G A A G T T G C G G G C G G



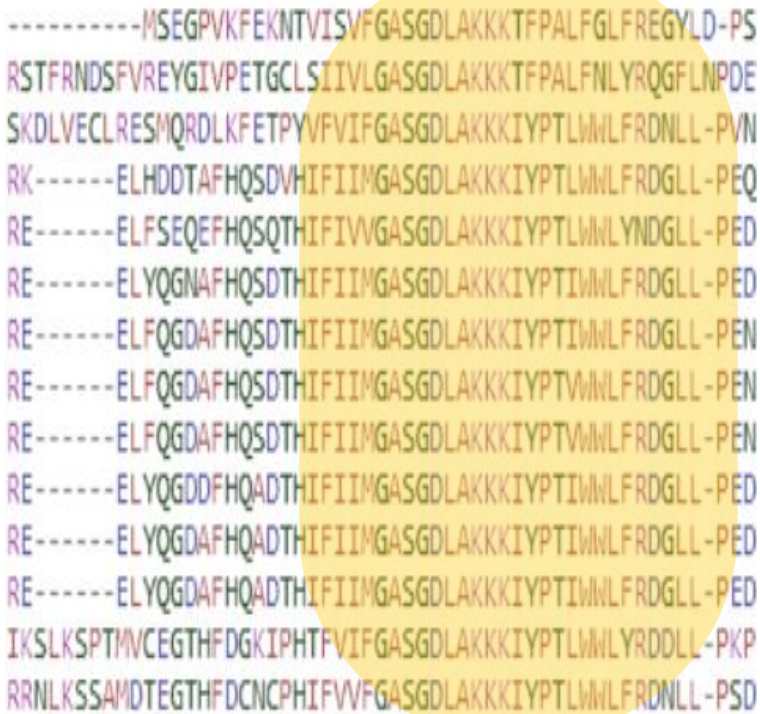
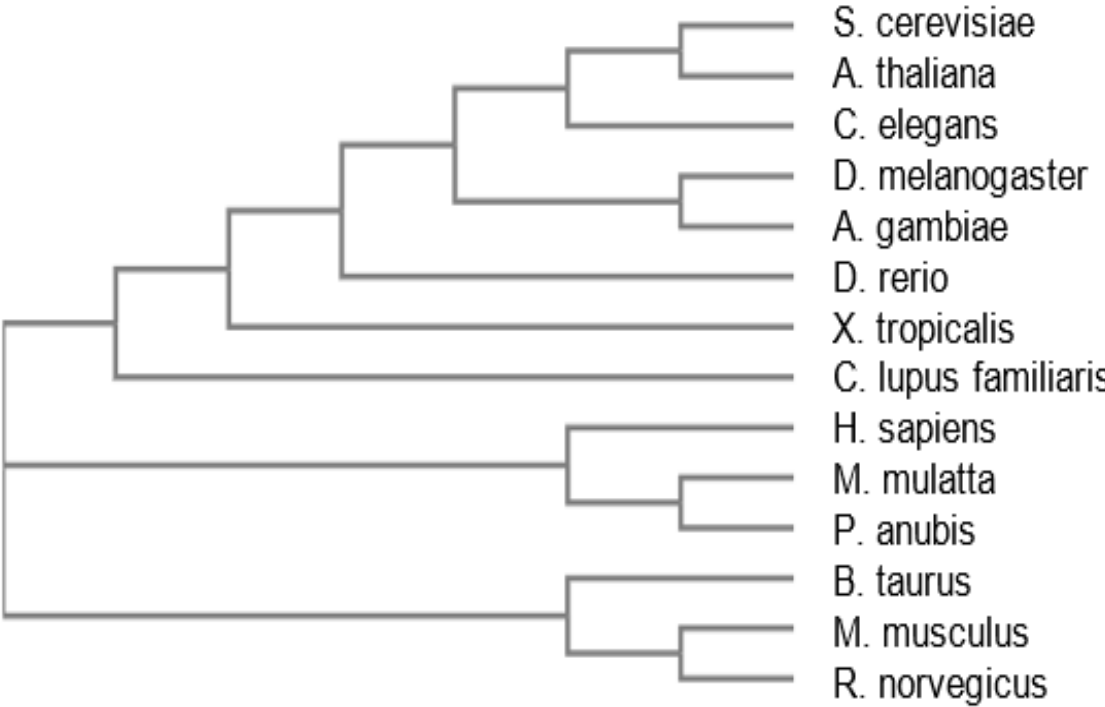
Unmethylated C

