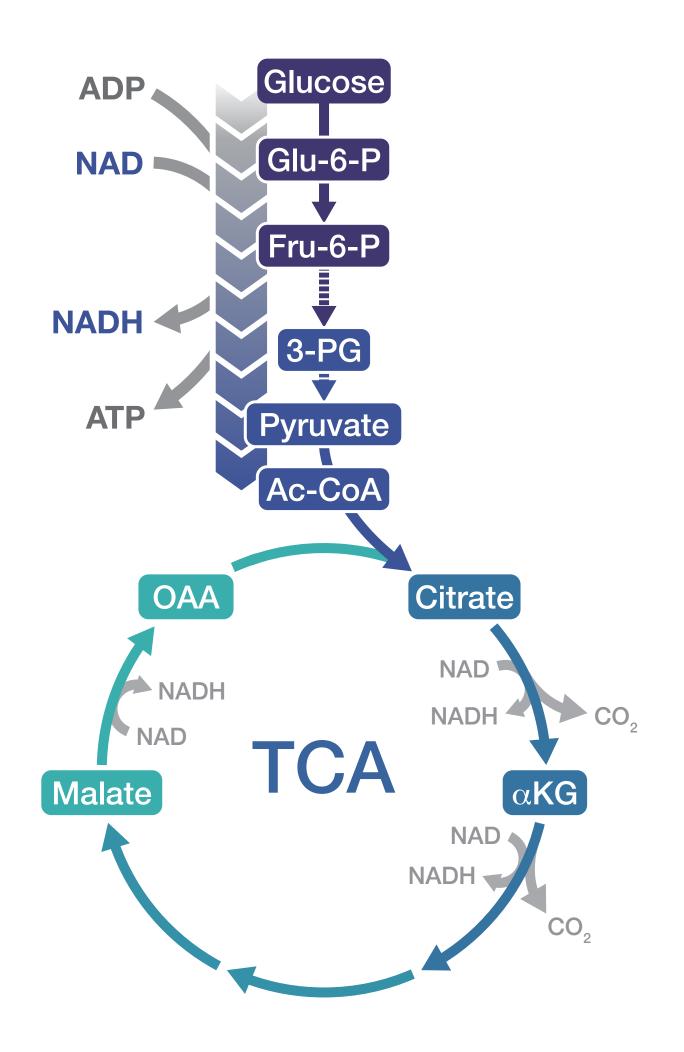


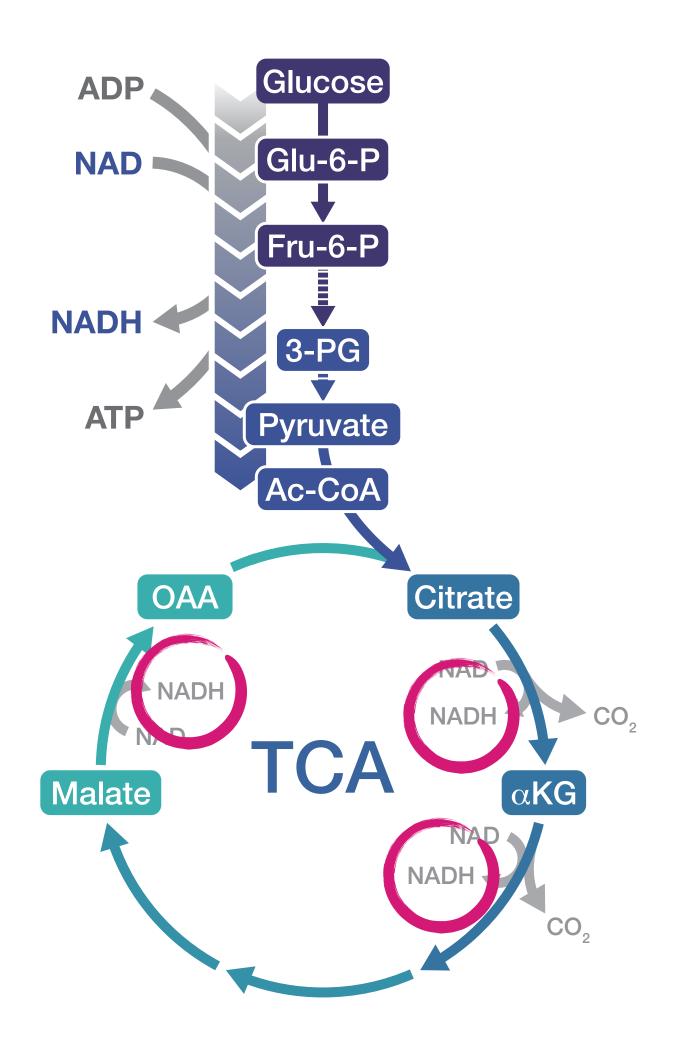


Hans Krebs





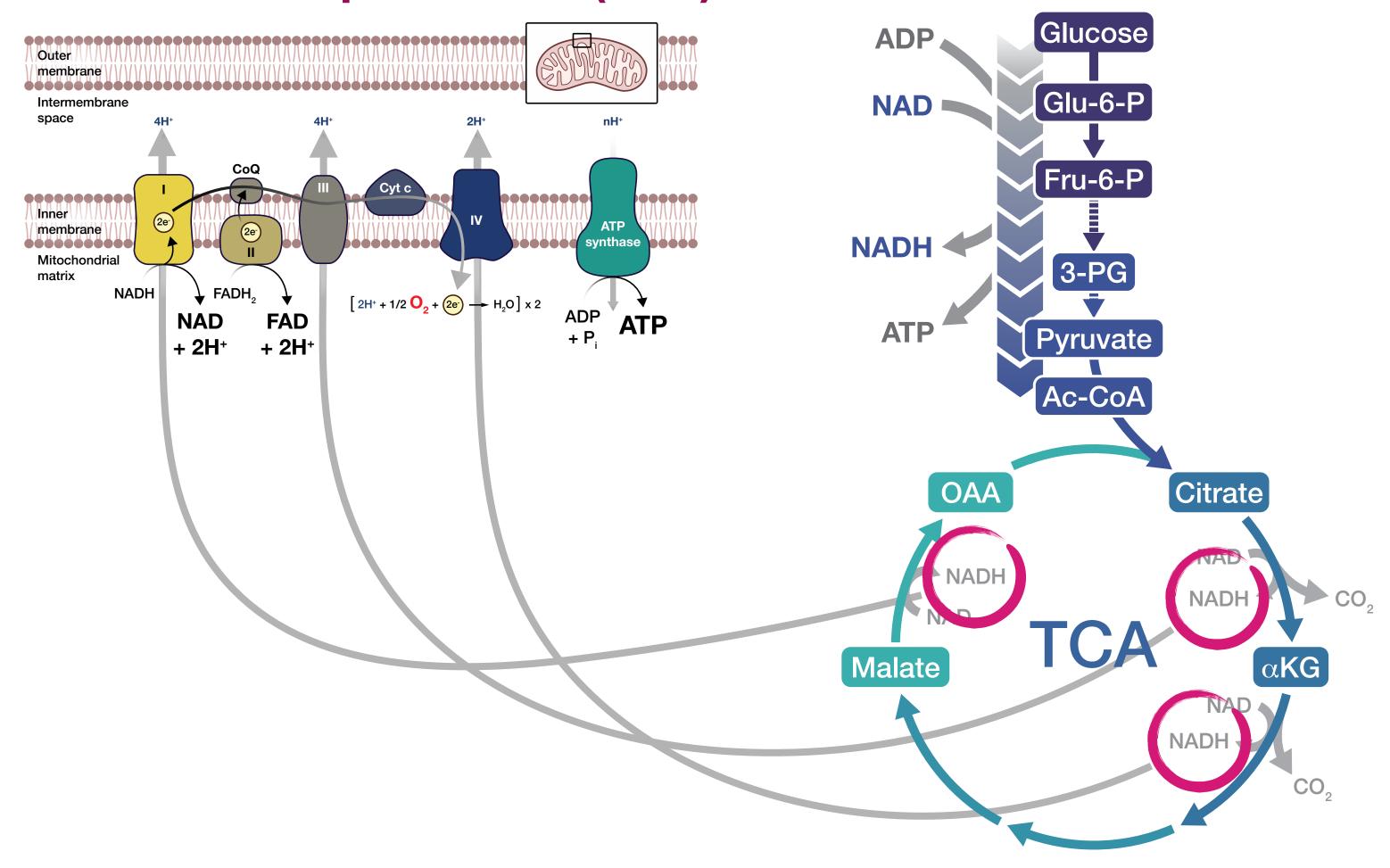
Hans Krebs





Hans Krebs

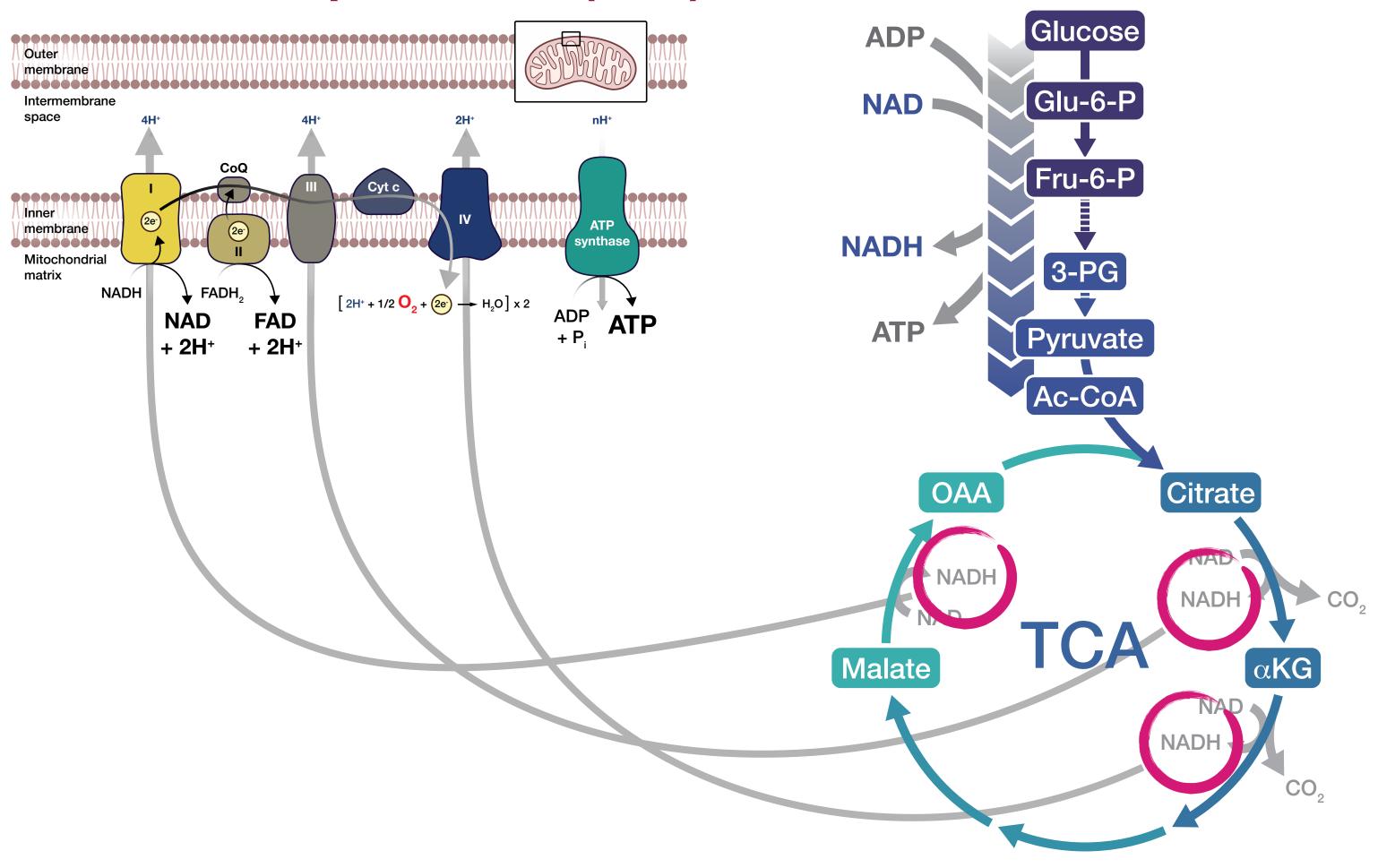
#### **Electron Transport Chain (ETC)**



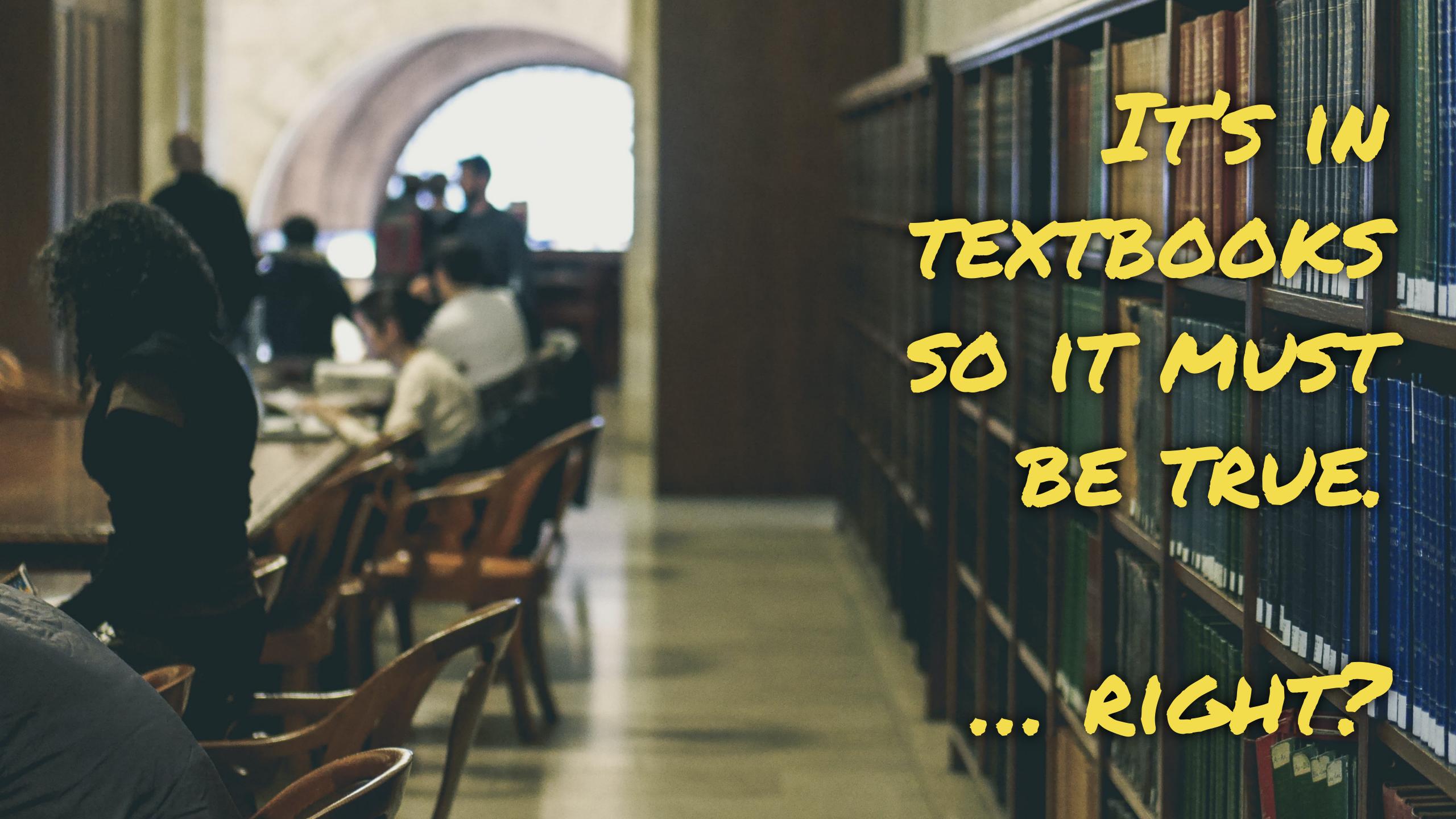


Hans Krebs

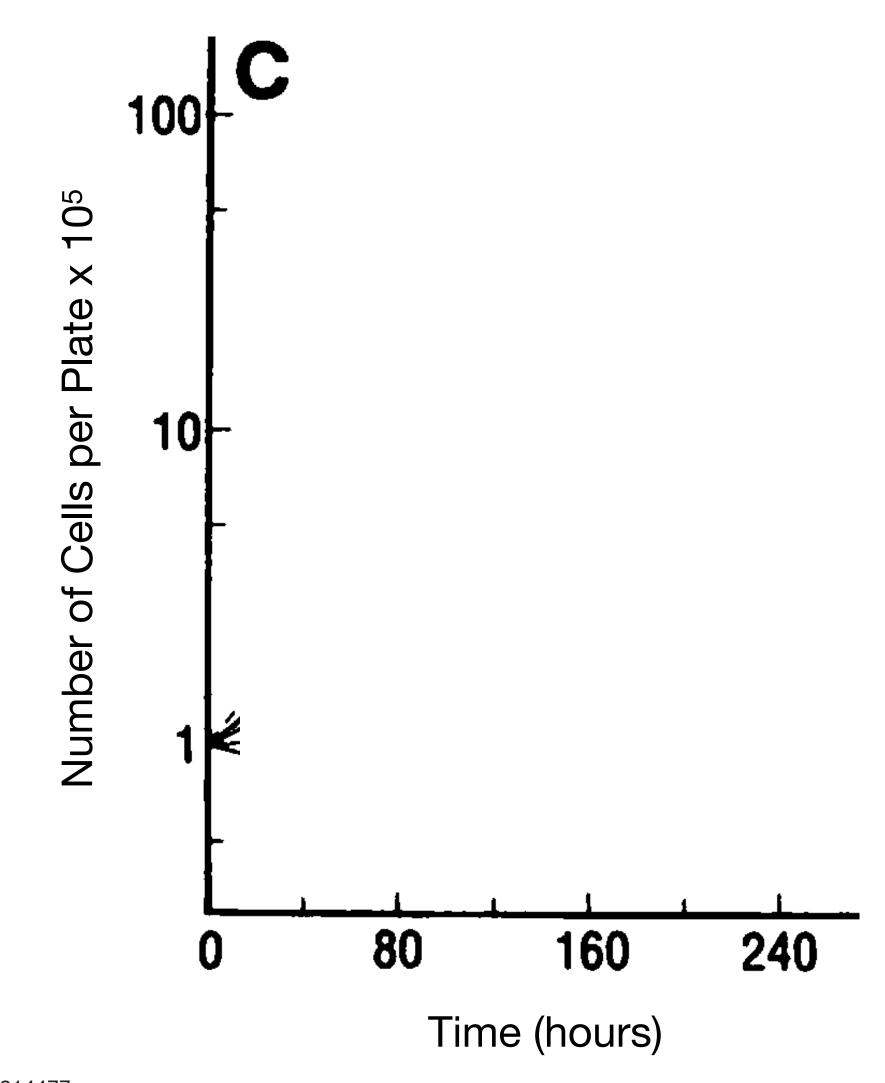
#### **Electron Transport Chain (ETC)**







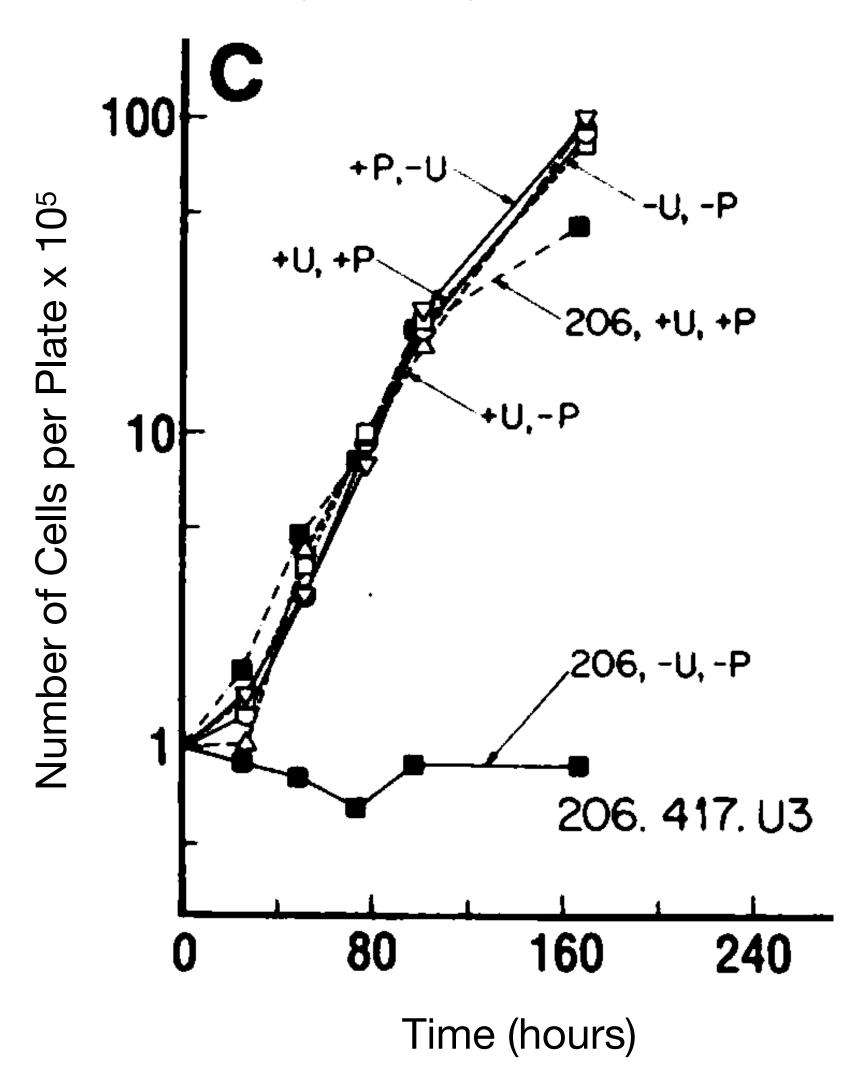
U = uridine P = pyruvate 206 = 143B Rho0 cells (no mtDNA)



# Oxidative Phosphorylation is dispensable for cell proliferation

U = uridine P = pyruvate

206 = 143B Rho0 cells (no mtDNA)

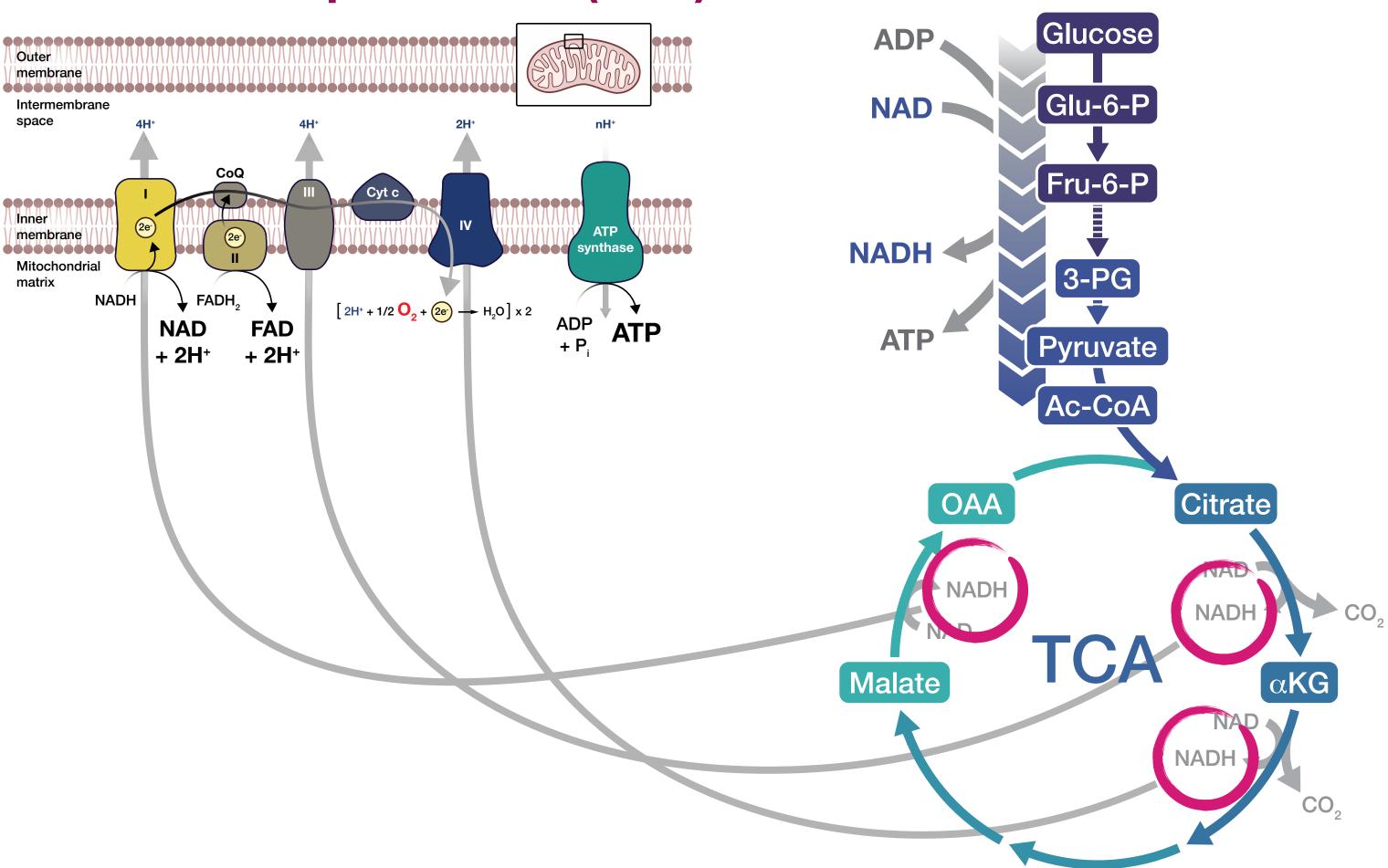


# CELLS WITHOUT AN ELECTRON TRANSPORT CHAIN CAN GROW, AS LONG AS THEY HAVE PYRUVATE AND URIDINE

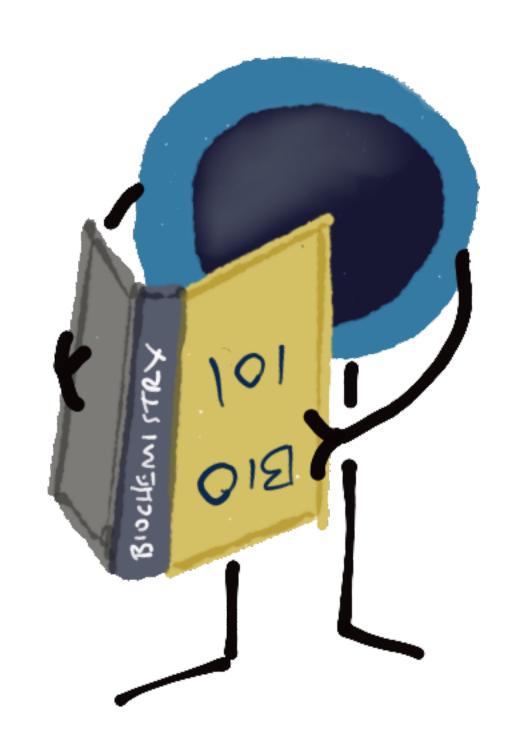
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PMID: 2814477

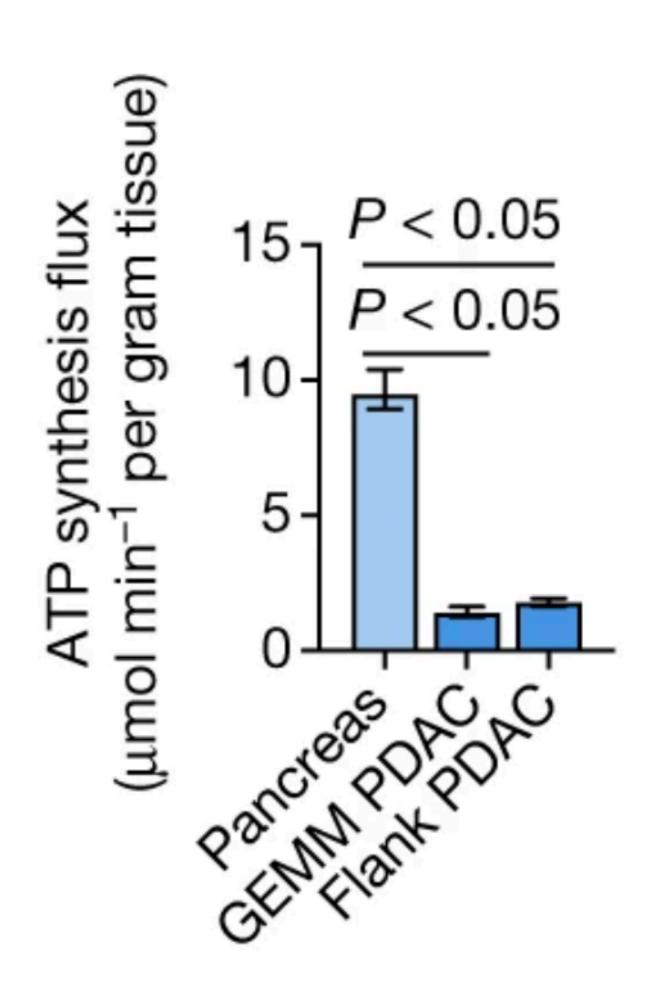
#### **Electron Transport Chain (ETC)**



# TURNS OUT CELLS DON'T READ TEXTBOOKS

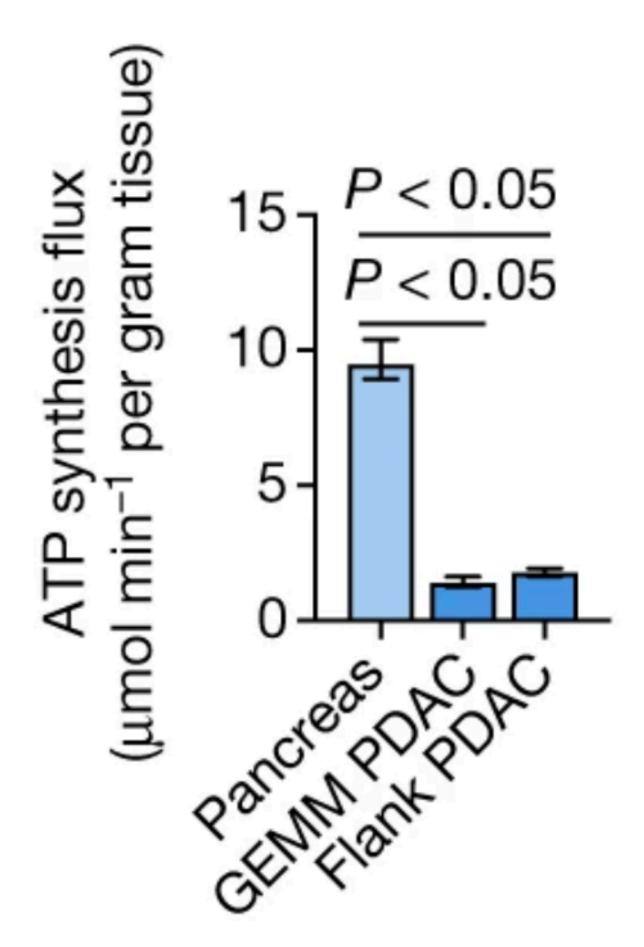


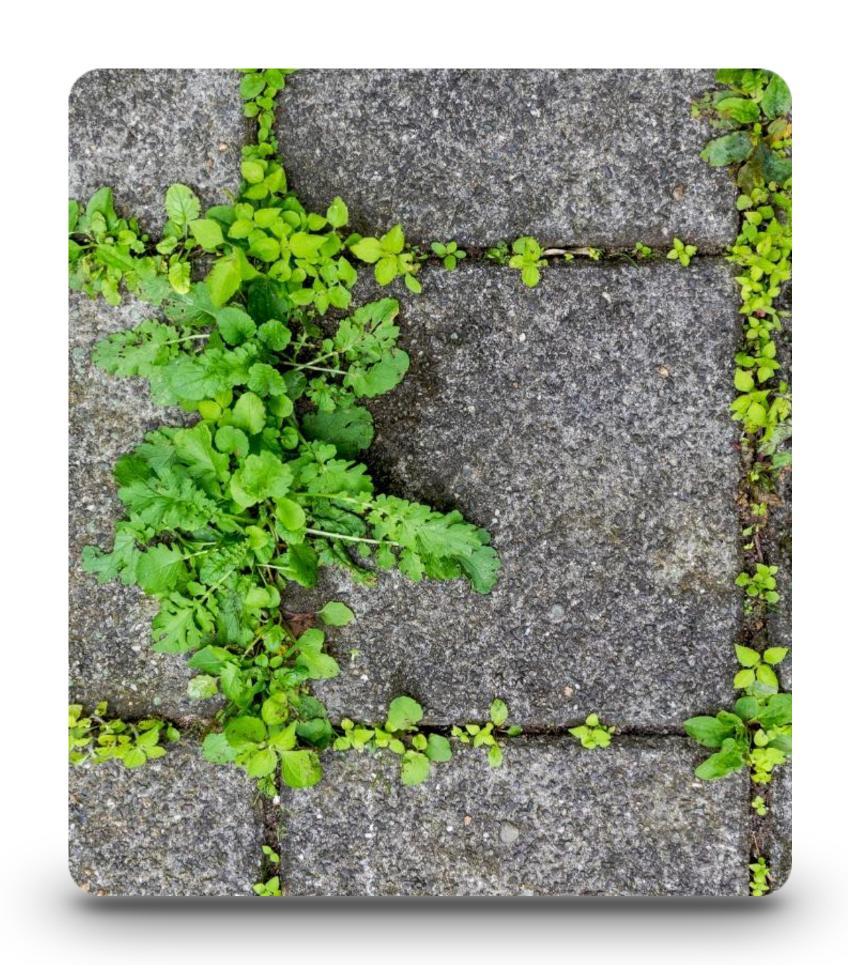
#### Proliferating cells need LESS ATP than differentiated tissues



#### Proliferating cells need LESS ATP than differentiated tissues

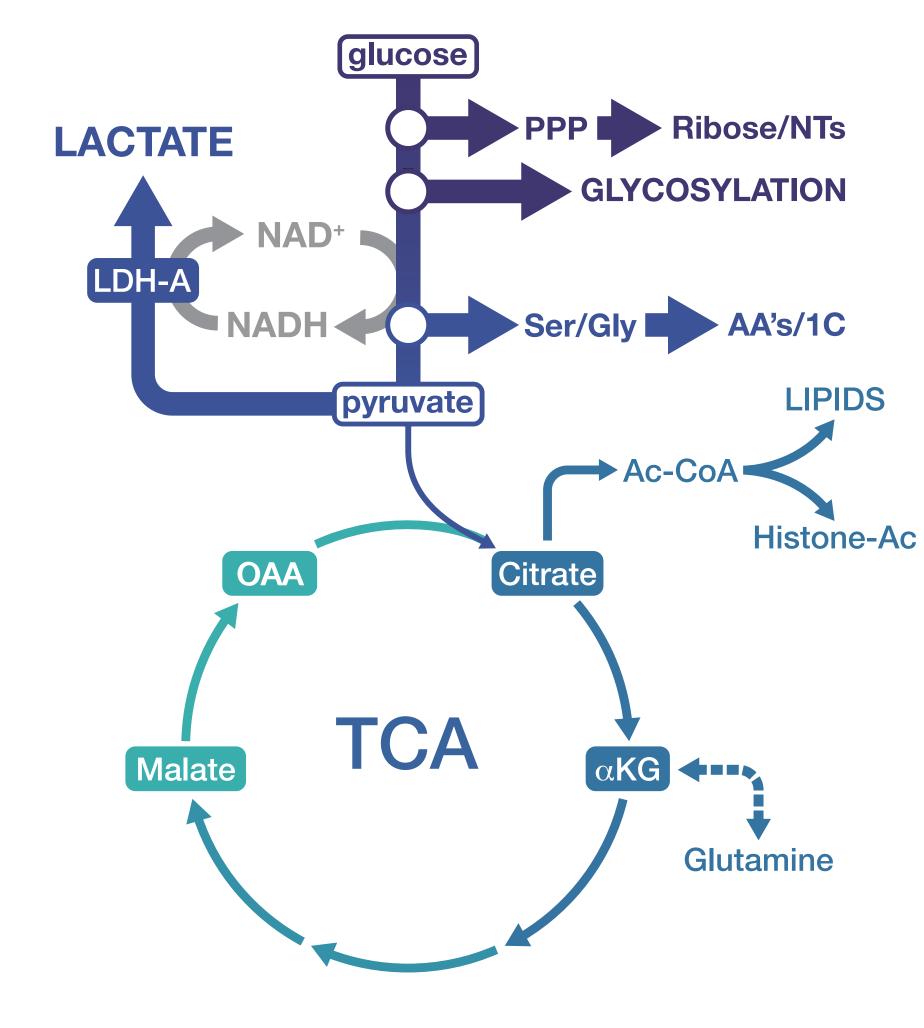


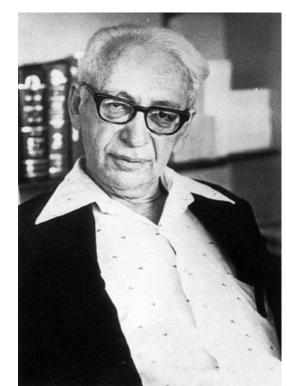




#### Classic Warburg Aerobic Glycolysis

(e.g. highly glycolytic tumors)







Ephraim Racker

Otto Warburg

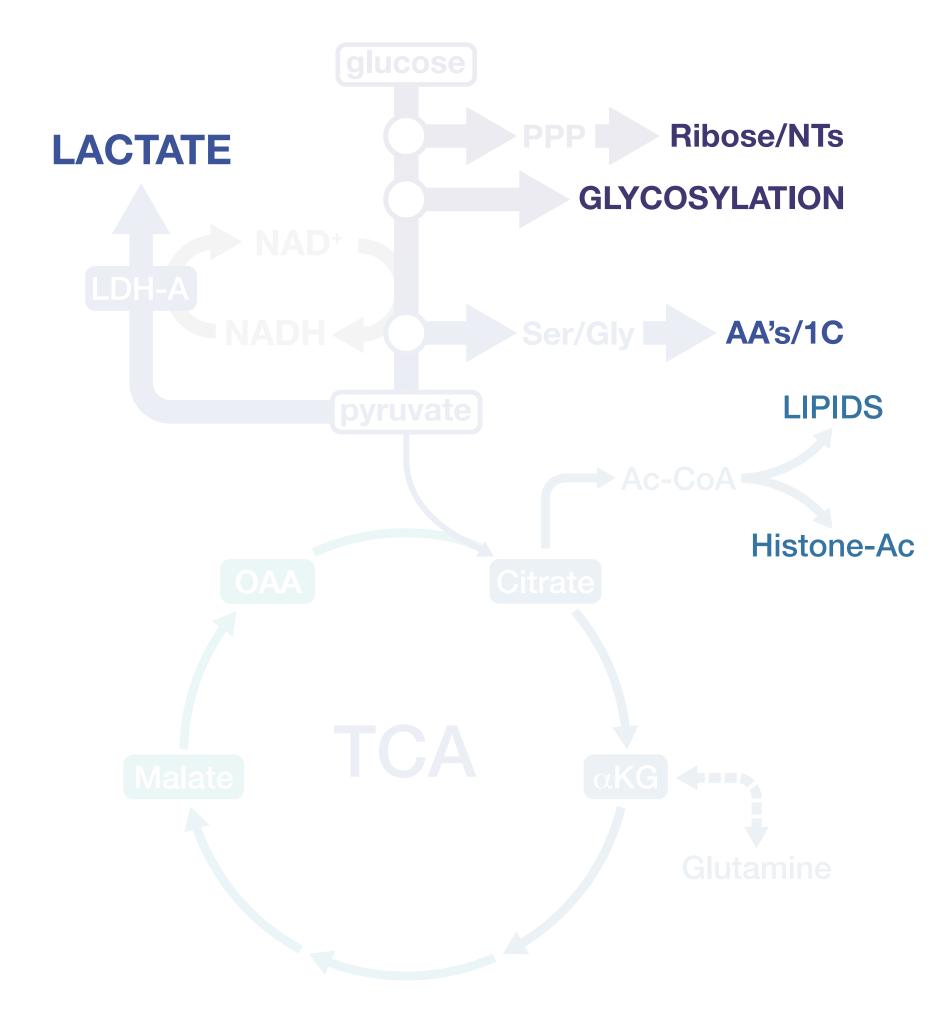
Bioenergetics and the Problem of Tumor Growth

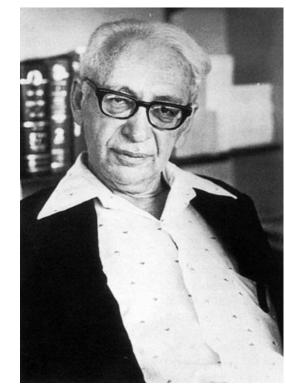
An understanding of the mechanism of the generation and control of biological energy may shed light on the problem of tumor growth



#### Classic Warburg Aerobic Glycolysis

(e.g. highly glycolytic tumors)





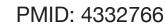


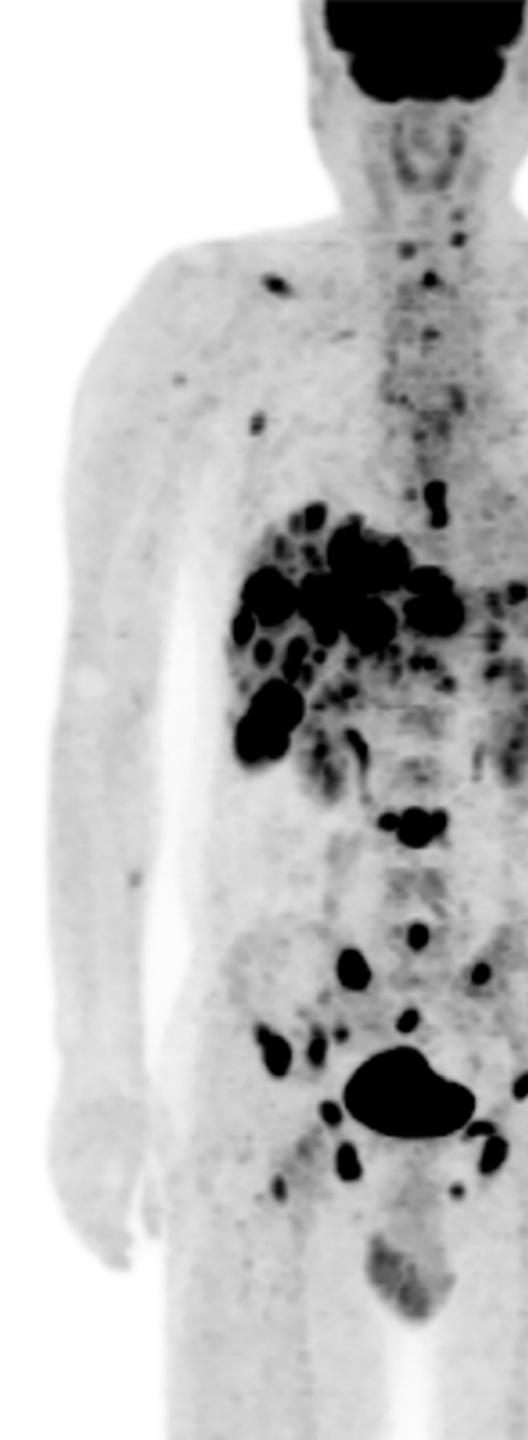
Ephraim Racker

Otto Warburg

Bioenergetics and the Problem of Tumor Growth

An understanding of the mechanism of the generation and control of biological energy may shed light on the problem of tumor growth

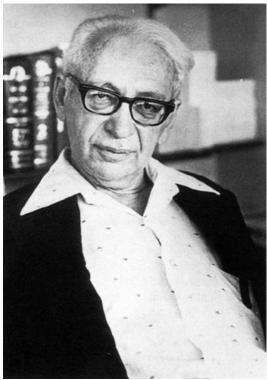




#### Classic Warburg Aerobic Glycolysis

(e.g. highly glycolytic tumors)

#### 





**Ephraim Racker** 

Otto Warburg

Bioenergetics and the Problem of Tumor Growth

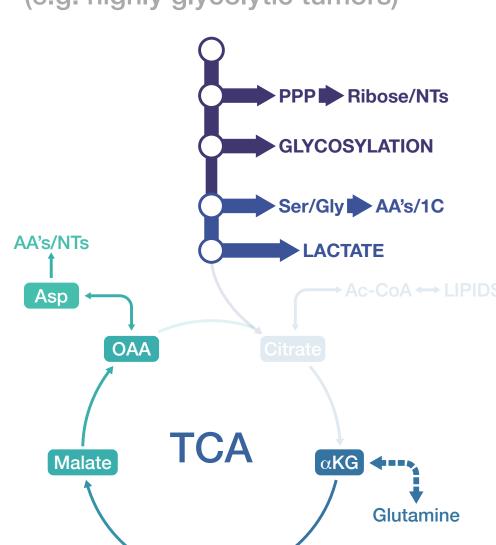
An understanding of the mechanism of the generation and control of biological energy may shed light on the problem of tumor growth



# Cells rewire metabolic pathways to fulfill anabolic, catabolic and regulatory needs

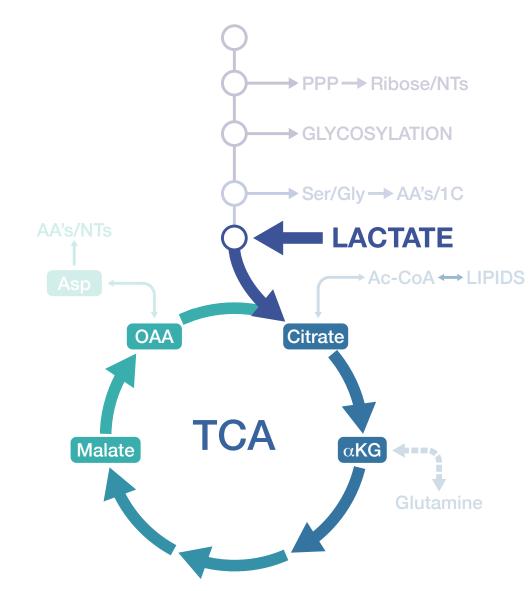
#### Canonical Krebs Cycle **Oxidative Phosphorylation** (e.g. muscle, neurons) → PPP → Ribose/NTs → GLYCOSYLATION Ser/Gly → AA's/1C → LACTATE Ac-CoA ← LIPIDS Citrate TCA NAD + 2H<sup>+</sup> FAD + 2H<sup>+</sup> **Mitochondrial** matrix Intermembrane space

### Classic Warburg Aerobic Glycolysis (e.g. highly glycolytic tumors)



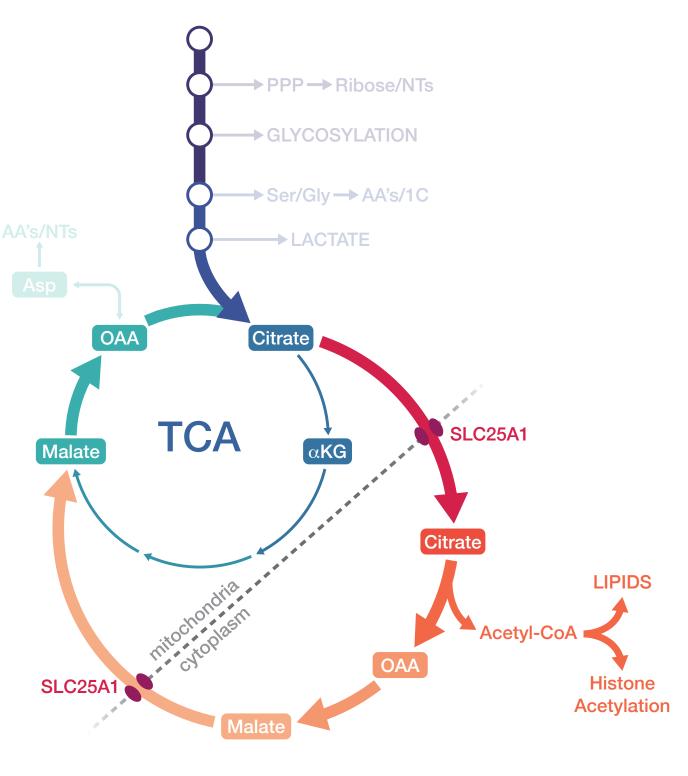
#### Lactate Oxidation

(e.g. cancer cells in normoxia, neurons)



#### Non-Canonical Krebs Cycle

(e.g. ES cells, myoblasts)