Methylated C

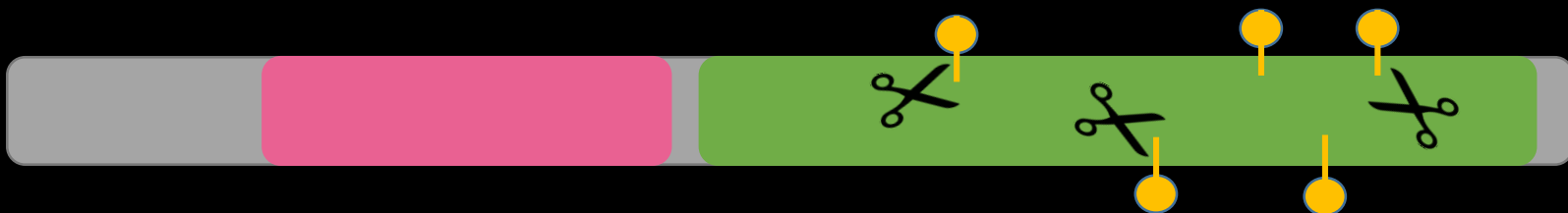
C Cytosine not in CpG site C G CpG site



Aim 1: Identify conserved methylation sites in G6PD normal pregnant zebrafish using Phylogenomics and ClustalOmega

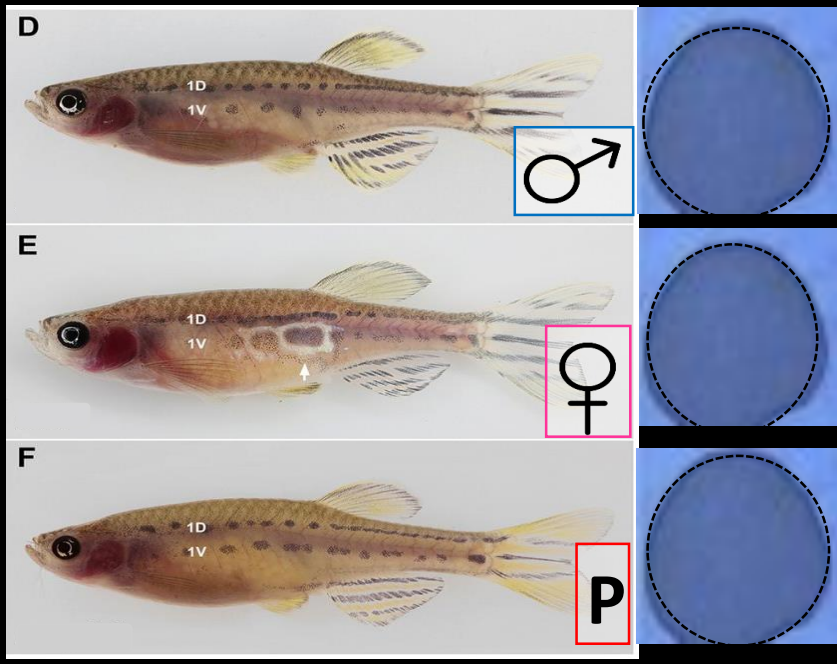
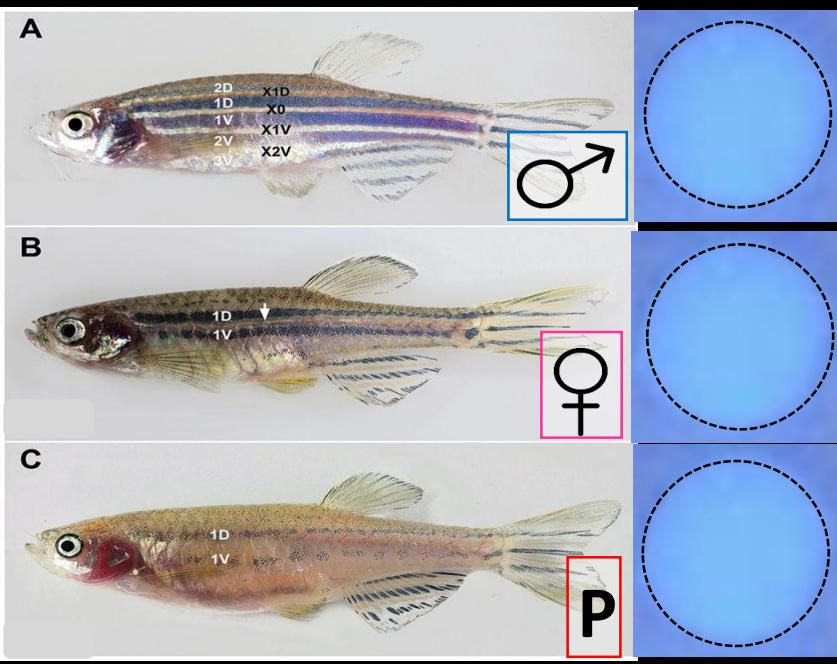