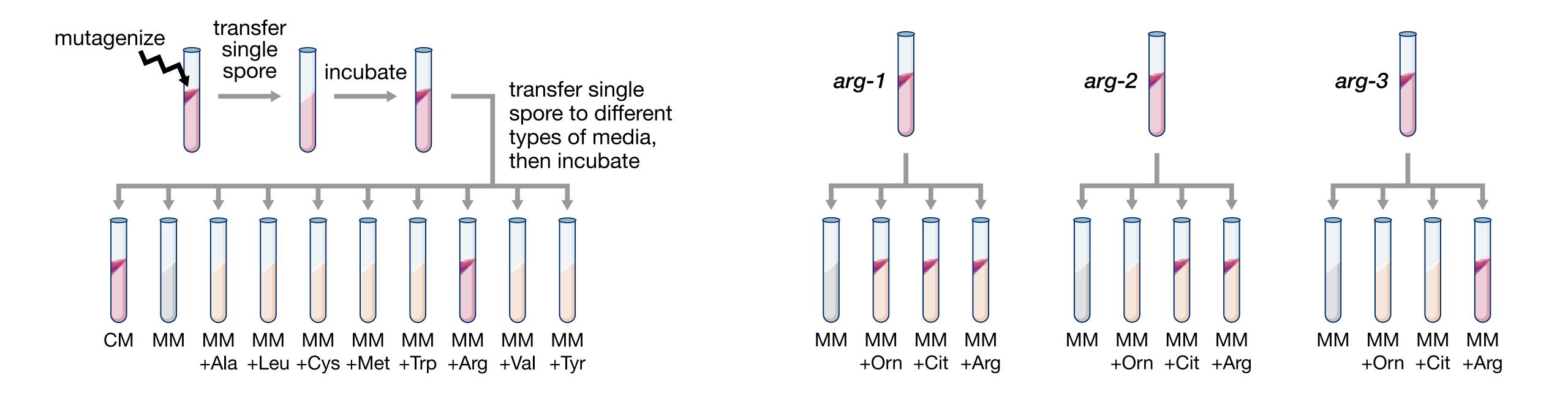


## Takehome points 1

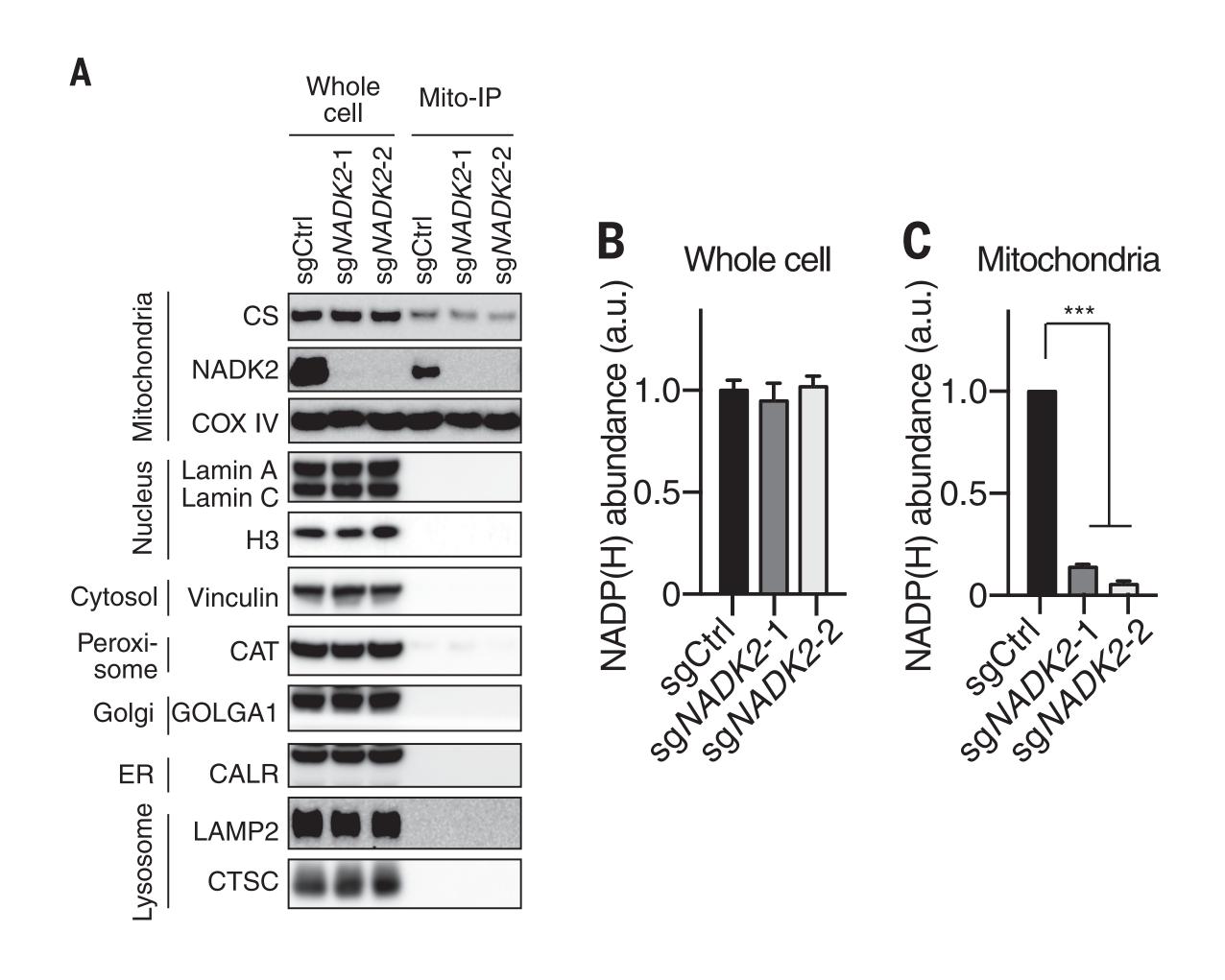
- 1. Metabolic flux is malleable (many ways to make a product)
- 2. Metabolism is not ALWAYS about energy
- 3. Growth (anabolic metabolism) requires SUBSTRATES >>> ENERGY
- 4. Cells (and organisms) rewire their metabolism to suit their needs

# METABOLISM AFFECTS ALL CELLULAR PROCESSES... BUT HOW DO WE FIND OUT WHICH METABOLITES INFLUENCE SPECIFIC PROCESSES AND HOW?

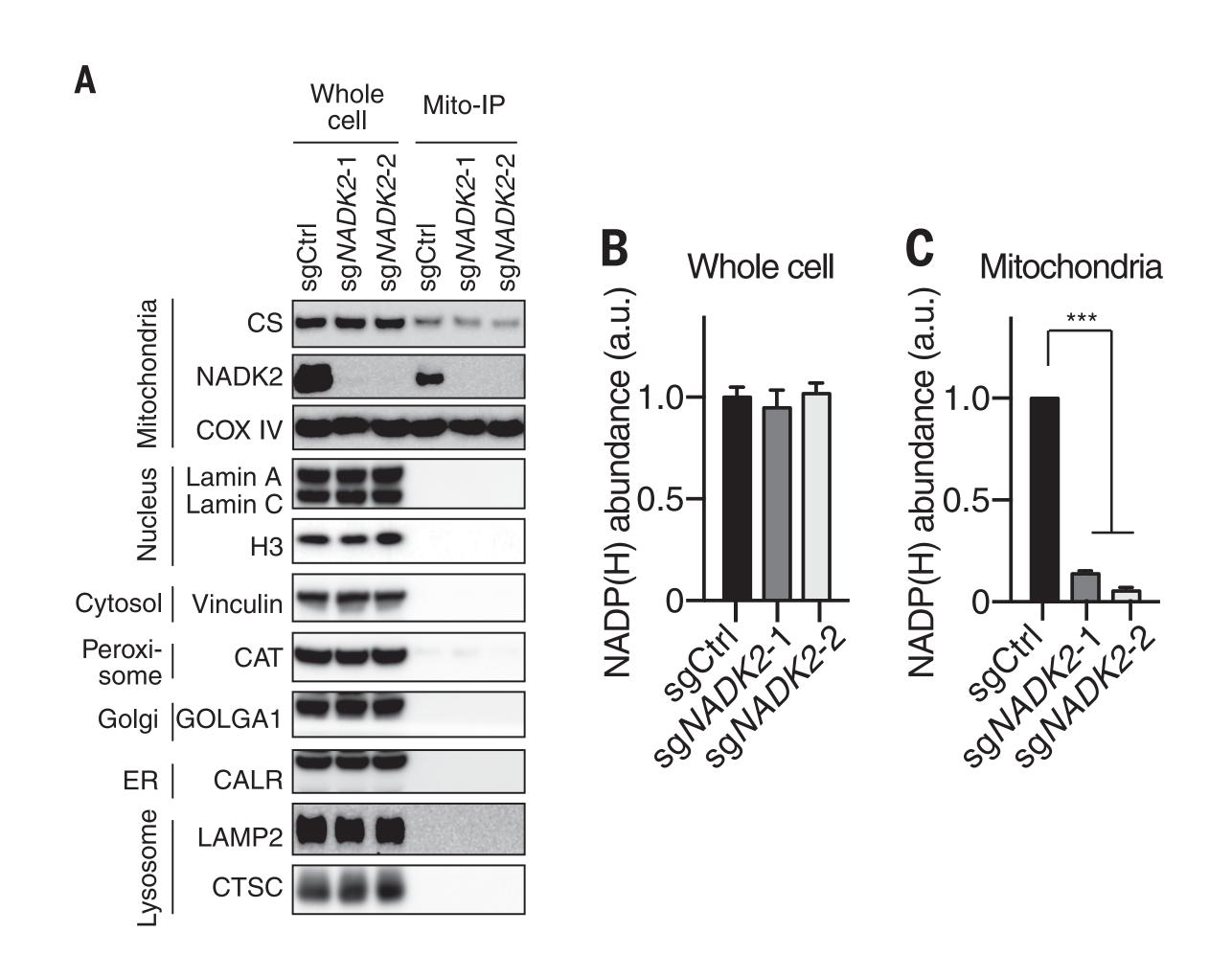
#### Auxotrophs & Prototrophs: the genetics of biochemistry

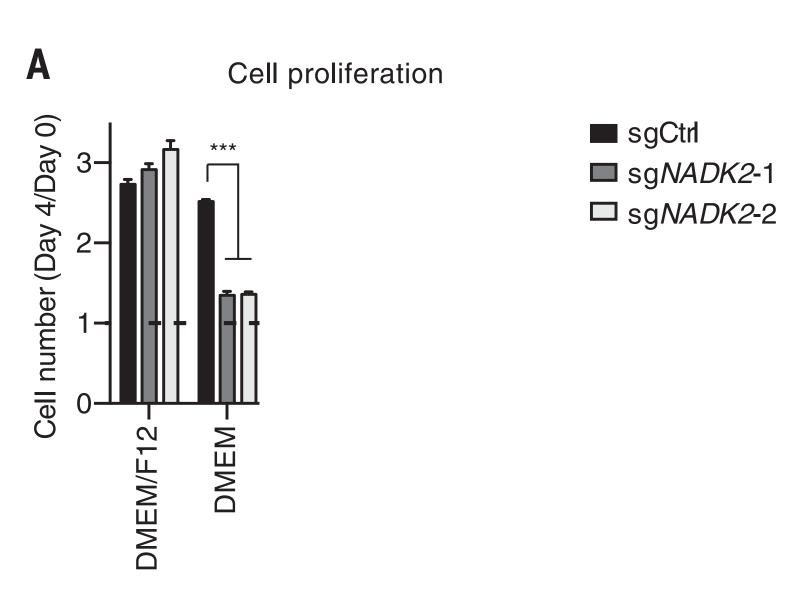


Precursor 
$$\underbrace{\begin{array}{c} Gene A \\ arg-1 \end{array}}_{\text{NH}_2} \underbrace{\begin{array}{c} Gene B \\ arg-2 \end{array}}_{\text{NH}_2} \underbrace{\begin{array}{c} Gene C \\ NH_2 \end{array}}_{\text{NH}_2} \underbrace{\begin{array}{c} Gene C \\ arg-3 \end{array}}_{\text{NH}_2} \underbrace{\begin{array}{c} NH \\ NH_2 \end{array}}_{\text{NH}_2} \underbrace{\begin{array}{c$$

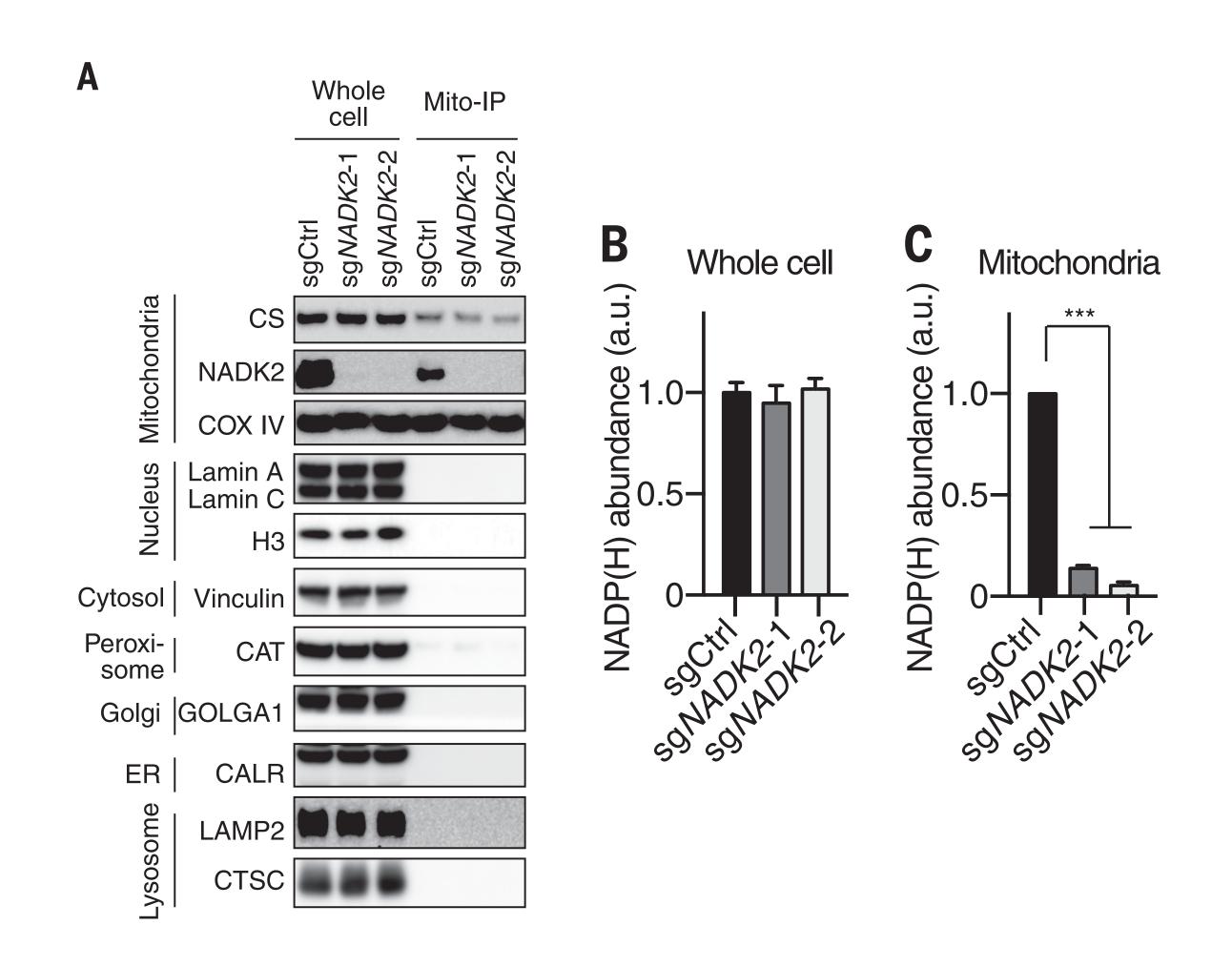


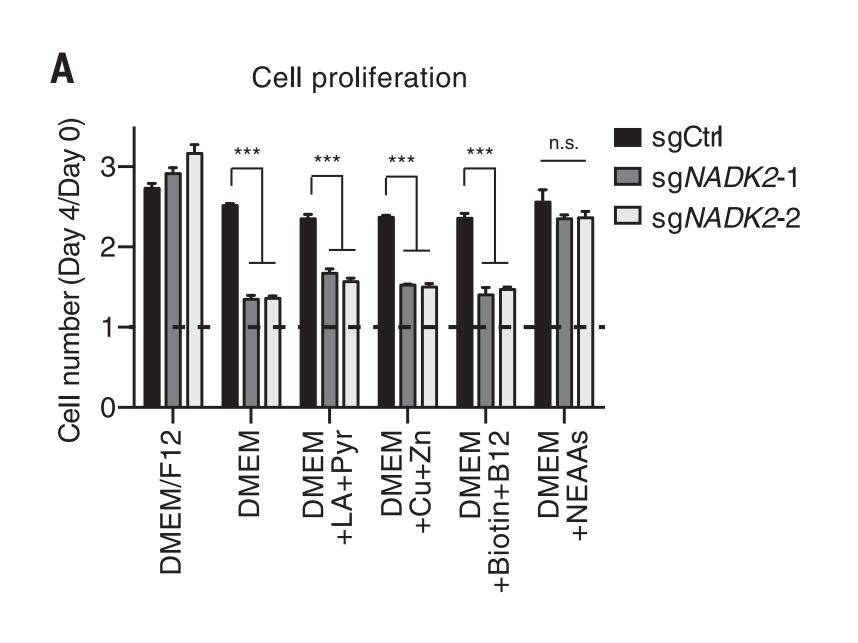
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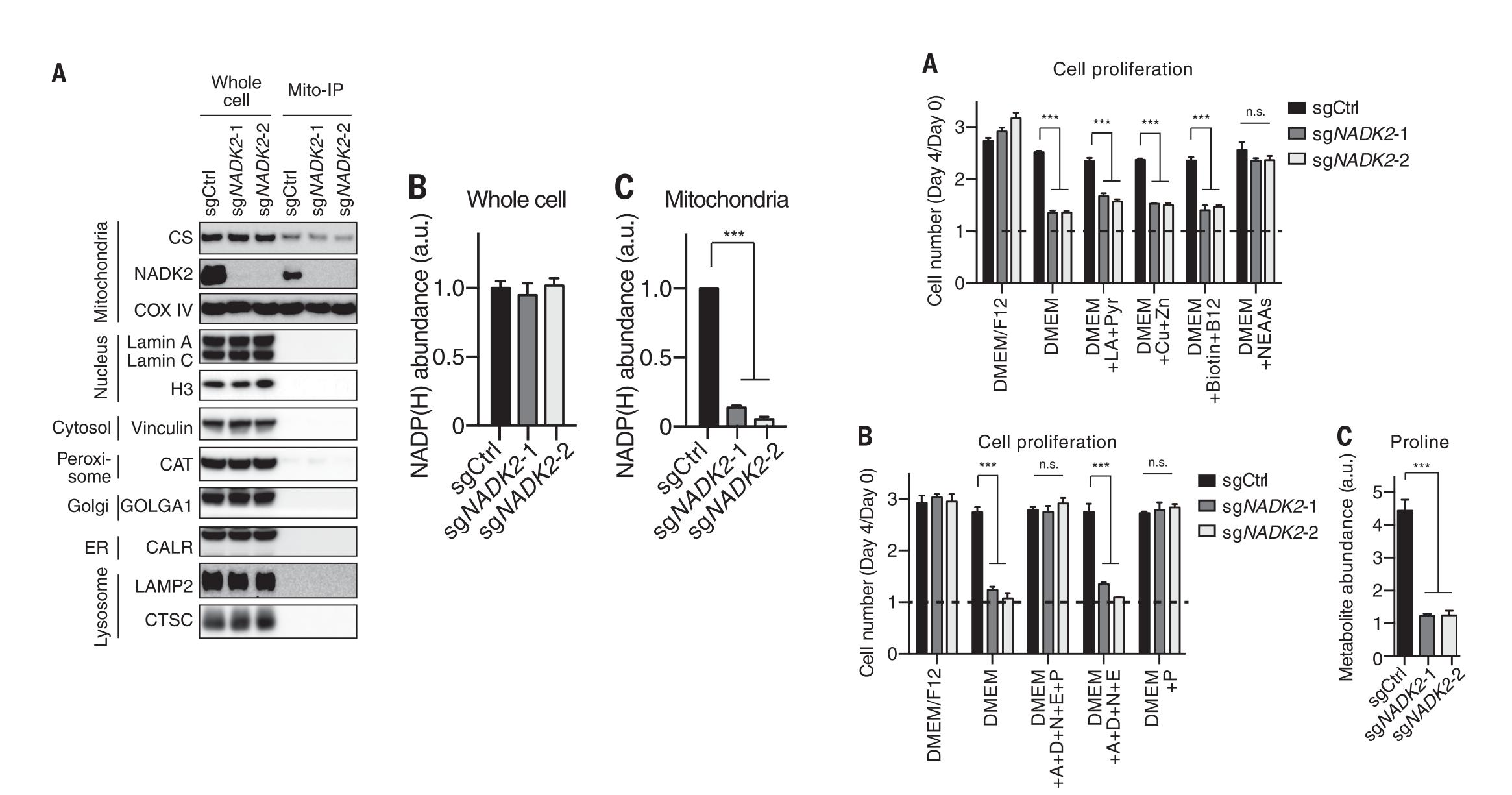




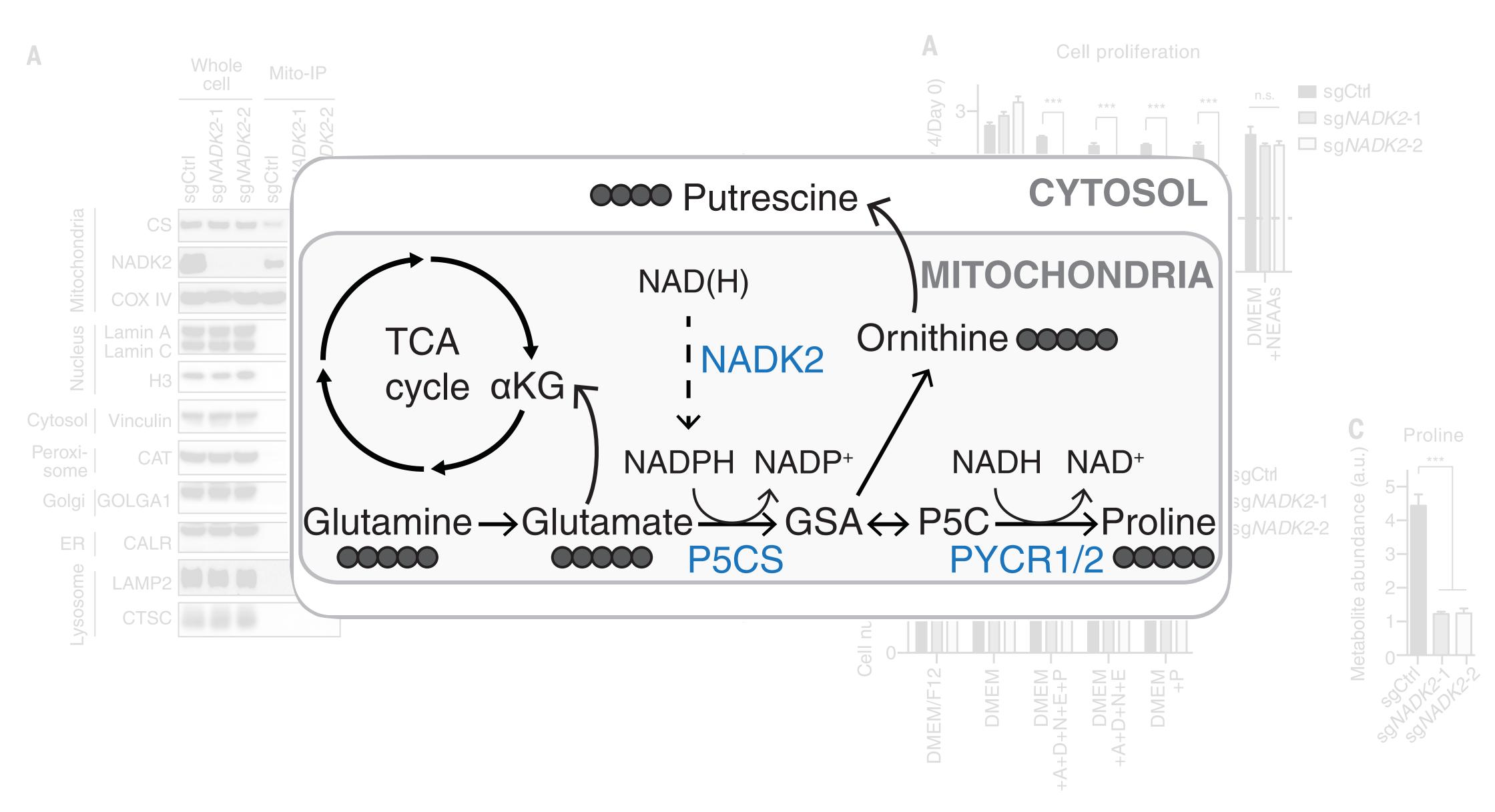
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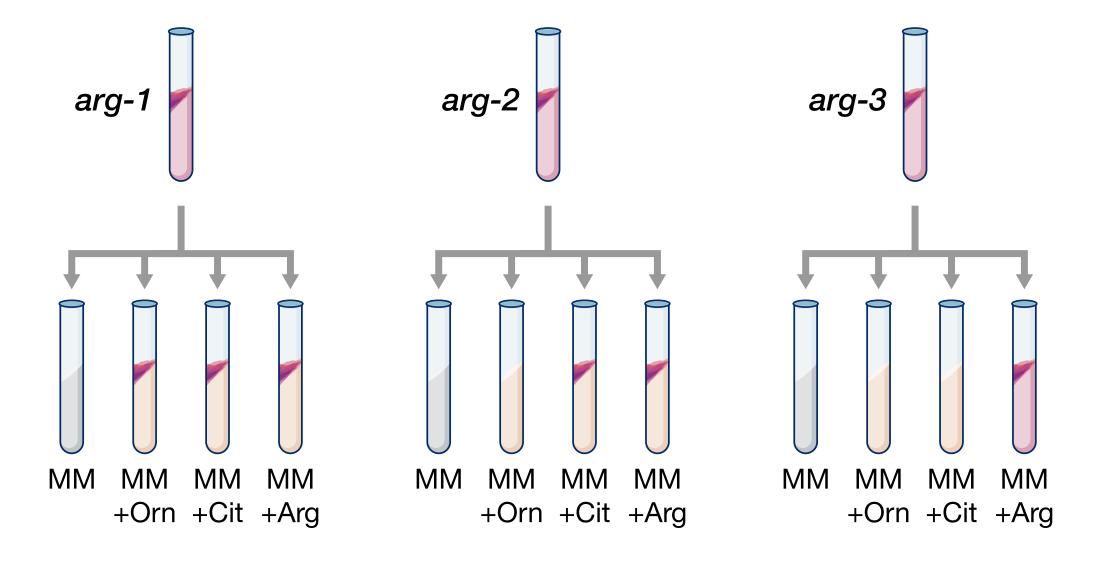
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pmid: 33888598

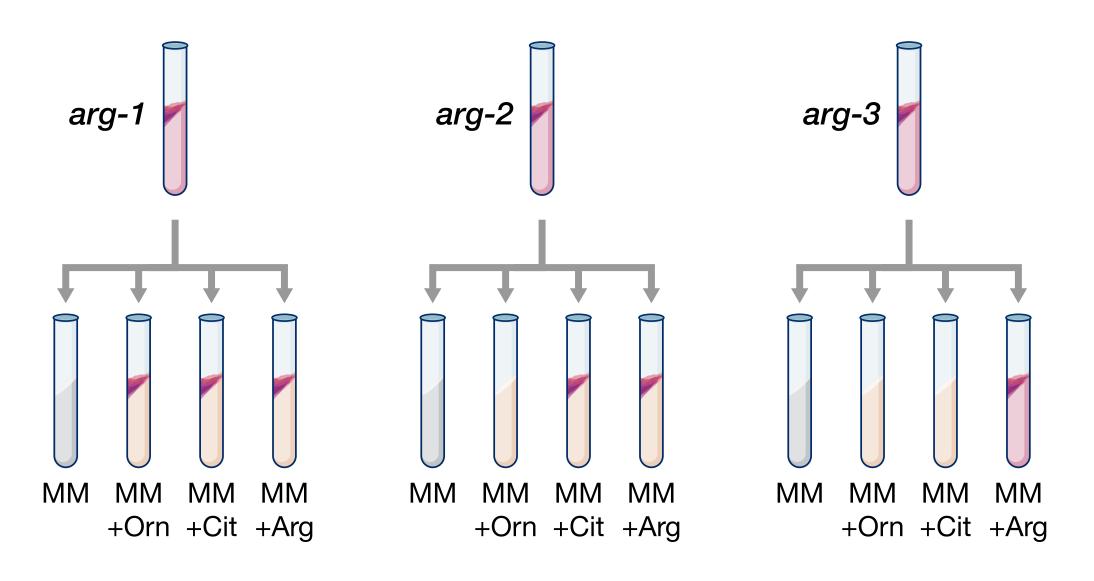
#### What about screens? (1)

#### SINGLE CELL CLONES (1941)

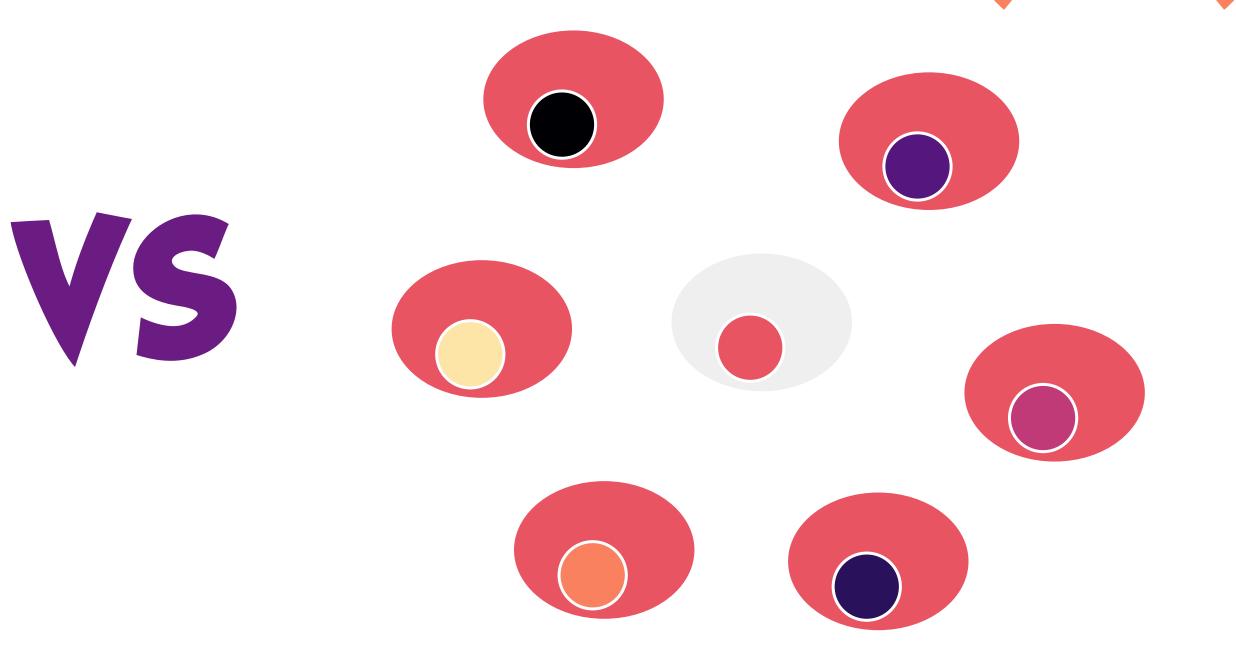


#### What about screens? (1)

#### SINGLE CELL CLONES (1941)

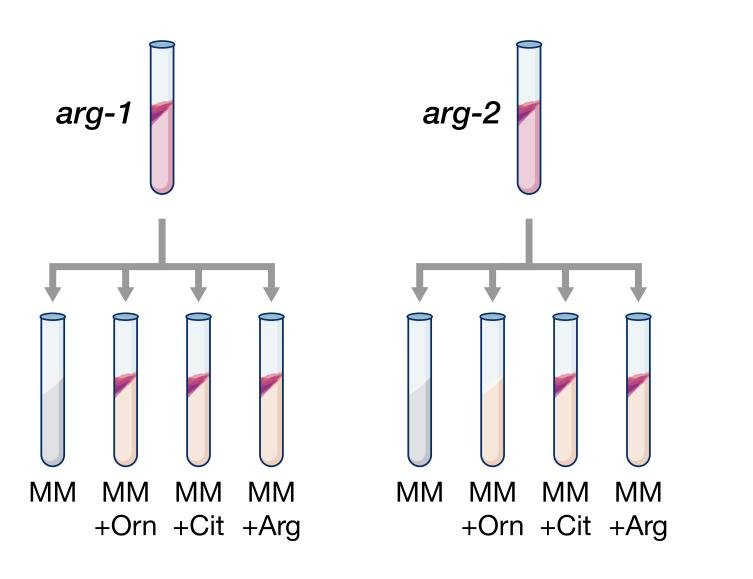


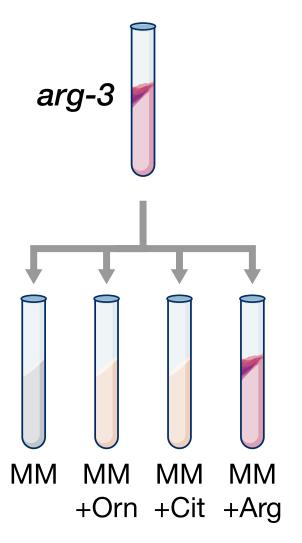
#### POOLED LIBRARIES (2025)



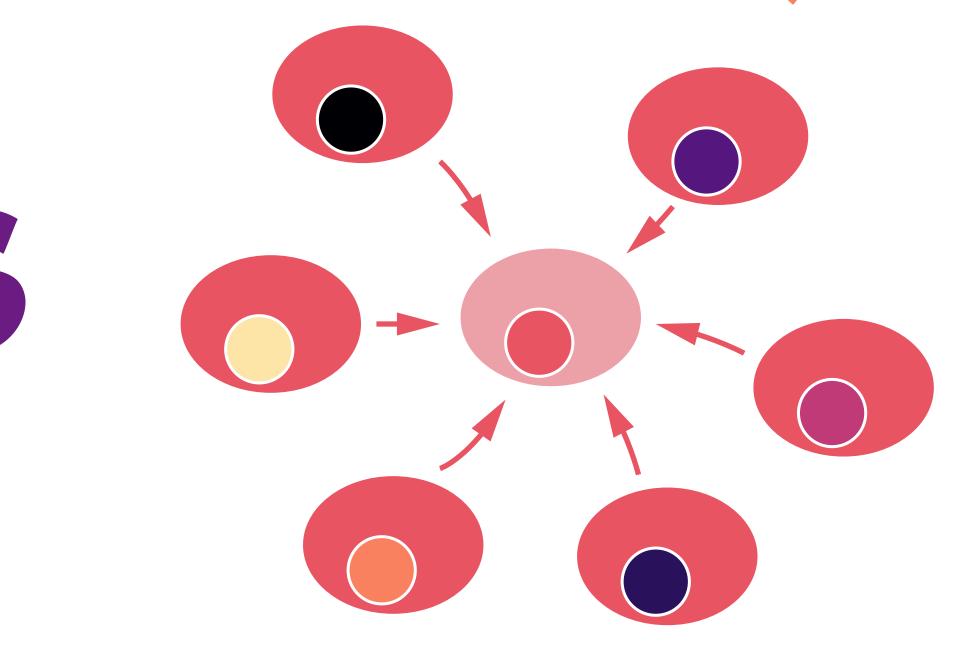
#### What about screens? (1)

#### SINGLE CELL CLONES (1941)





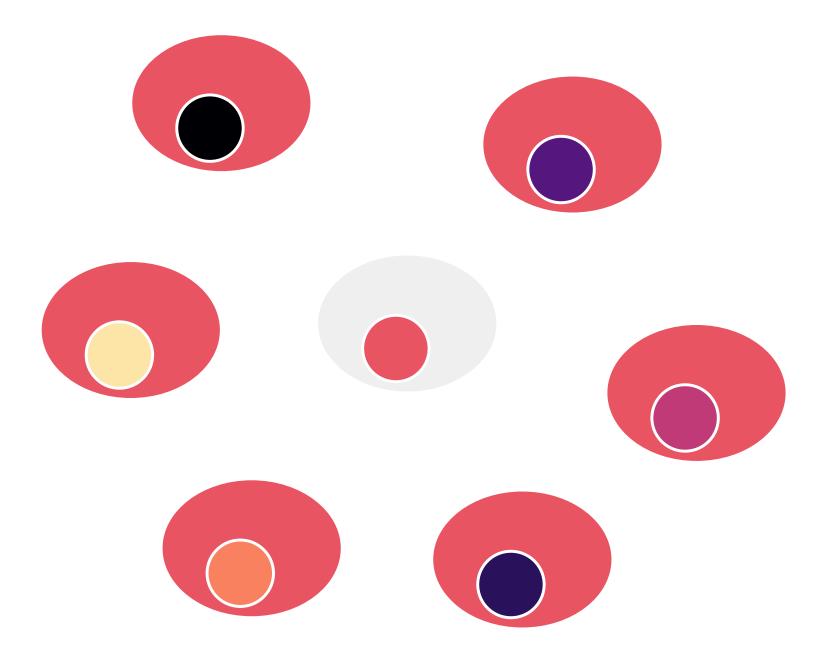
POOLED LIBRARIES (2025)



## NEIGHBORS CAN RESCUE

#### What about screens? (2)



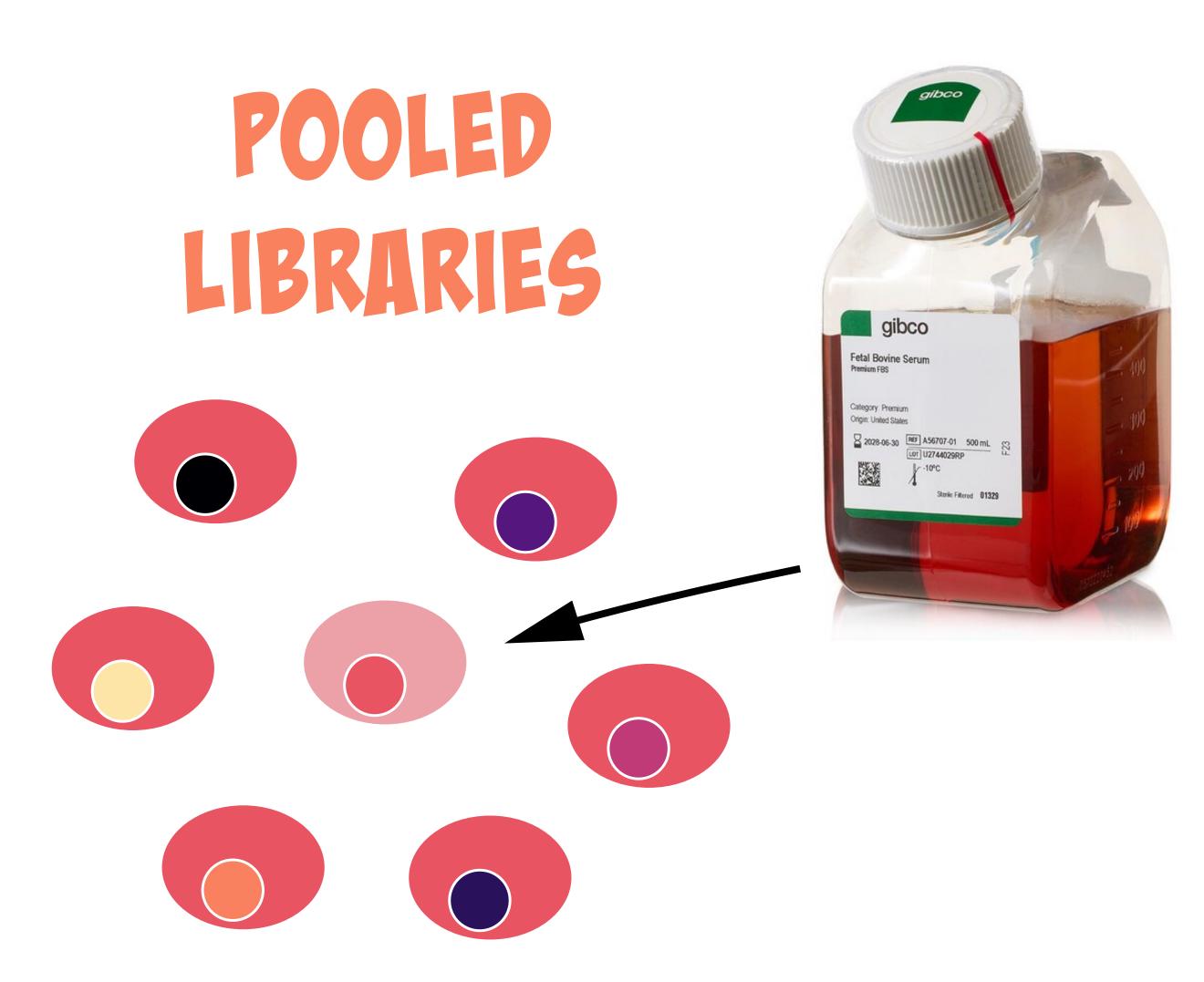


#### What about screens? (2)



#### What about screens? (2)

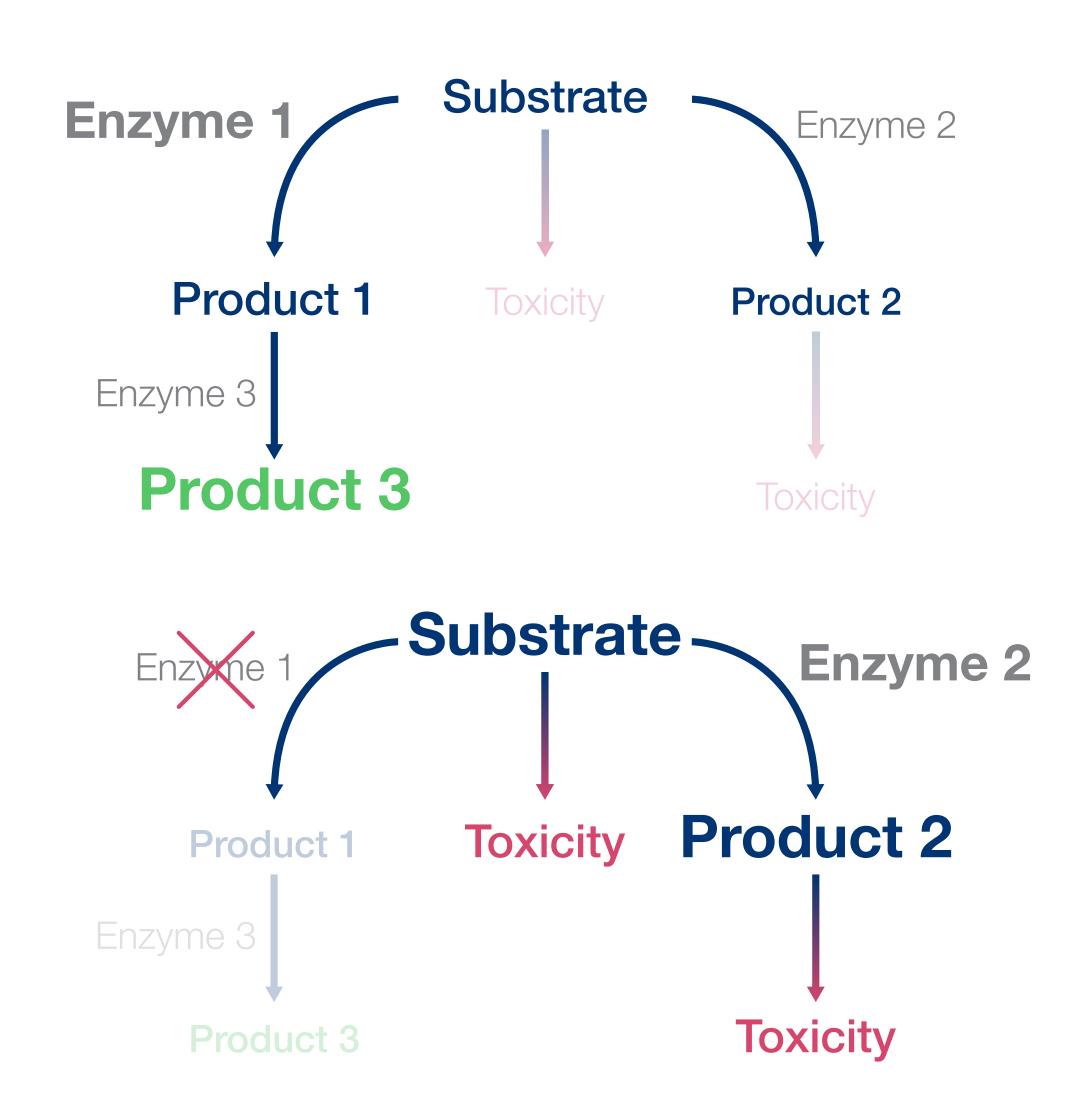
SERUM METABOLITES CAN RESCUE



## Inborn errors of metabolism (IEMs): a window in to what matters and where

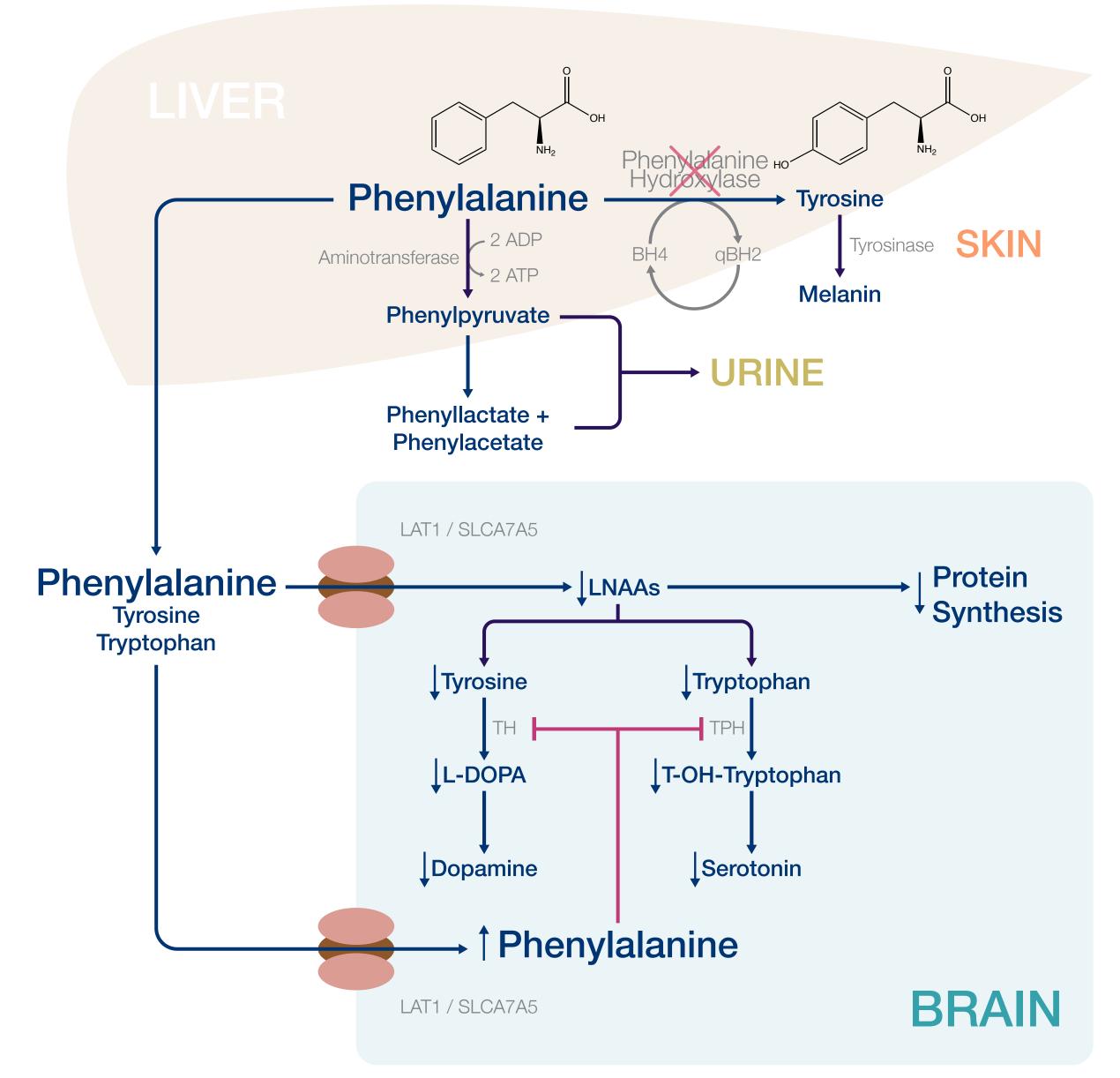
Inborn errors of metabolism (IEMs): a genetic aberration that disrupts a single metabolic pathway