Aim 1: Identify conserved methylation sites in G6PD normal pregnant zebrafish using CRISPR/Cas9 and Fluorescent Spot Test



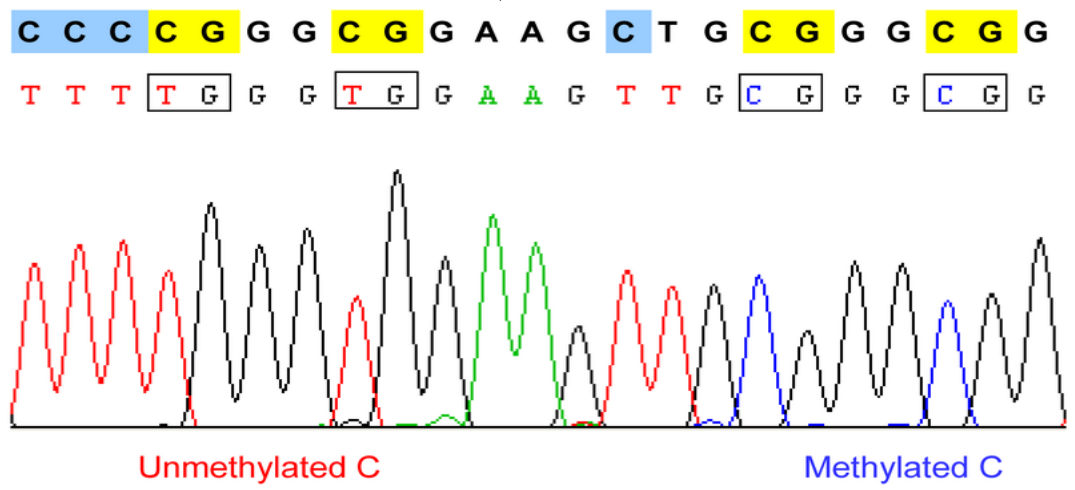
WT (Normal NADPH)

G6PDd (Low NADPH)



NADPH is naturally fluorescent and represents G6PD level

Aim 2: Identify a small molecule that affects G6PD methylation pattern of Pregnant Zebrafish