- 1. End-product depletion
- 2. Substrate accumulation
- Activation of alternative pathways producing toxic by-products
- 4. A combination of the above



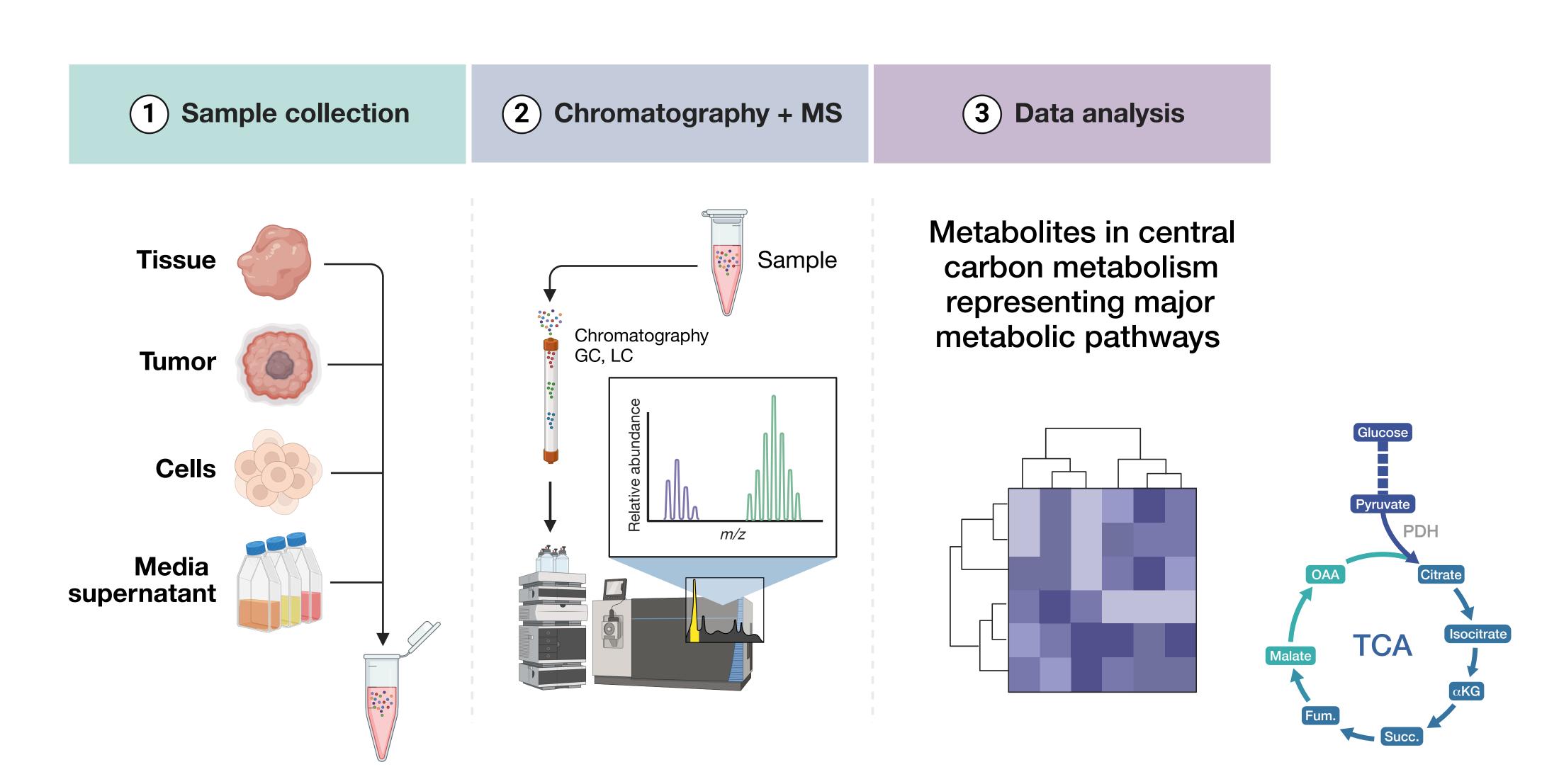
#### Phenylketonuria

- 1. Deficiency of **Phenylalanine Hydroxylase** from mutations in *PAH*
- 2. Inheritance: autosomal recessive
- 3. 111 levels of dietary phenylalanine in plasma
- 4. Clinical features:
  - CNS: intellectual disability, seizures, behavioral problems
  - Urine: musty smell (breakdown products)
  - **Skin**: decreased melanin
- 5. Treatment: dietary phenylalanine avoidance



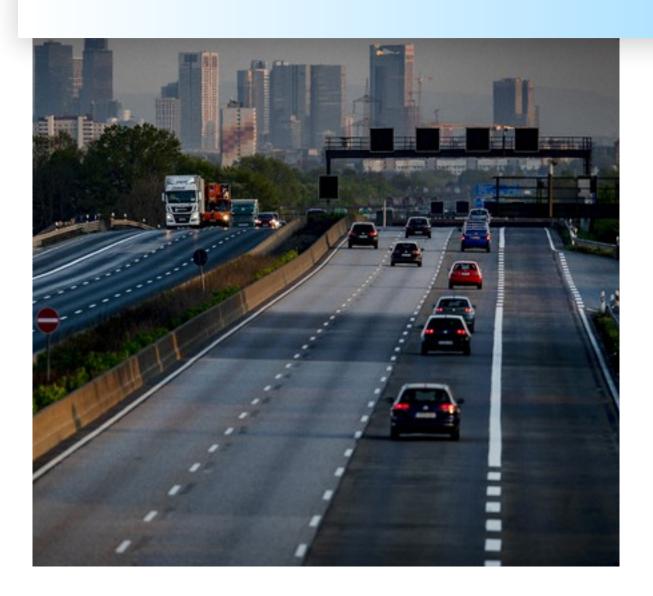
#### Measuring Metabolism

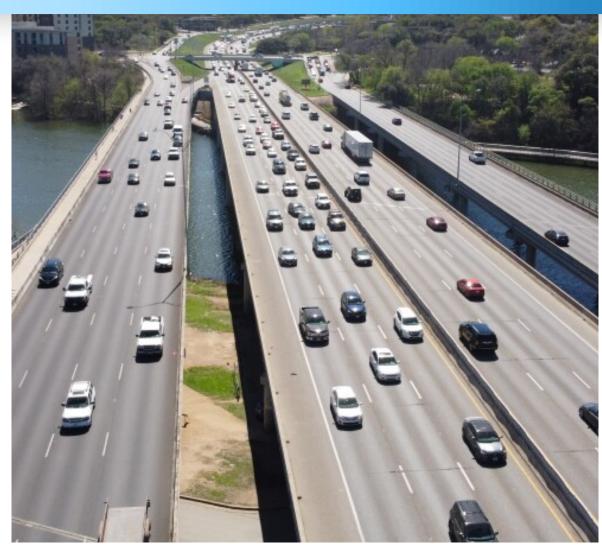
Bulk metabolomics (GC or LC/MS)

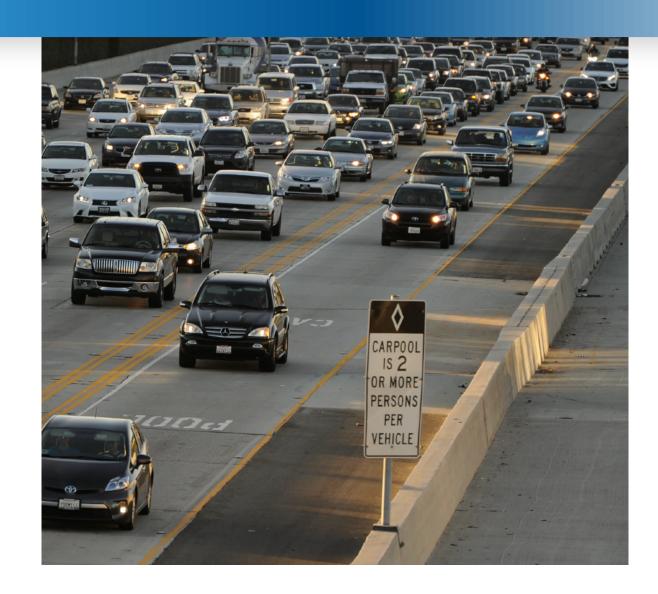


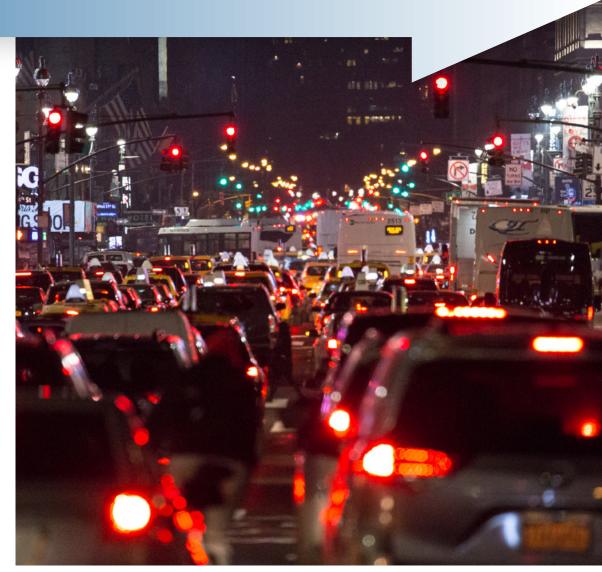
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## What is metabolic flux? (And how to measure)