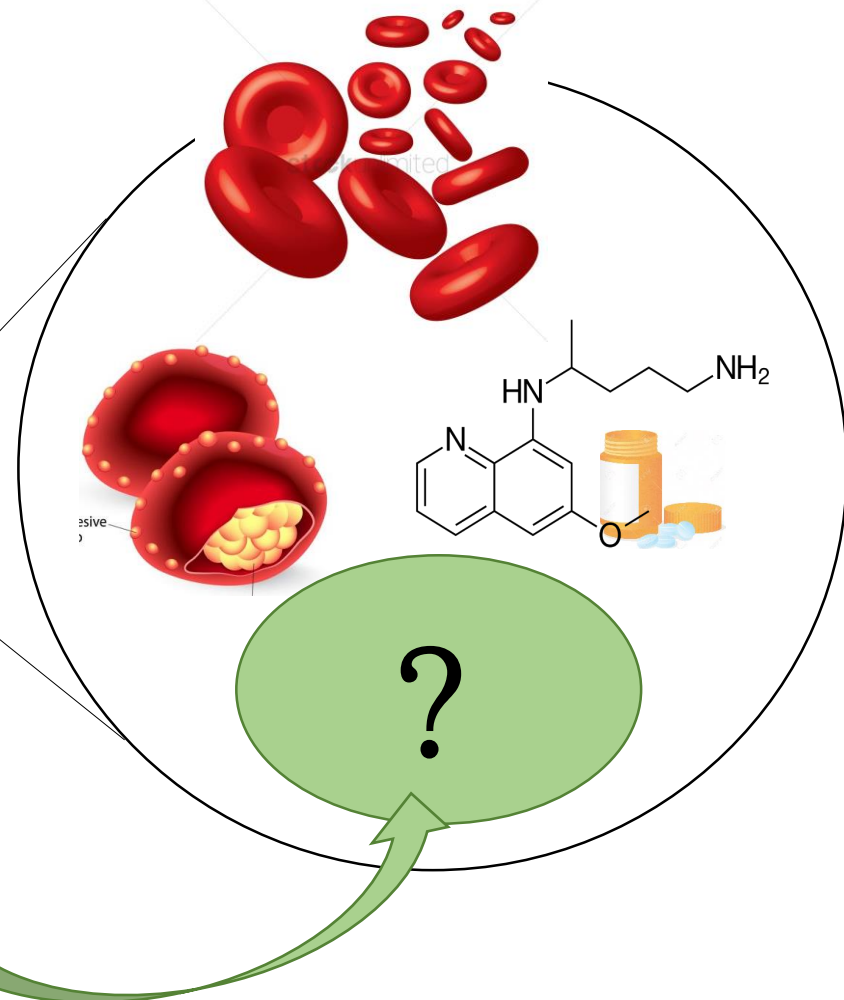


Aim 2: Identify a **small molecule** that affects methylation, destroys malaria, and can be taken with Primaquine without hemolysis