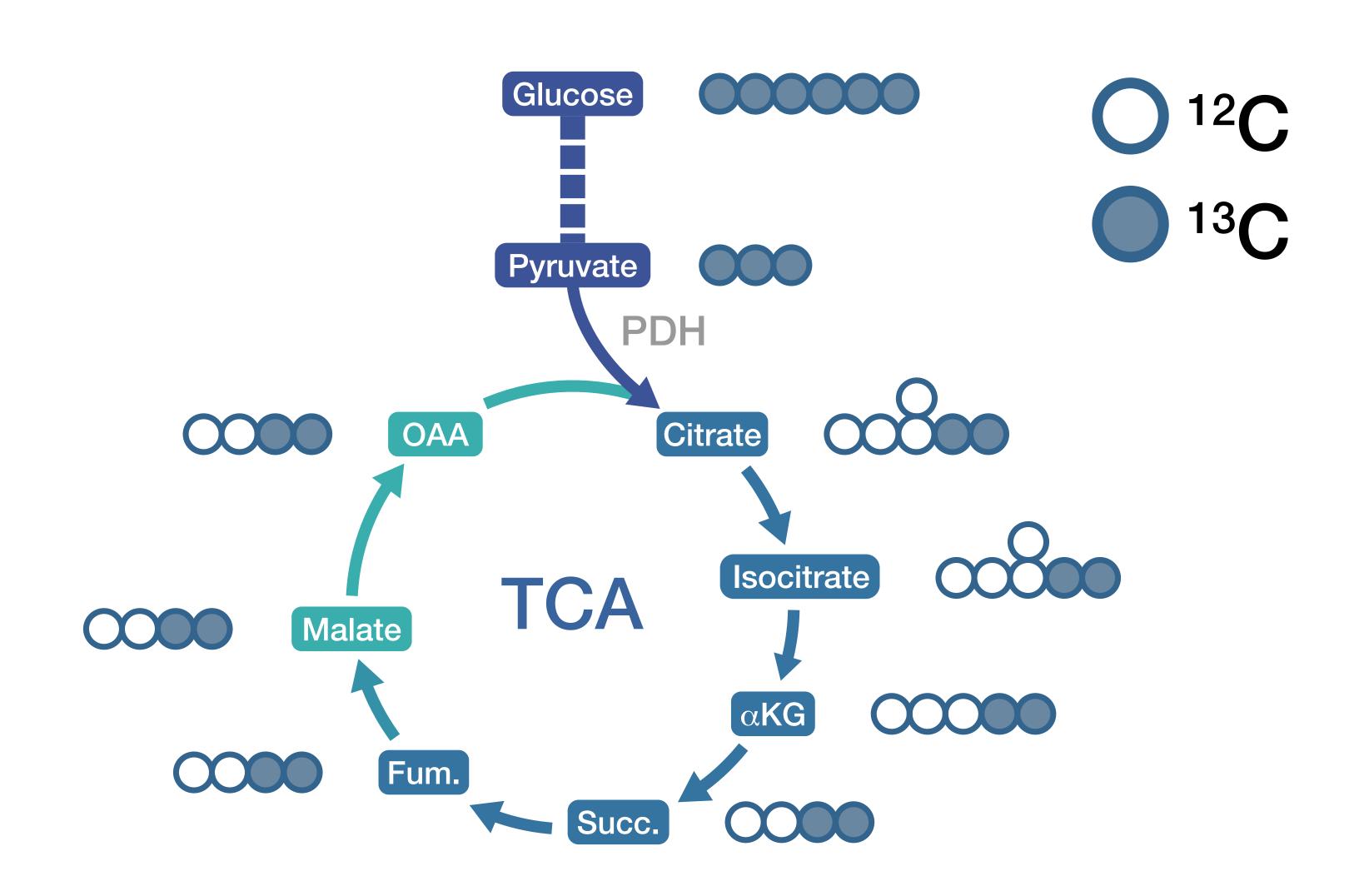


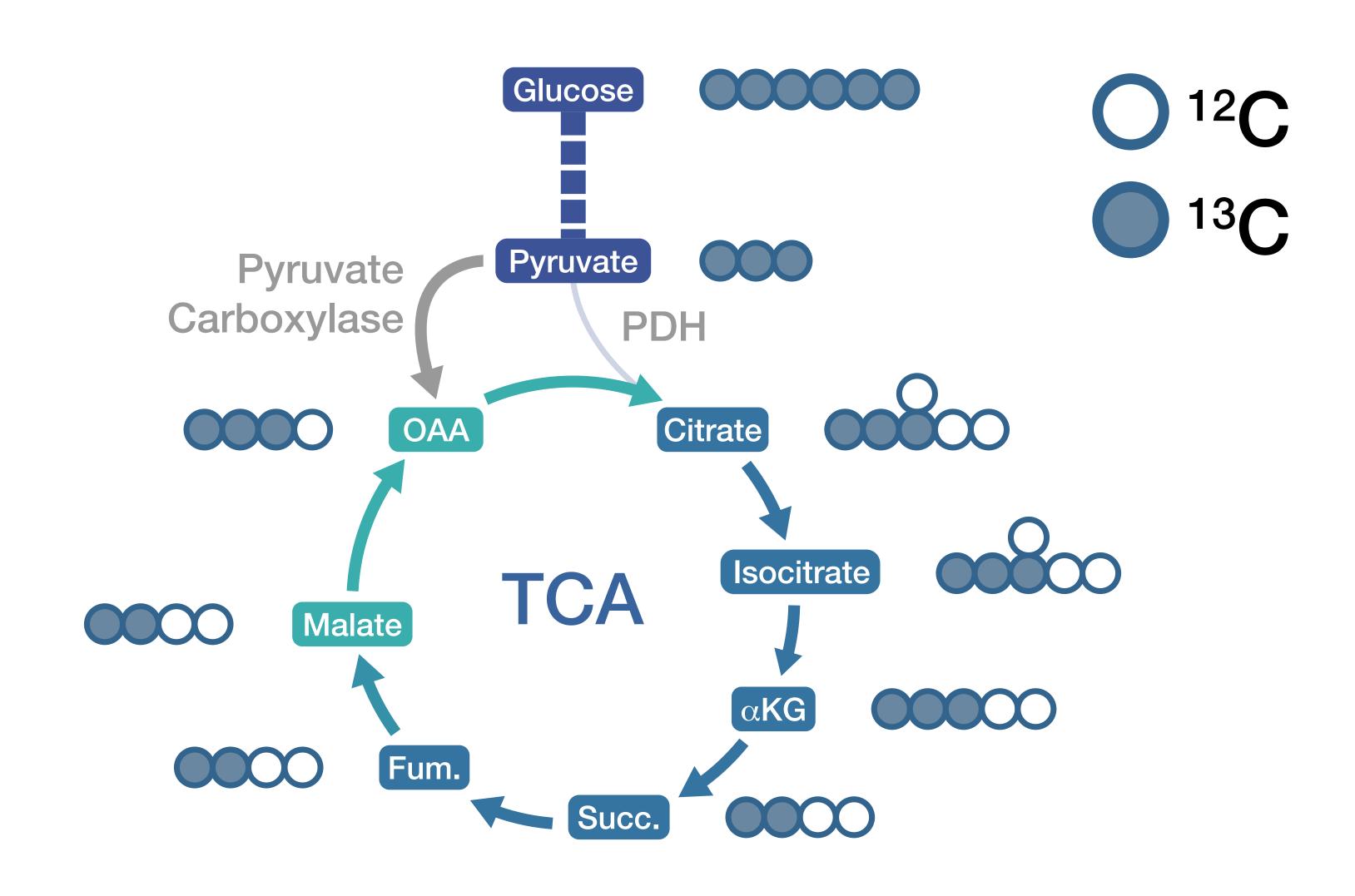


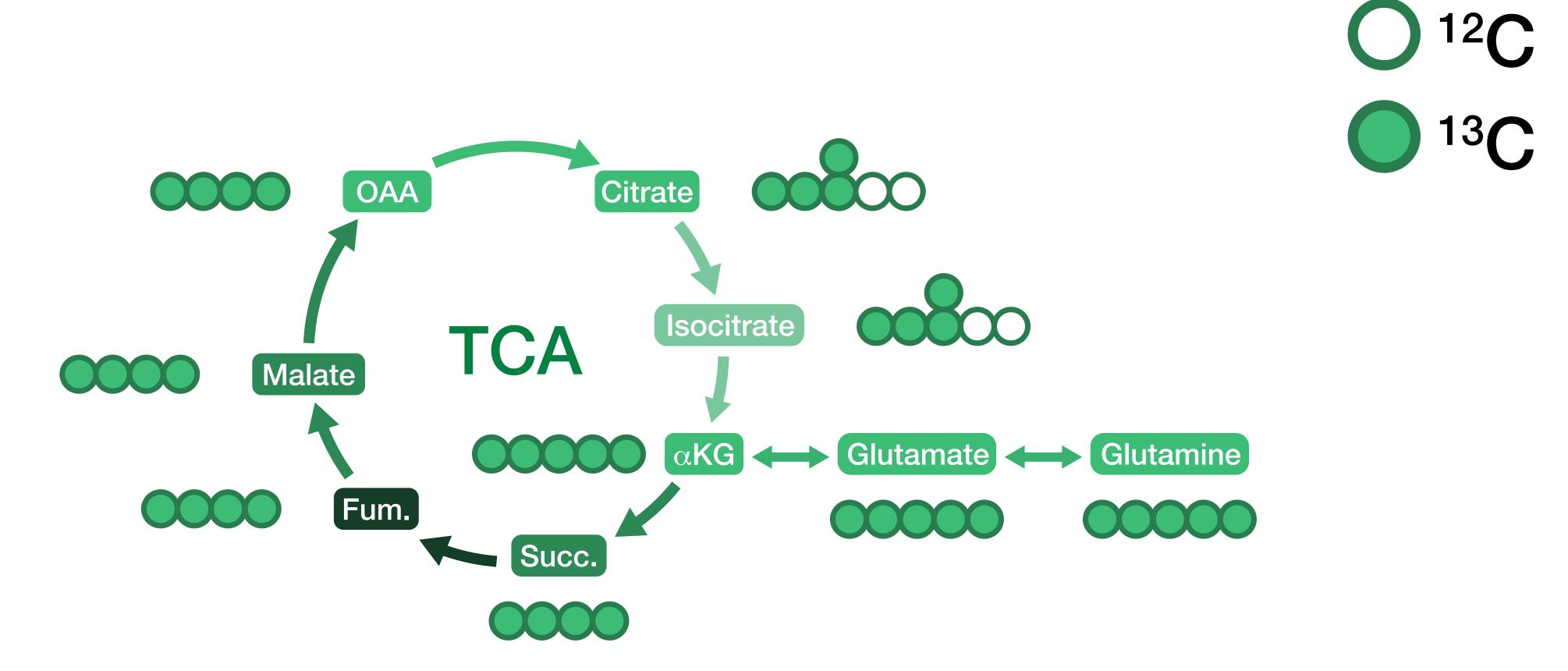


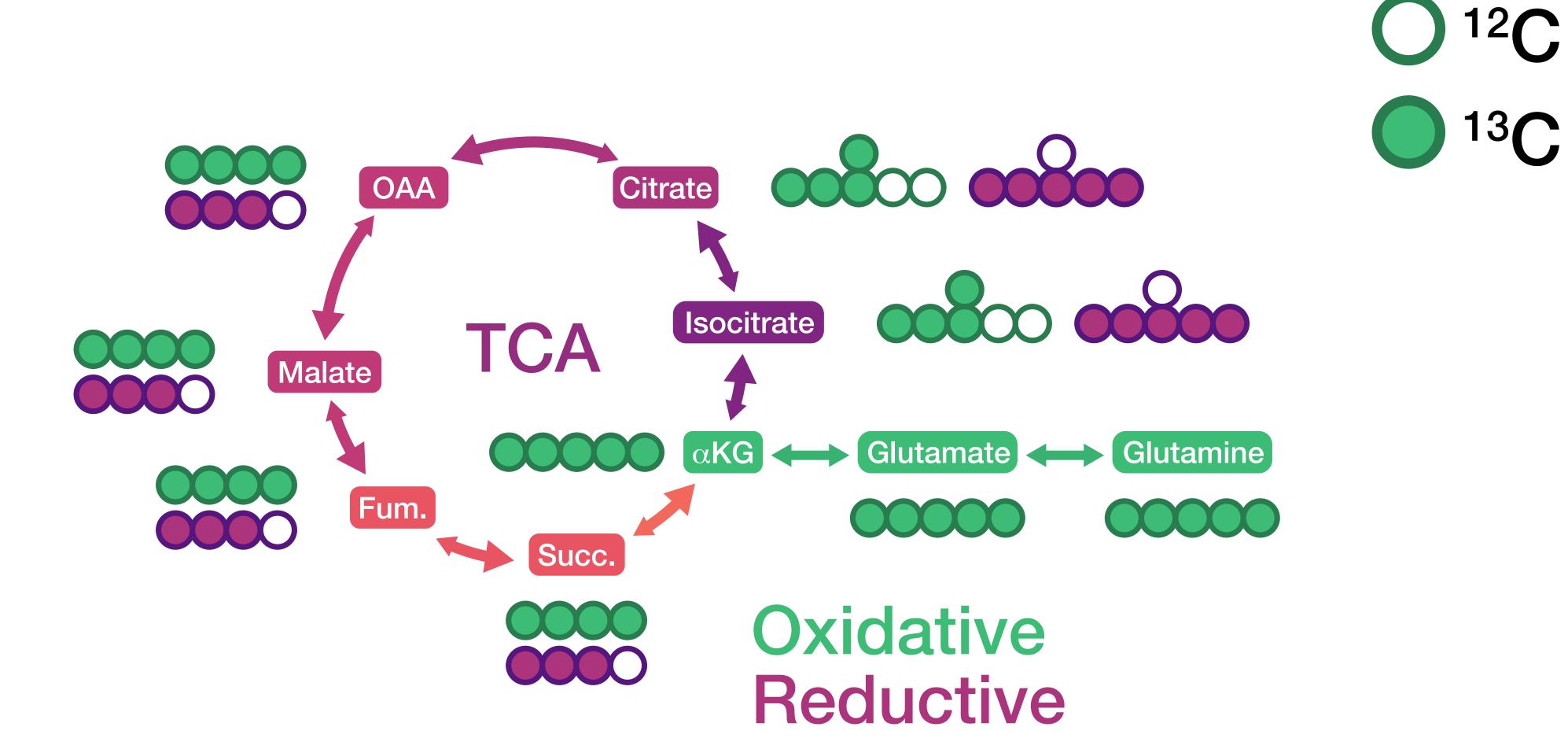




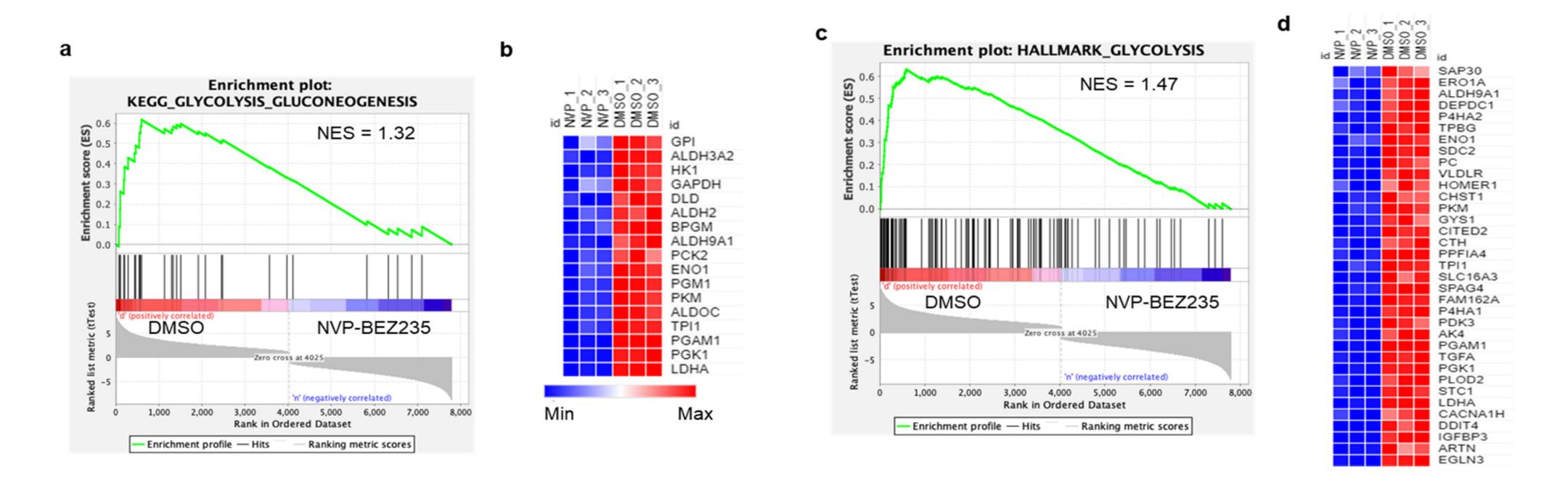








#### Gene expression as a proxy for metabolic flux

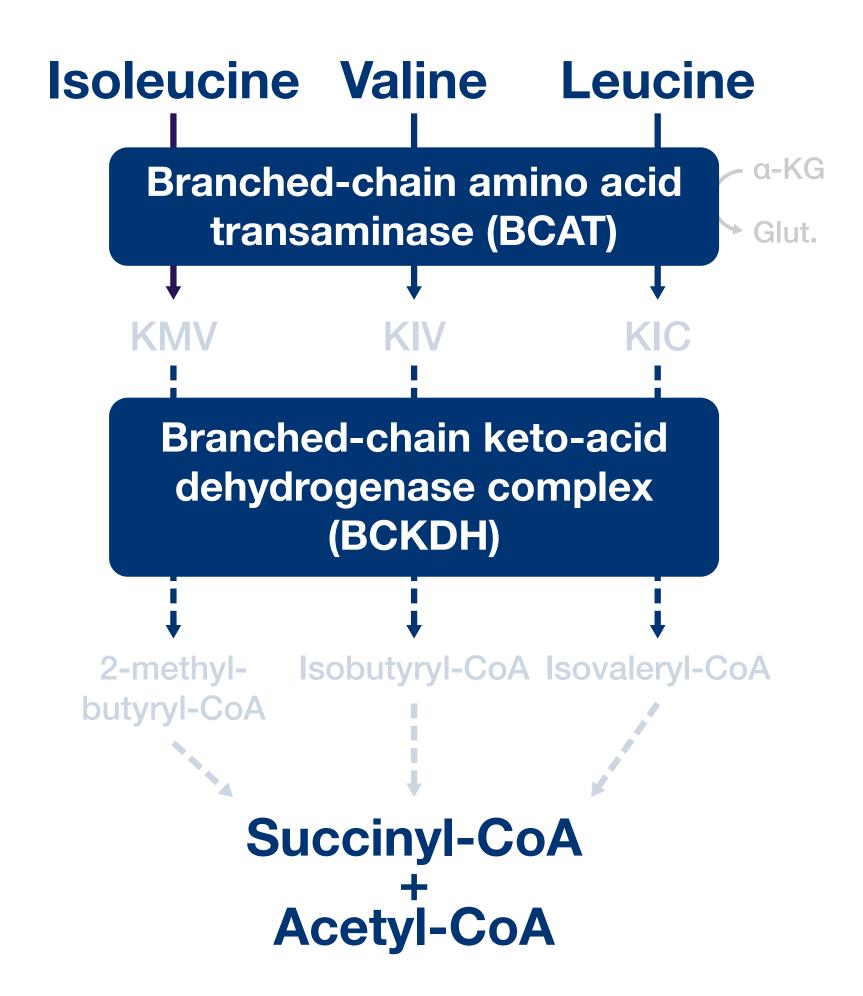


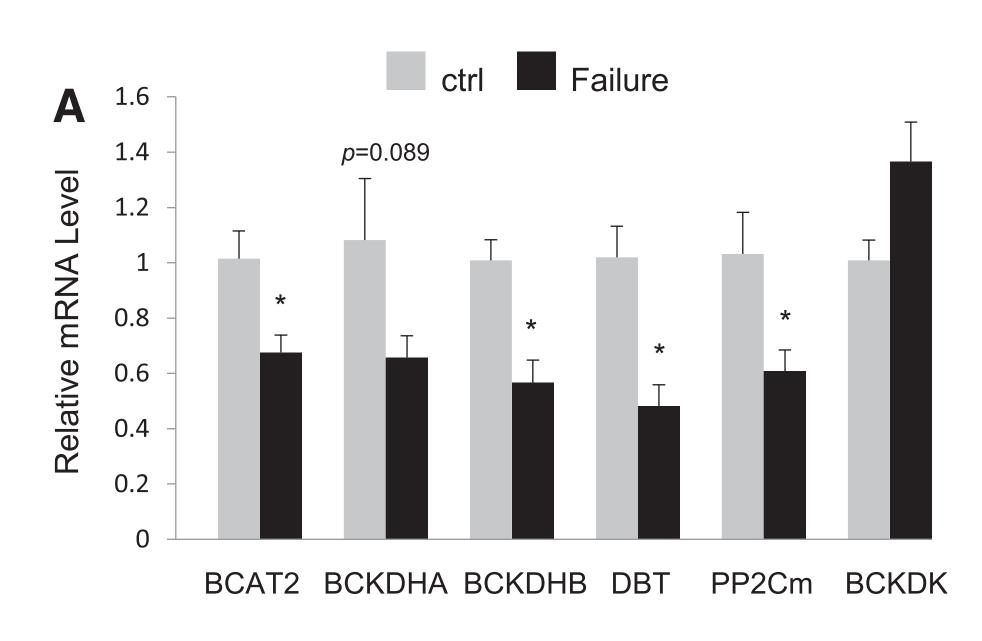
28,501 Metabolism-related Gene-Sets in GSEA Molecular Signatures Database

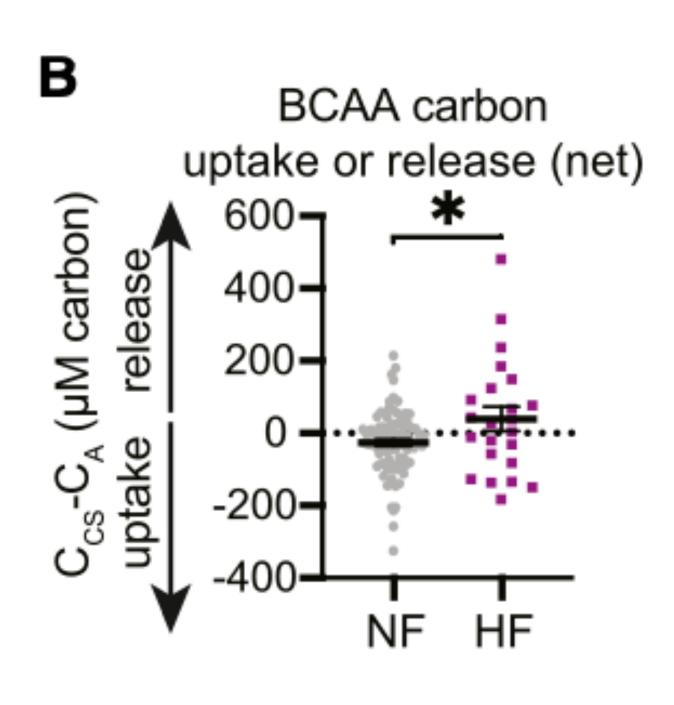


# Metabolic flux CANNOTIS

#### be extrapolated from expression of enzymatic gene loci



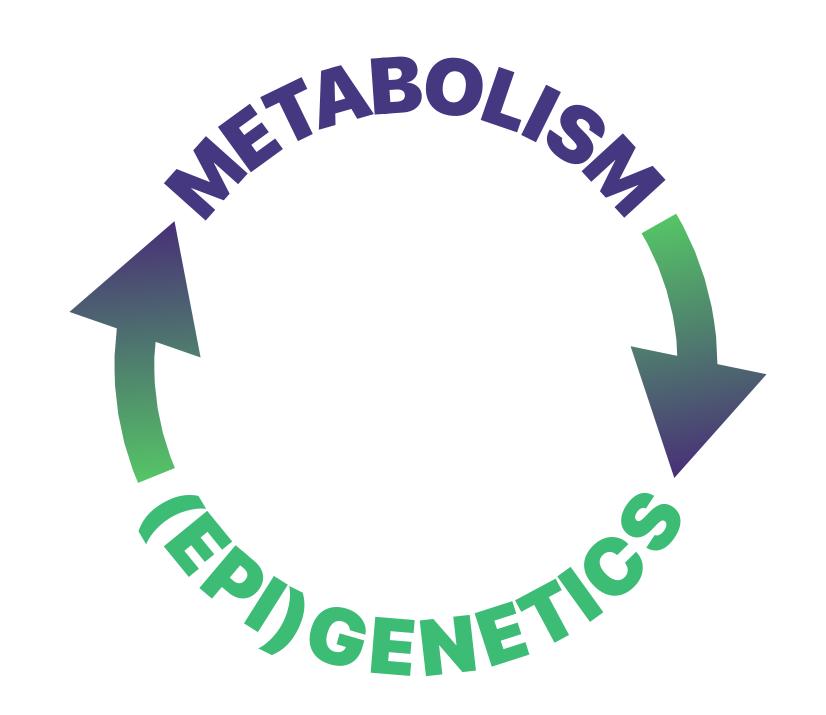




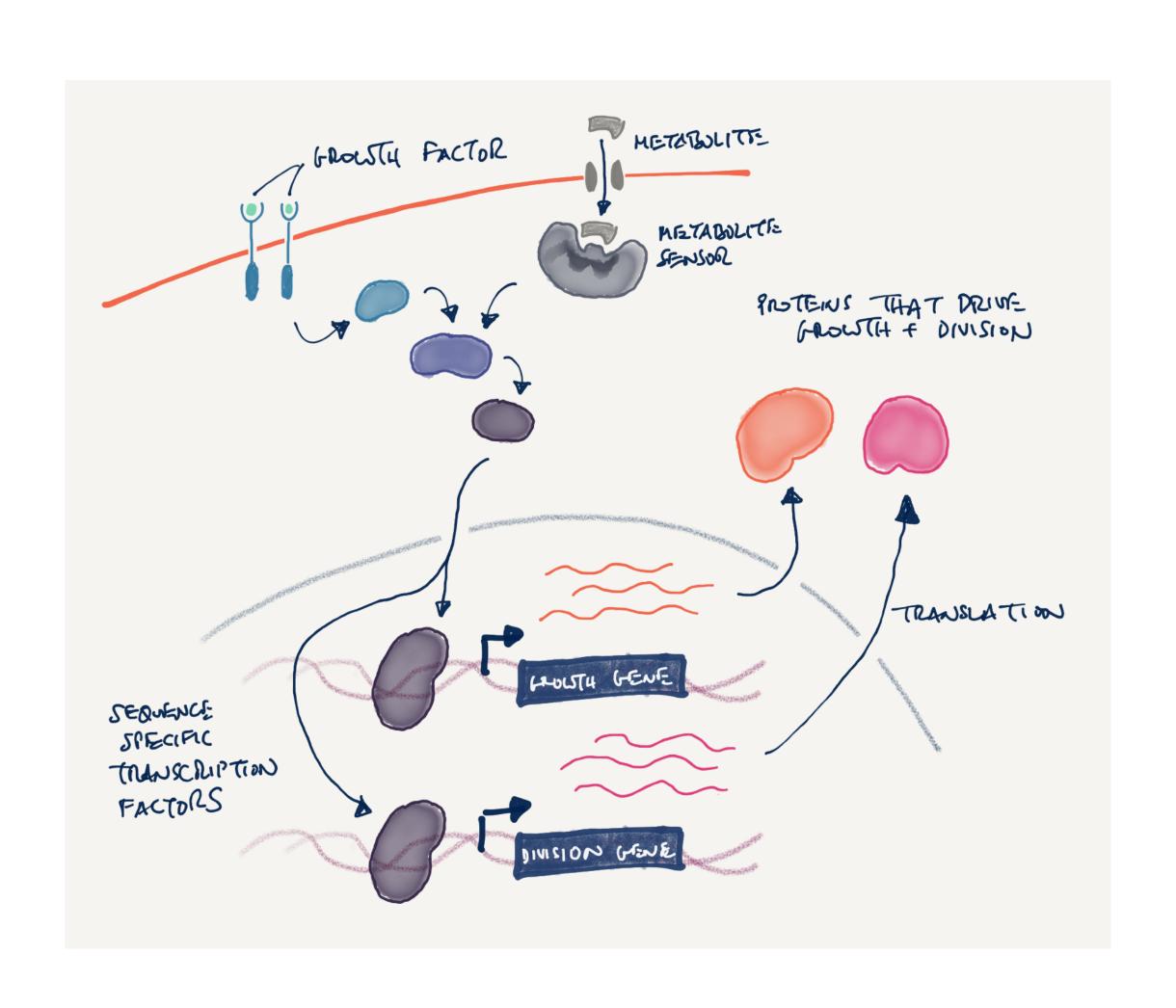
## Takehome points 2

- 1. Forward genetics can be used to study metabolic pathways in auxotrophic systems/organisms
- 2. Inborn errors of metabolism are the clinical manifestations of blocks in metabolic flux
- 3. Metabolomics shows us the snapshot AMOUNT of each metabolite in a population
- 4. Flux analysis reveals directionality and the contribution of one metabolite to different pathways
- 5. Metabolic flux CANNOT be extrapolated from gene expression data

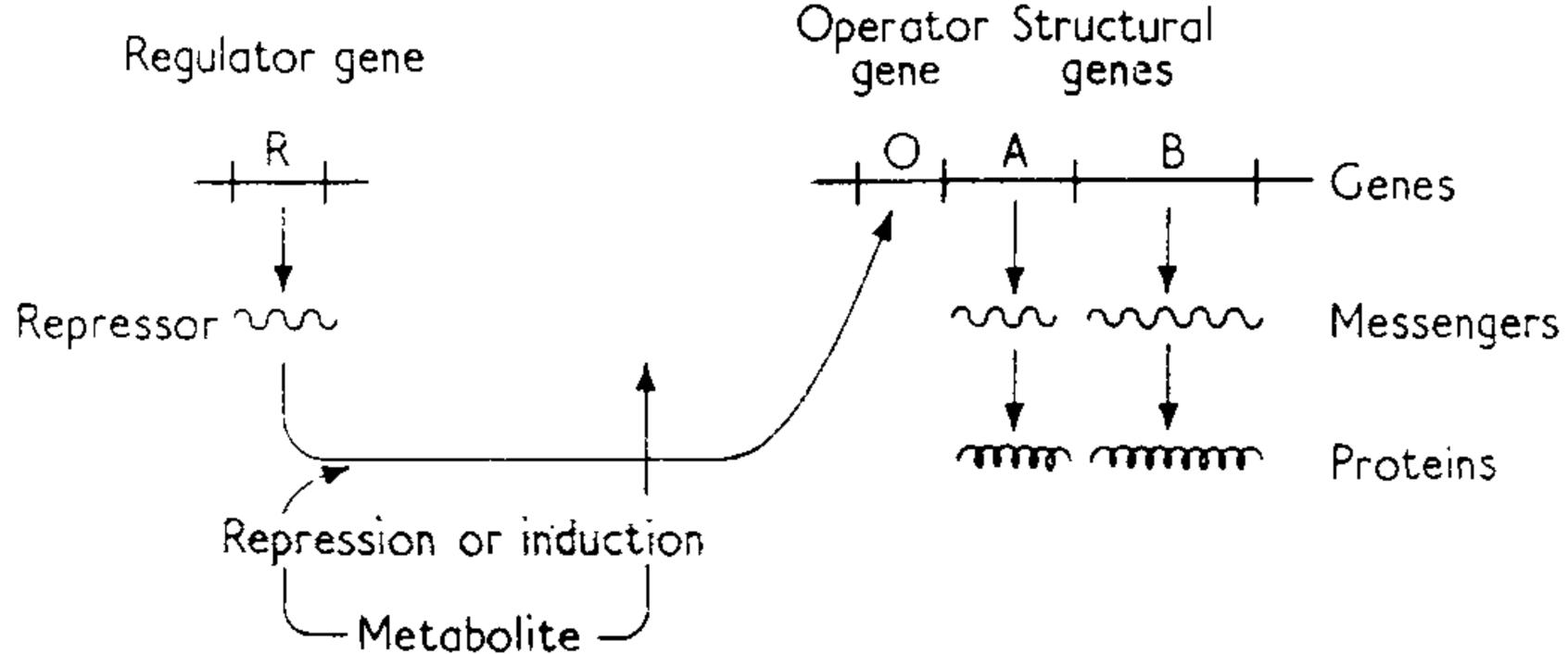
#### A DIVE INTO HOW METABOLISM AFFECTS CHROMATIN

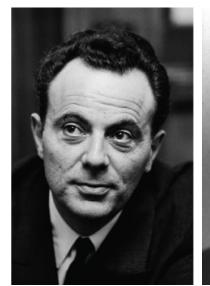


# The classic view of transcriptional regulation hinges on DNA-sequence specificity of activators and repressors



#### Bacterial gene-expression is DIRECTLY regulated by metabolites



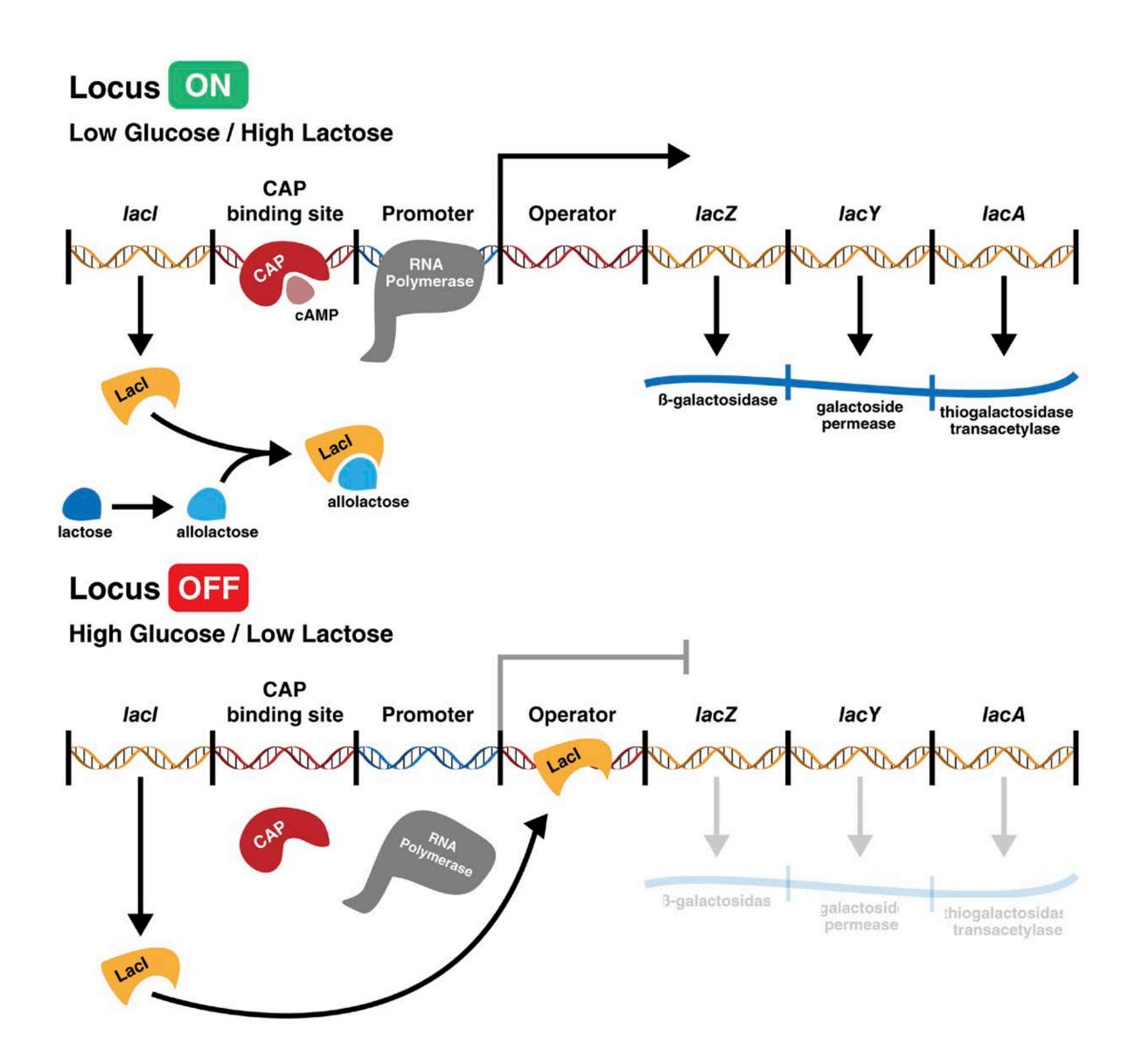






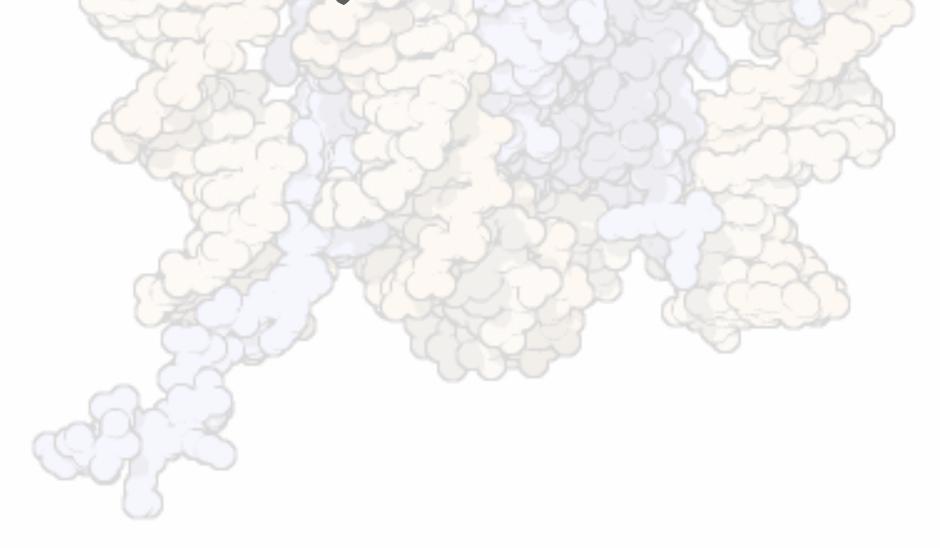
François Jacob Jacques Monod

#### Bacterial gene-expression is DIRECTLY regulated by metabolites



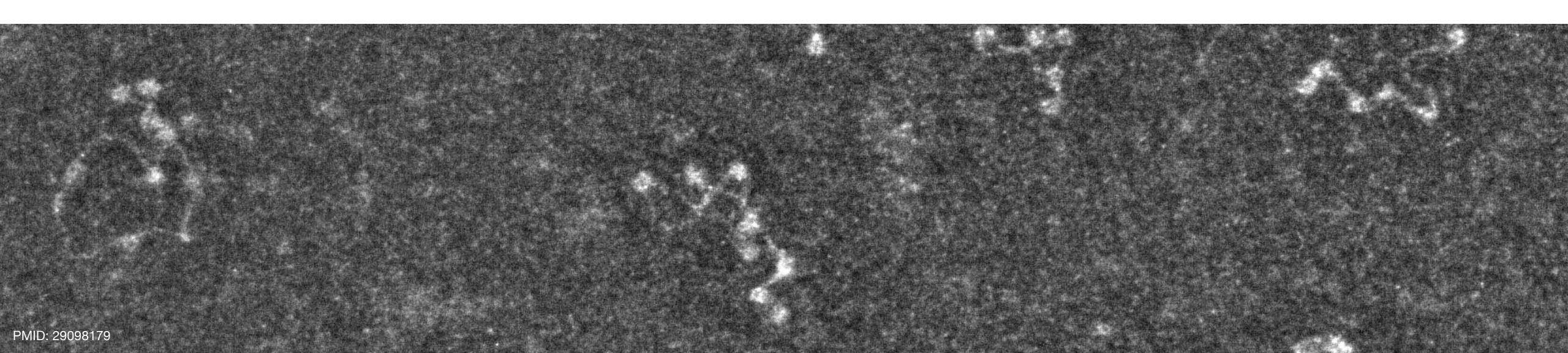


OK BUT THAT'S PROKARYOTES, NO NUCLEUS AND NO CHROMATIN...



#### Chromatin is heavily decorated with the products of metabolic reactions

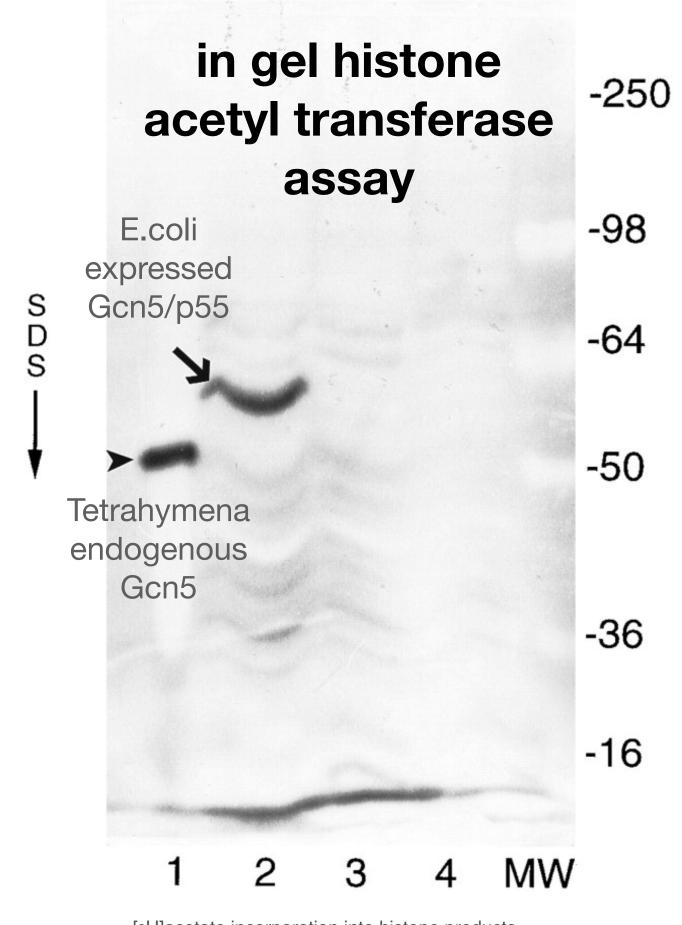




#### David Allis

Tetrahymena thermophila

#### So why are histones acetylated?

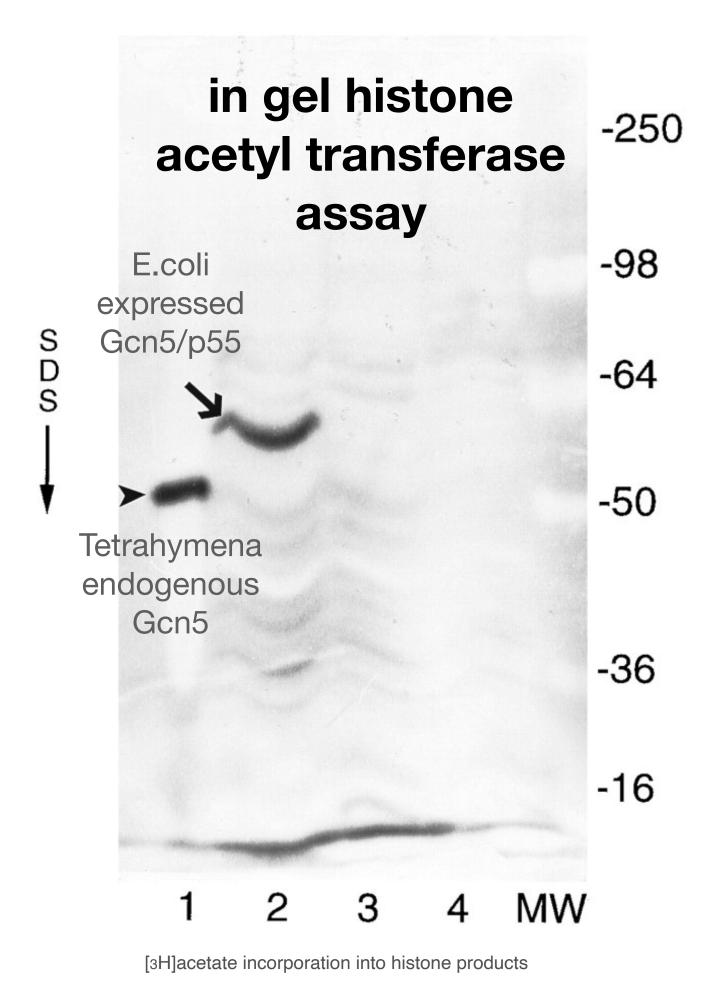


#### David Allis

# The state of the s

Tetrahymena thermophila

#### So why are histones acetylated?



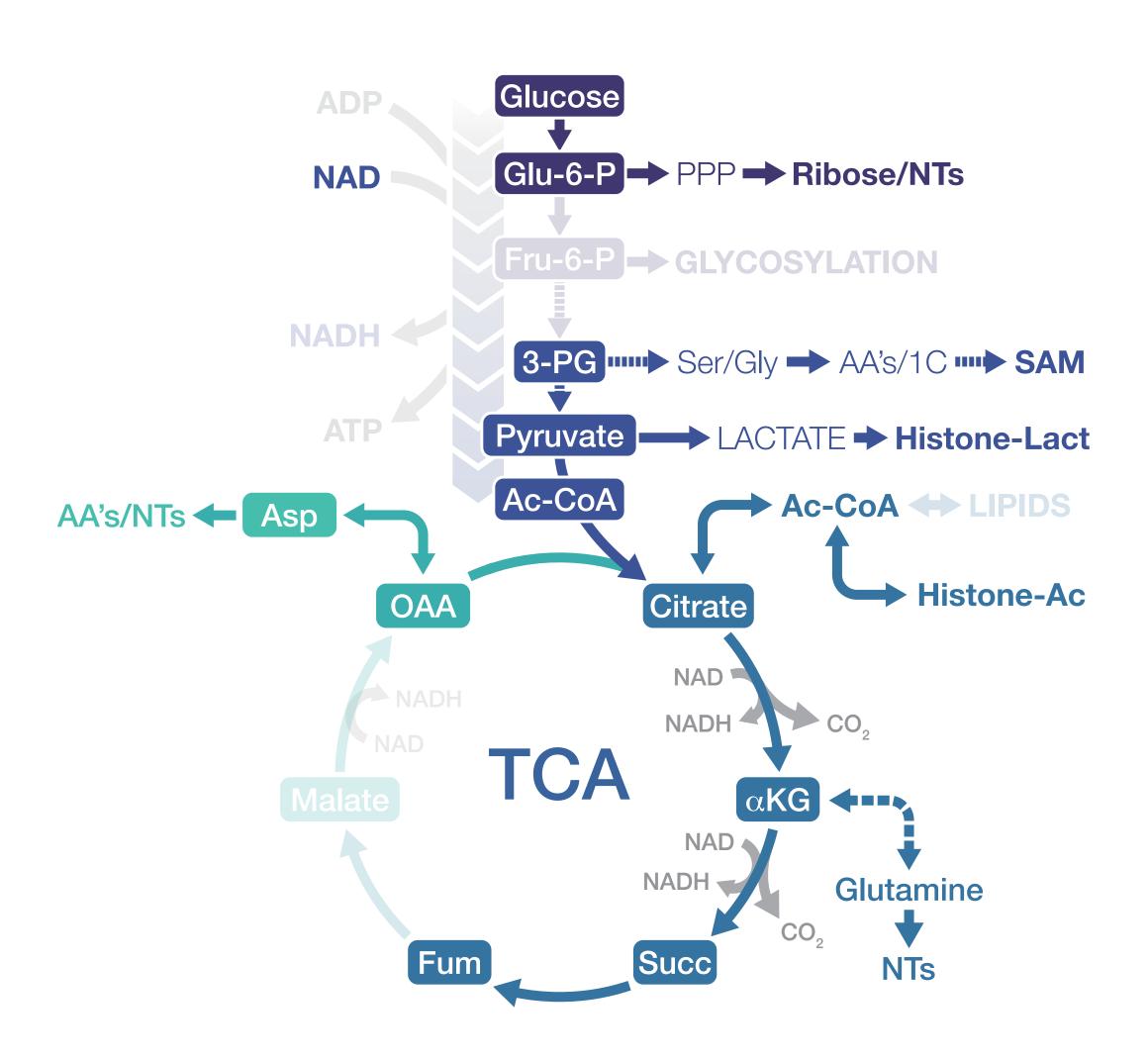
## METABOLITE-DEPENDENT ENZYMATIC MODIFICATION OF CHROMATIN CAN CHANGE GENE EXPRESSION



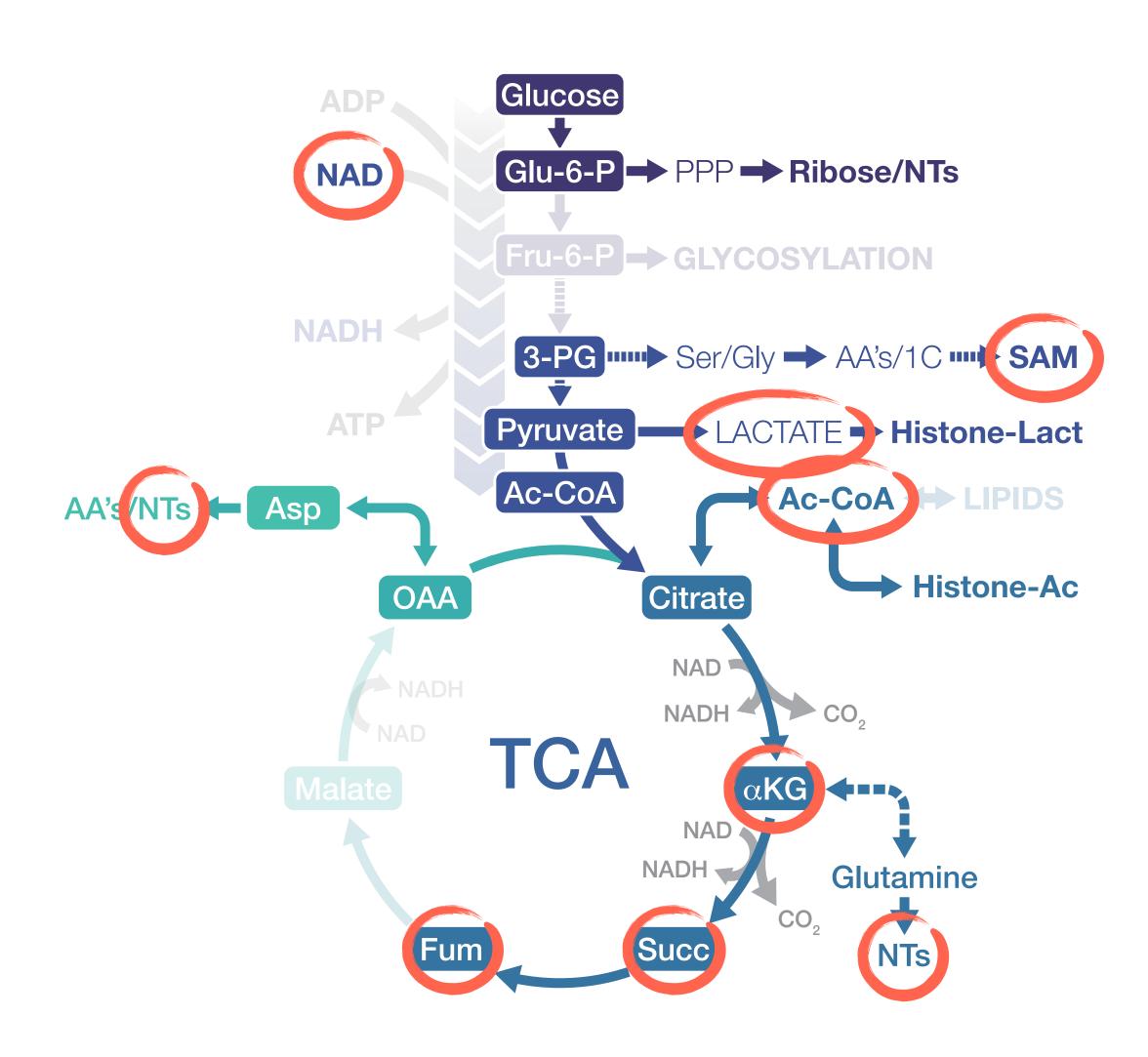
#### Implications:

- Changes in intracellular metabolism affect gene expression
- Gene expression programs can be altered through inhibition of the metabolic pathways they are dependent on
  - i.e. TFS CAN'T BE DRUGGED BUT ENZYMES CAN

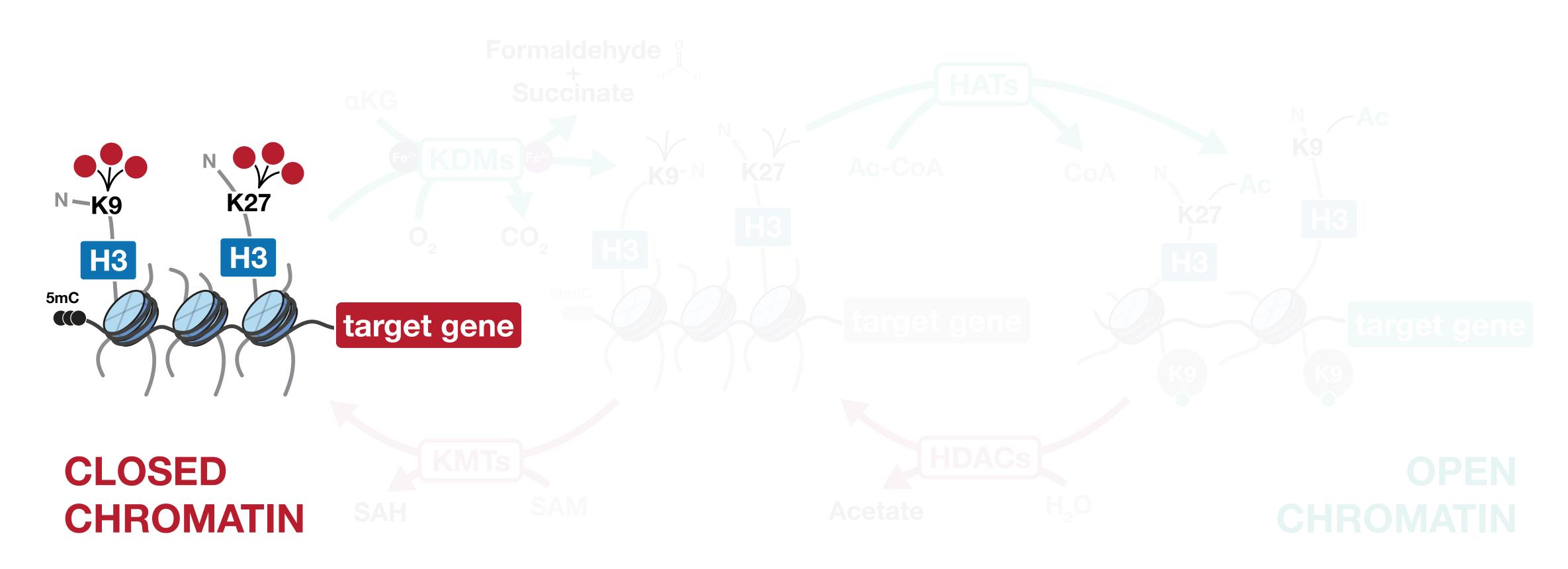
#### Central Carbon Metabolism: Links to Chromatin



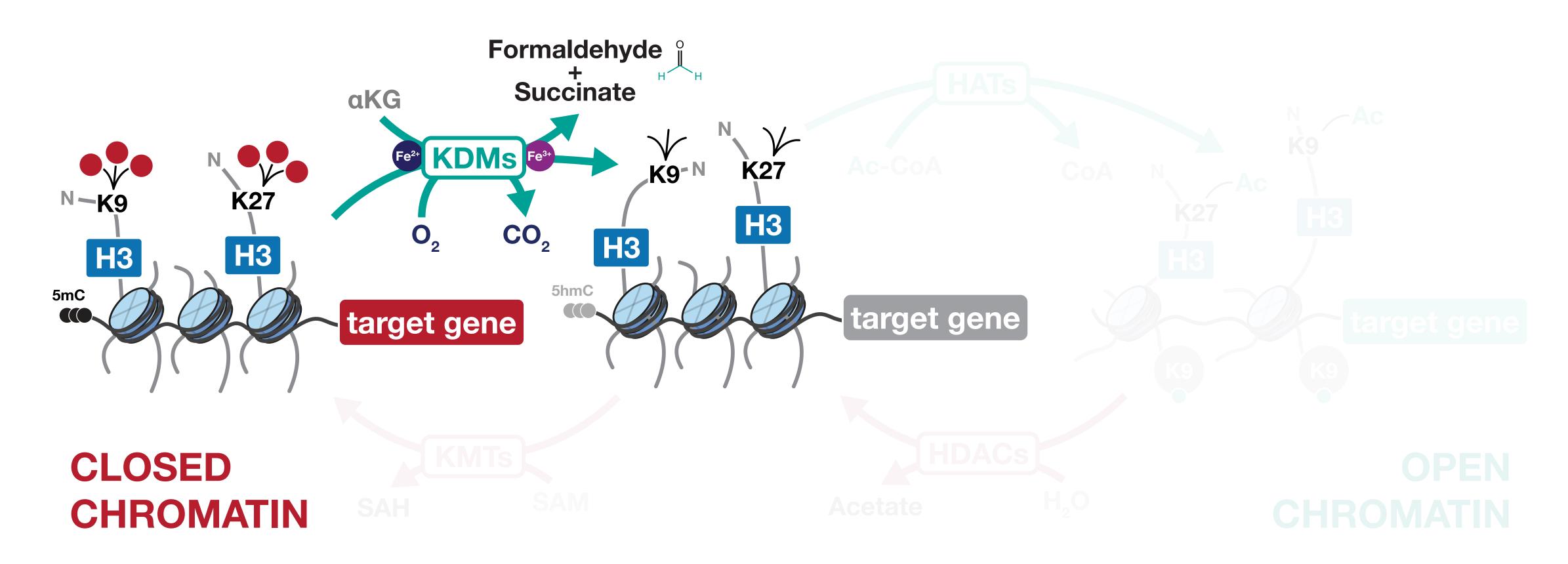
#### Central Carbon Metabolism: Links to Chromatin