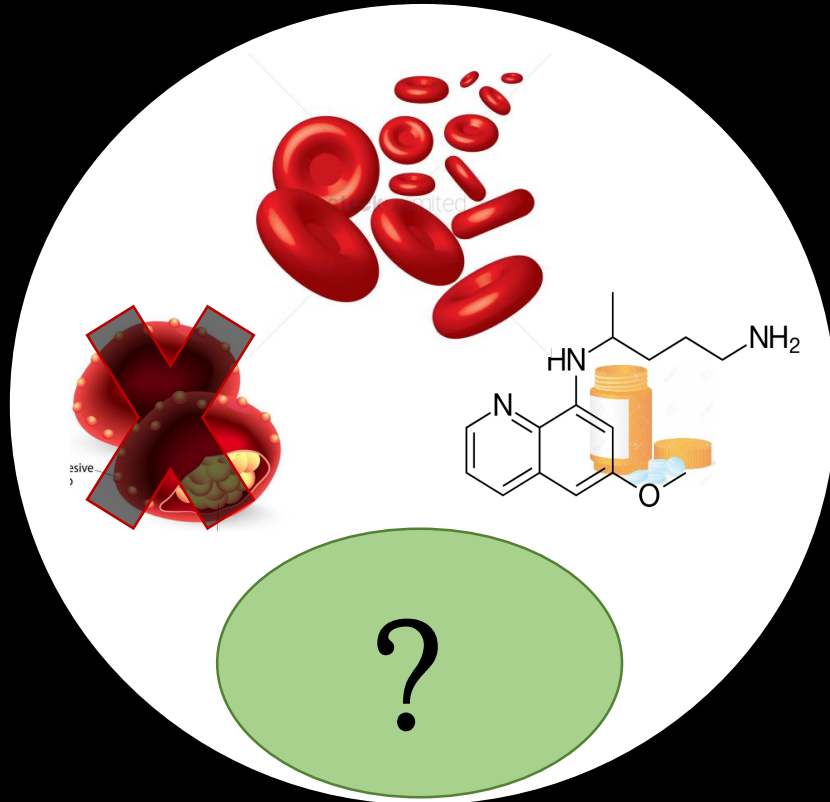


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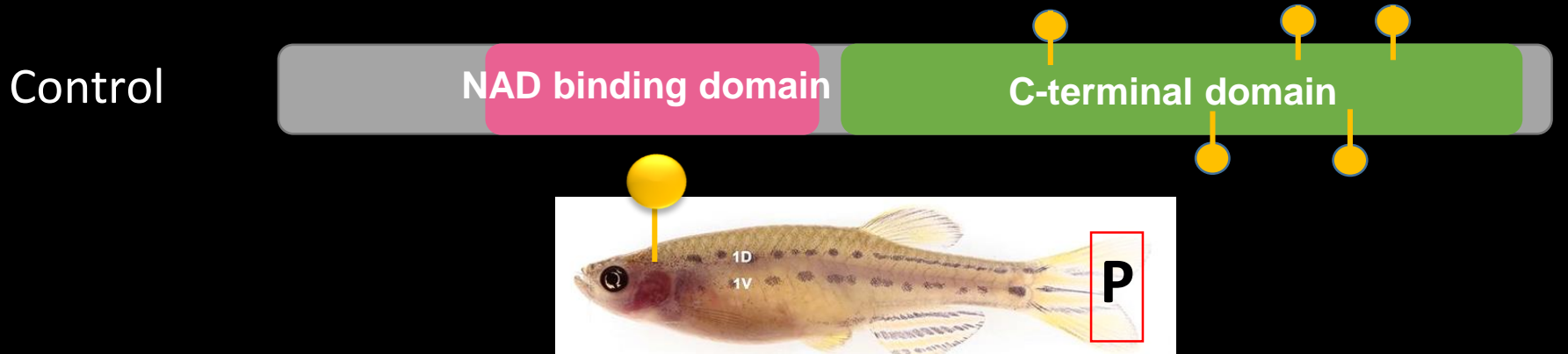
Copyright © 2009 Edita Aksentijevic



Aim 2: Identify a small molecule that affects G6PD methylation destroys malaria, and can be taken with Primaquine without hemolysis

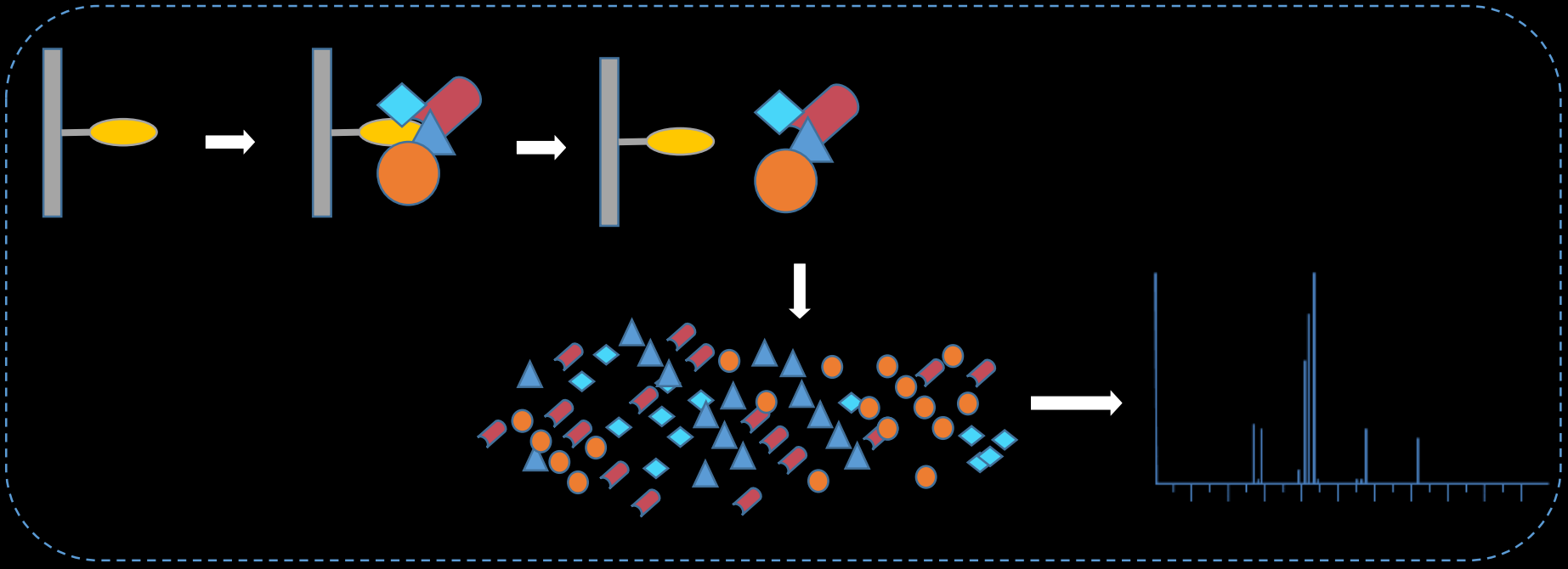


Aim 3: Determine differences in Protein-complexes associated with methylation changes in Pregnant Zebrafish

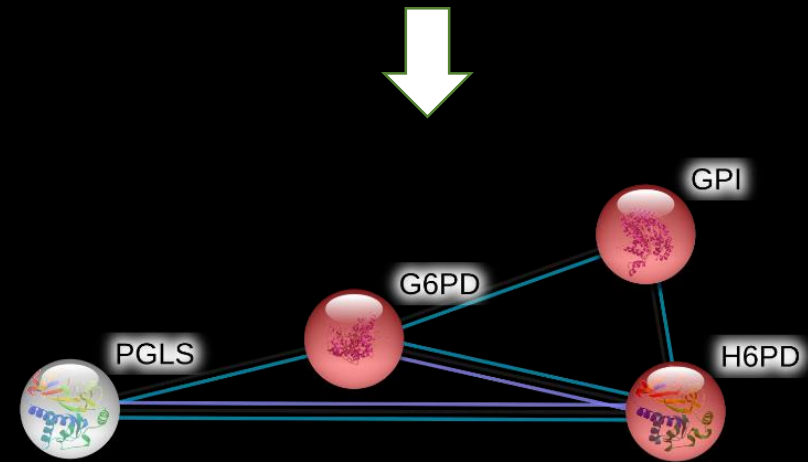
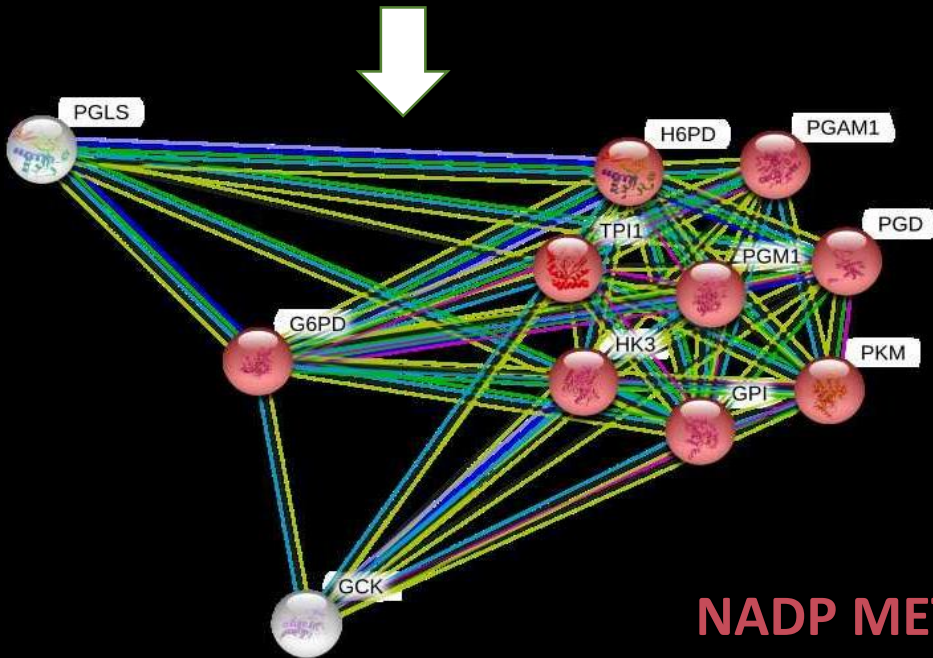


Use CRISPR/Cas9 to create treatment group with no methylation

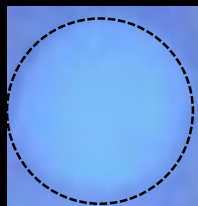
Aim 3: Determine differences in Protein-complexes associated with methylation changes



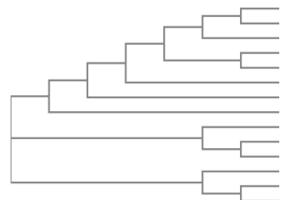
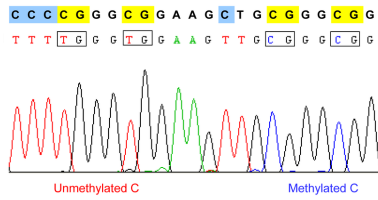
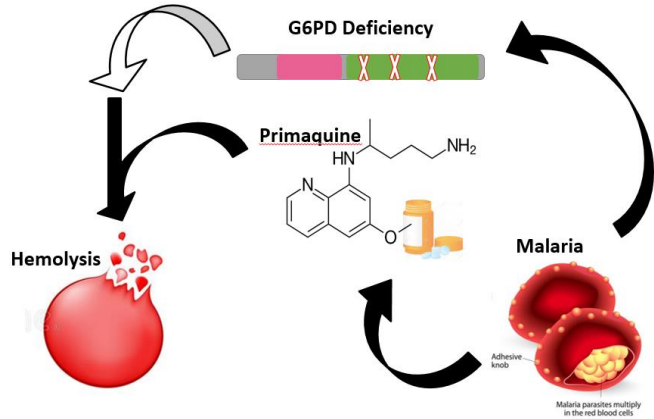
Aim 3: Determine differences in Protein-complexes associated with methylation changes



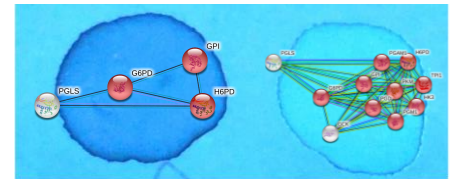
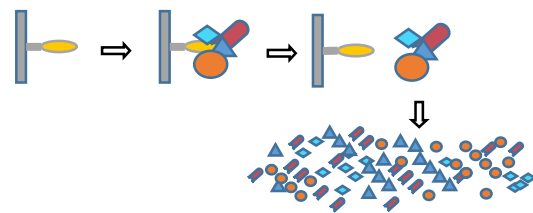
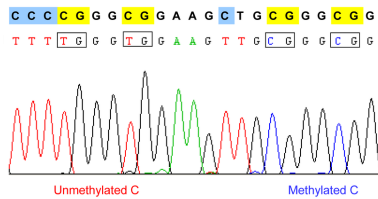
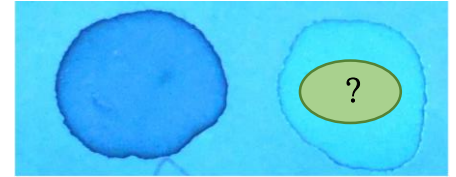
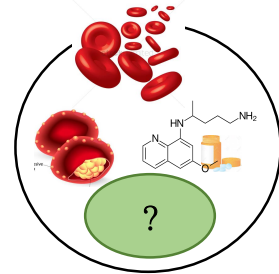
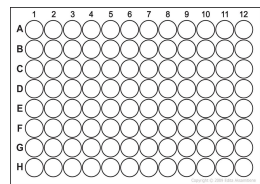
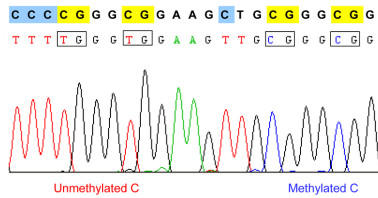
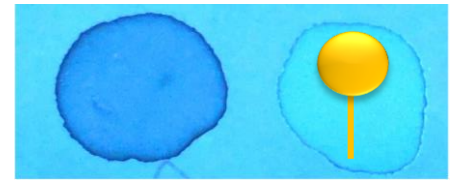
NADP METABOLISM



Conclusions



S. cerevisiae -----HSEGPVKFEKNTVIVFVGSGLAKKKTFFALFGLFREGYLD-PS
A. thaliana RSTFRNDSFVREYGIIVPETGCLSIIVLGASGLAKKKTFFALFGLFREGYLD-PS
C. elegans SKDLVECLRESIQDLKFTPEYVVFVGSGLAKKKIYPTLIMILFRDGLL-PVN
D. melanogaster RK-----ELHDDTAFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PEQ
A. gambiae RE-----ELFSEQEFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PED
X. tropicalis RE-----ELFQGDVAFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PEN
C. lupus familiaris RE-----ELFQGDVAFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PEN
H. sapiens RE-----ELFQGDVAFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PEN
M. mulatta RE-----ELFQGDVAFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PED
P. anubis RE-----ELFQGDVAFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PED
B. taurus RE-----ELFQGDVAFHQSDTHFIIVVGSGLAKKKIYPTLIMILFRDGLL-PED
M. musculus TISLKSPTMICEGTHFDGILPHTFVTFVGSGLAKKKIYPTLIMILFRDGLL-PKP
R. norvegicus RINLKS54HDTGETHFDGICMCPHFVTFVGSGLAKKKIYPTLIMILFRDGLL-PSD



Future Directions

Pharmacogenomics – tailoring drug treatment to one's genes

Determine if methylation effects are similar in humans and other organisms

Discover how to rescue the functional G6PD gene

References

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[https://www.translationalres.com/article/S1931-5244\(14\)00336-3/fulltext](https://www.translationalres.com/article/S1931-5244(14)00336-3/fulltext)

<https://string-db.org/cgi/network.pl?taskId=9ofOGJV7JRtJ>

<https://ourworldindata.org/malaria>

<https://www.ncbi.nlm.nih.gov/books/NBK65119/>

<https://www.nhlbi.nih.gov/health-topics/hemolytic-anemia>