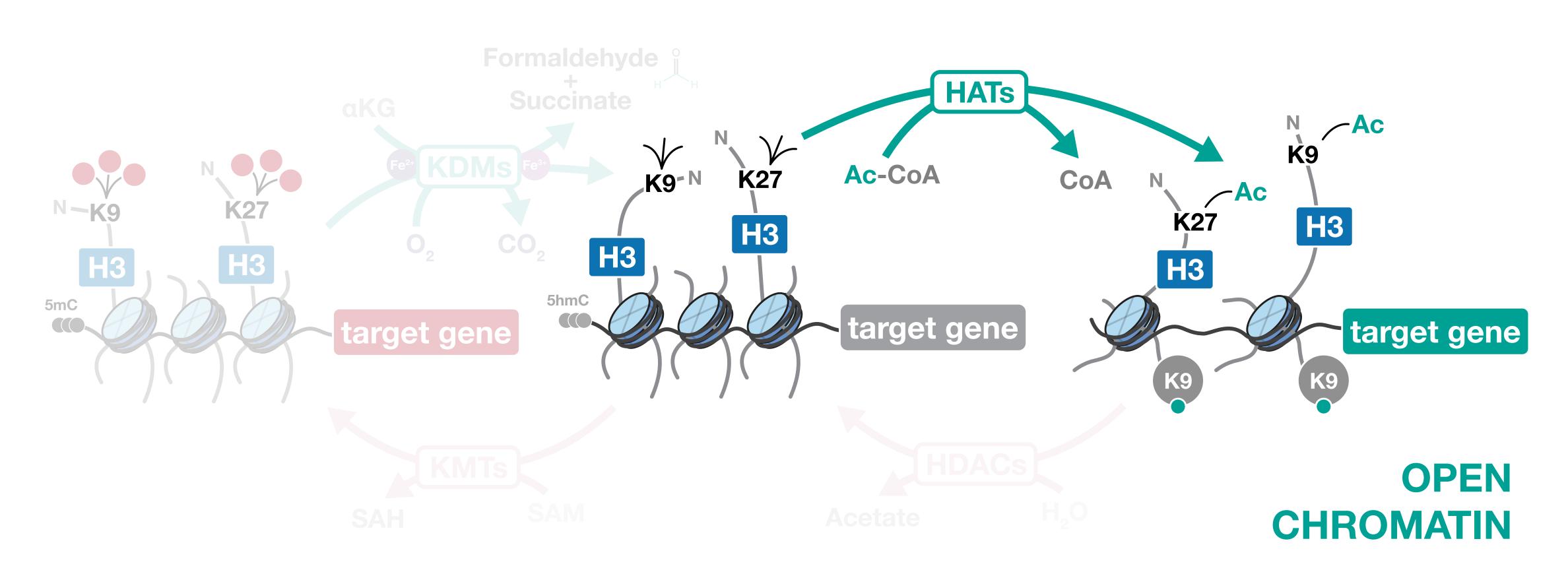




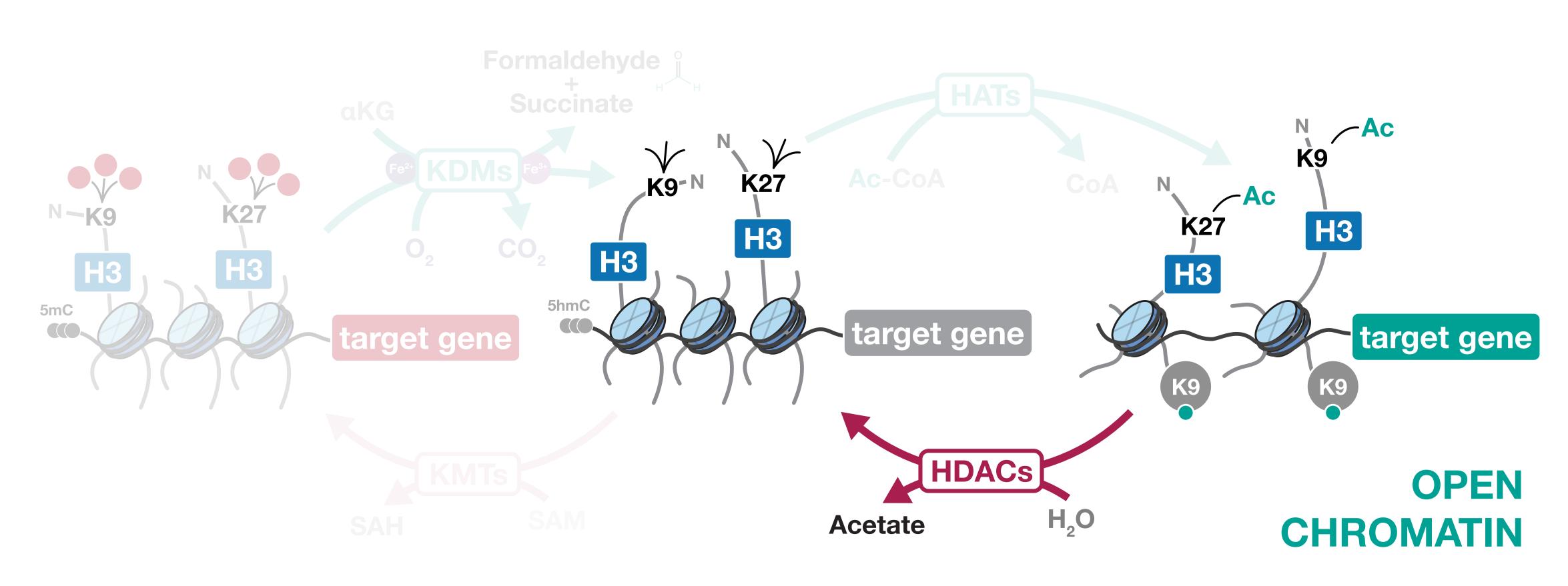




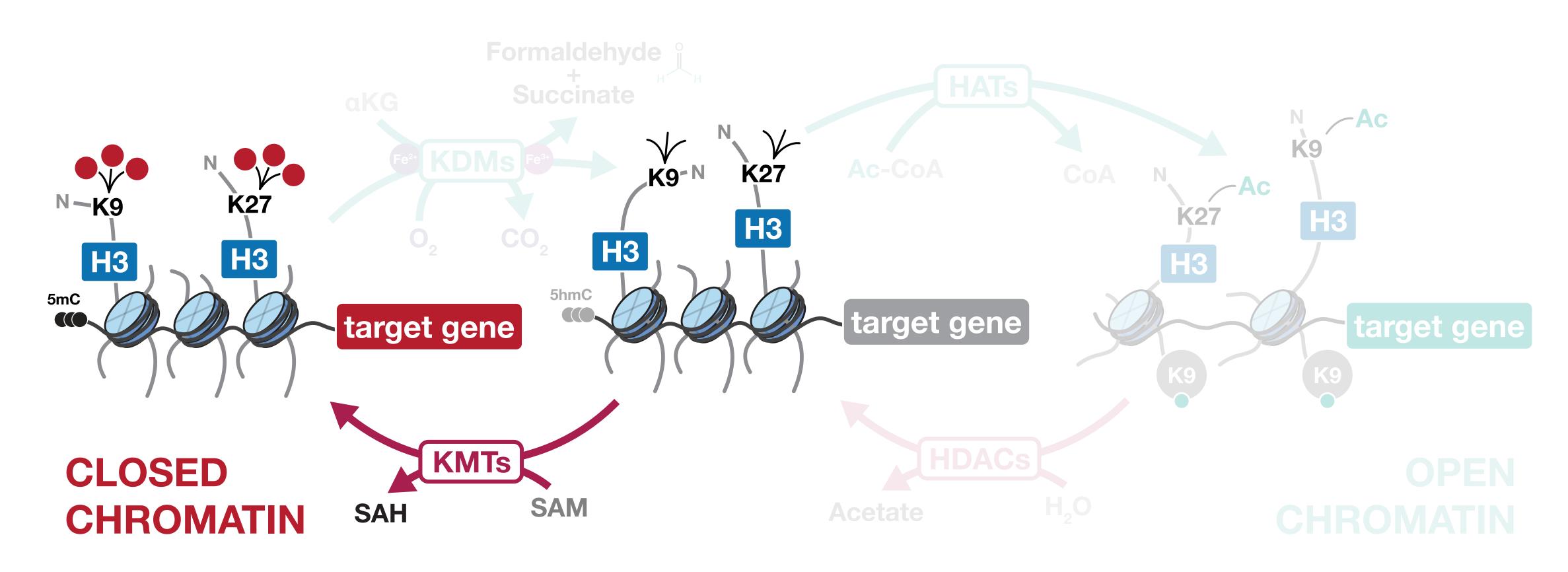




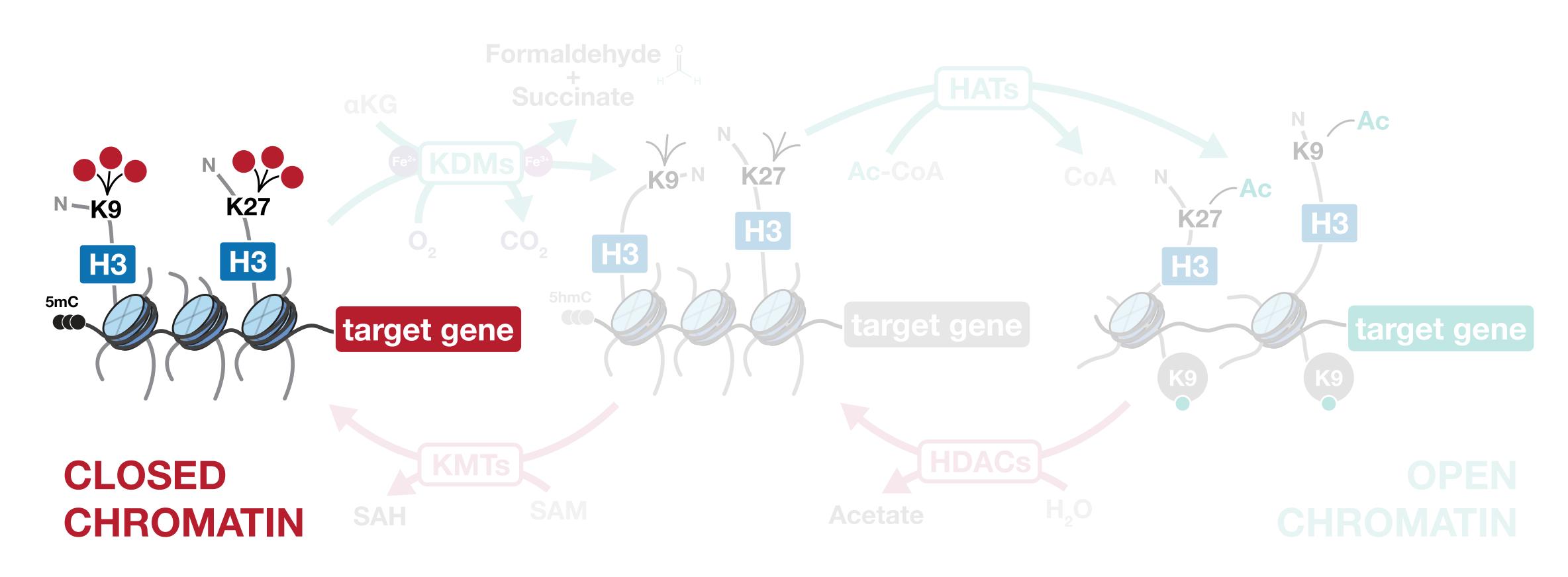








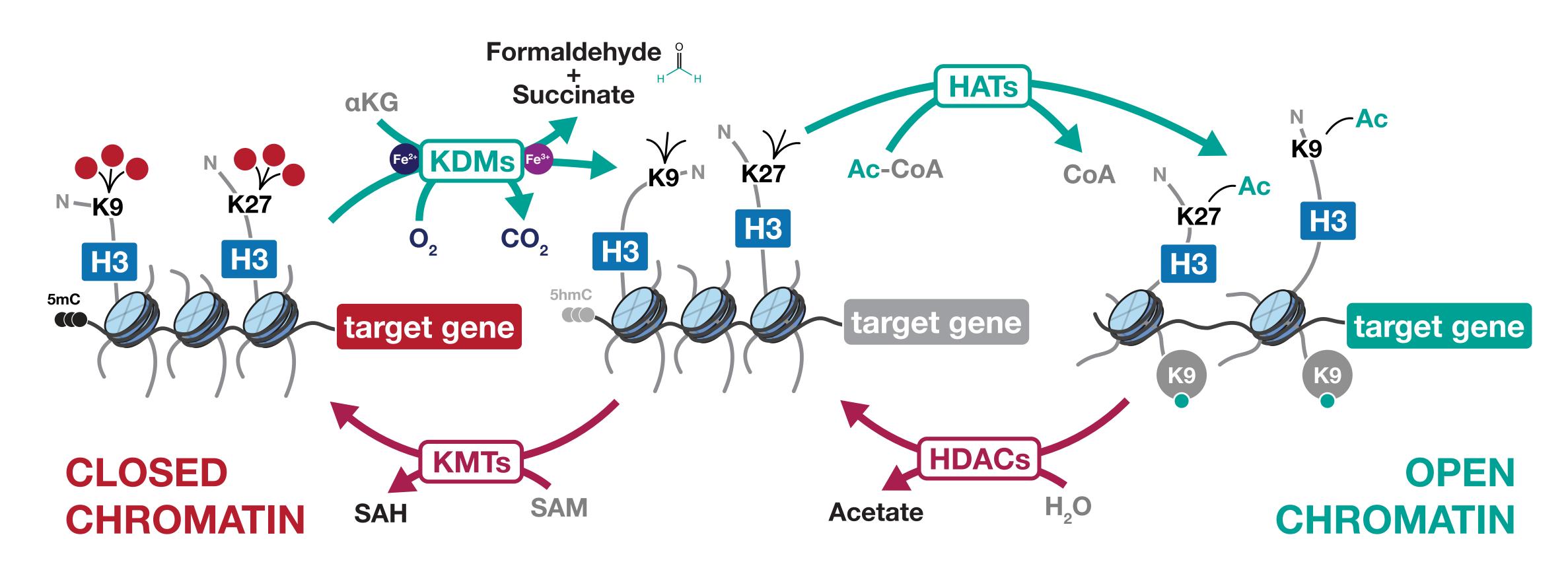




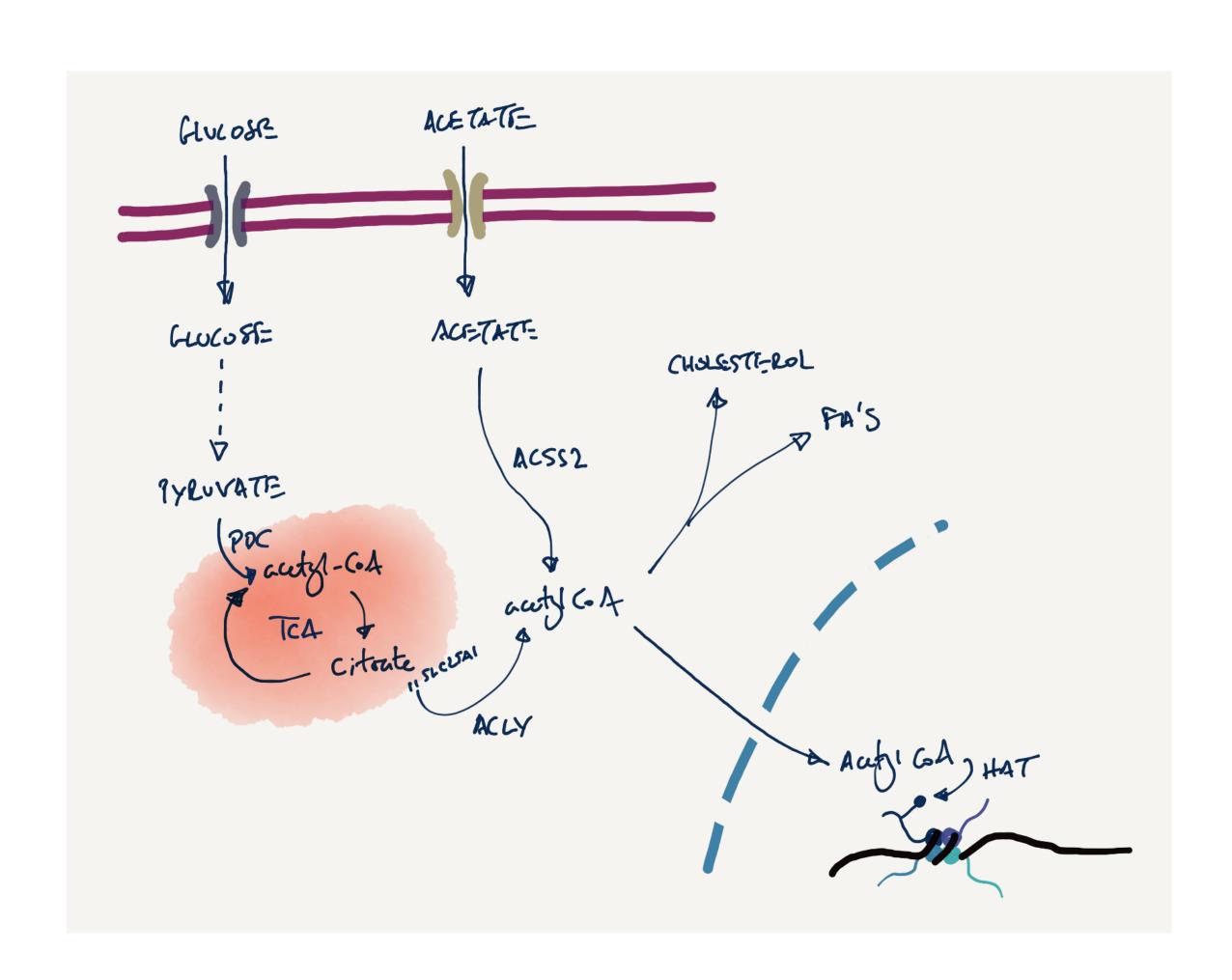
#### Chromatin modifications respond to the metabolic state of the cell

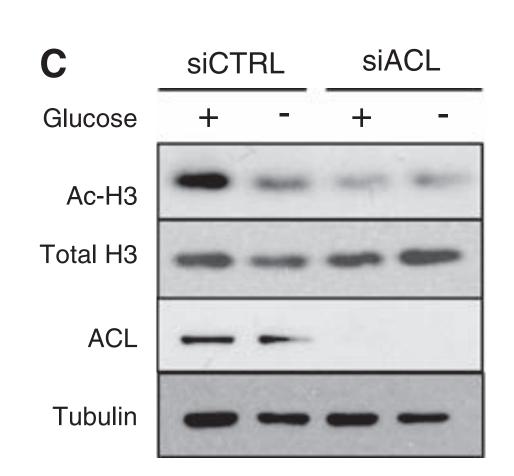






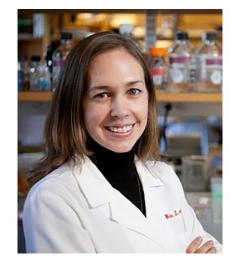
## Does acetyl-CoA concentration affect histone acetylation / gene expression?

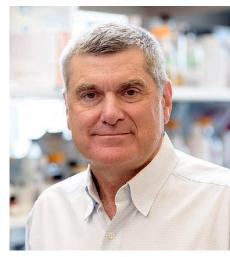




Adipocyte Differentiation (Oil Red O)



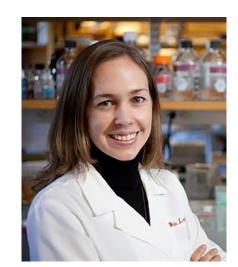


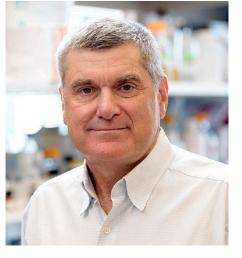


Katy Wellen Craig Thompson

## Does acetyl-CoA concentration affect histone acetylation / gene expression?

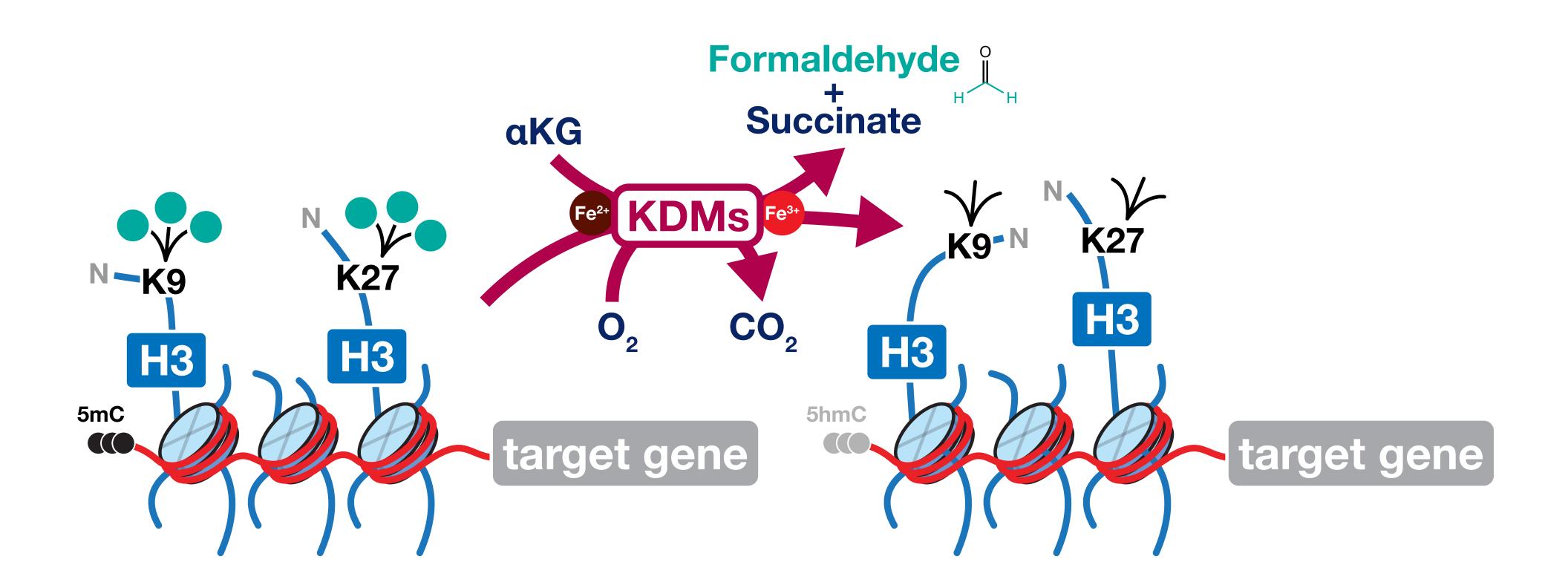
# [SUBSTRATES] MATTER!!!!



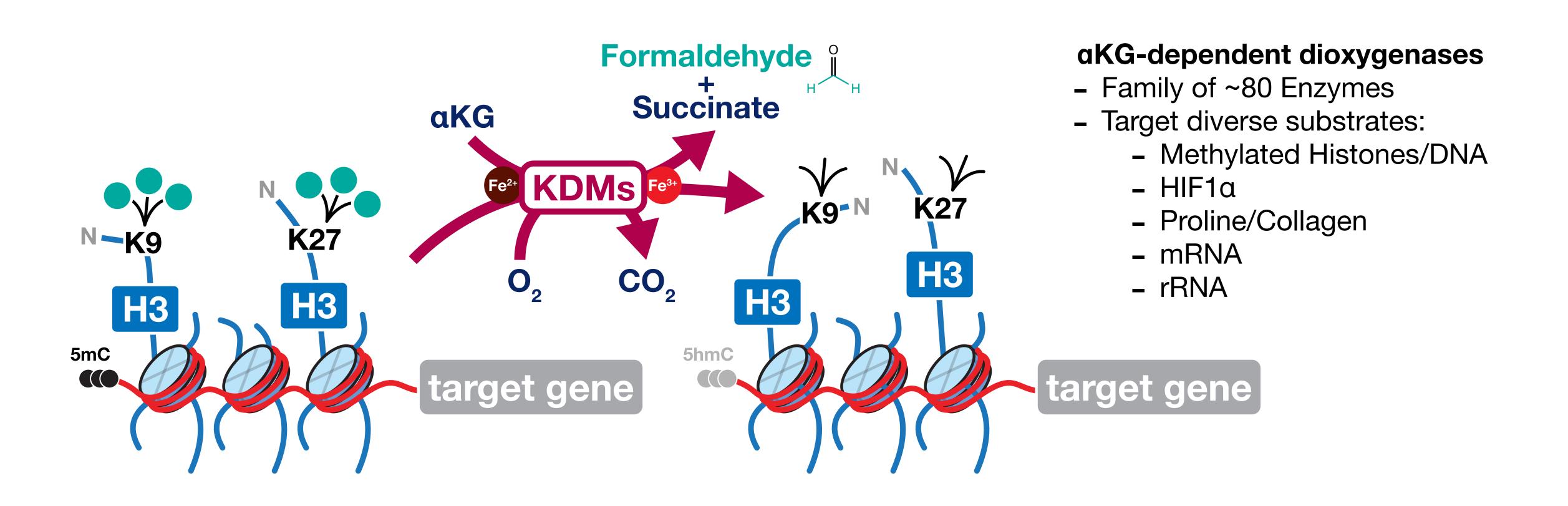


Katy Wellen Craig Thompson

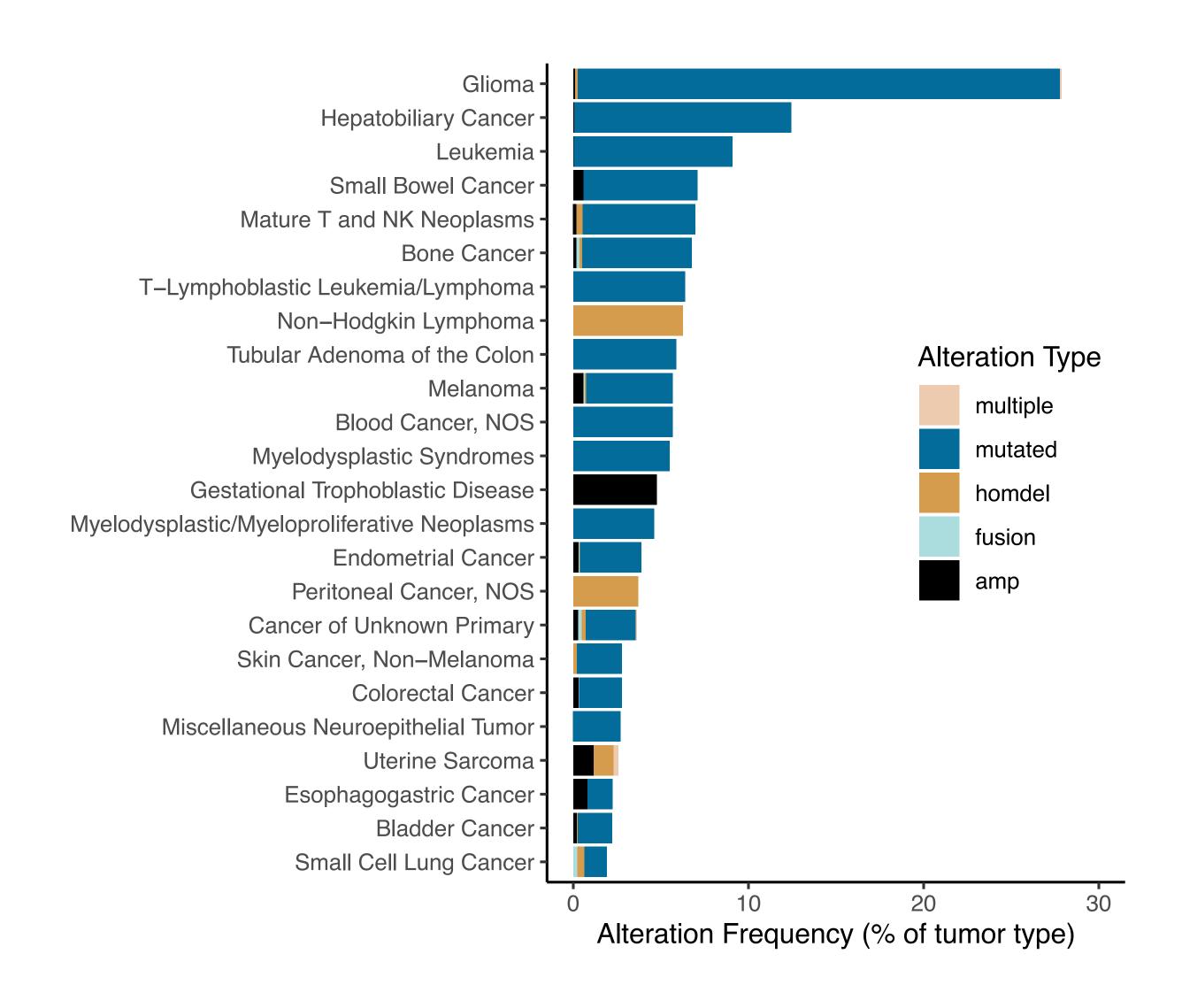
### aKG-dependent dioxygenases remove methyl marks from methylated histones



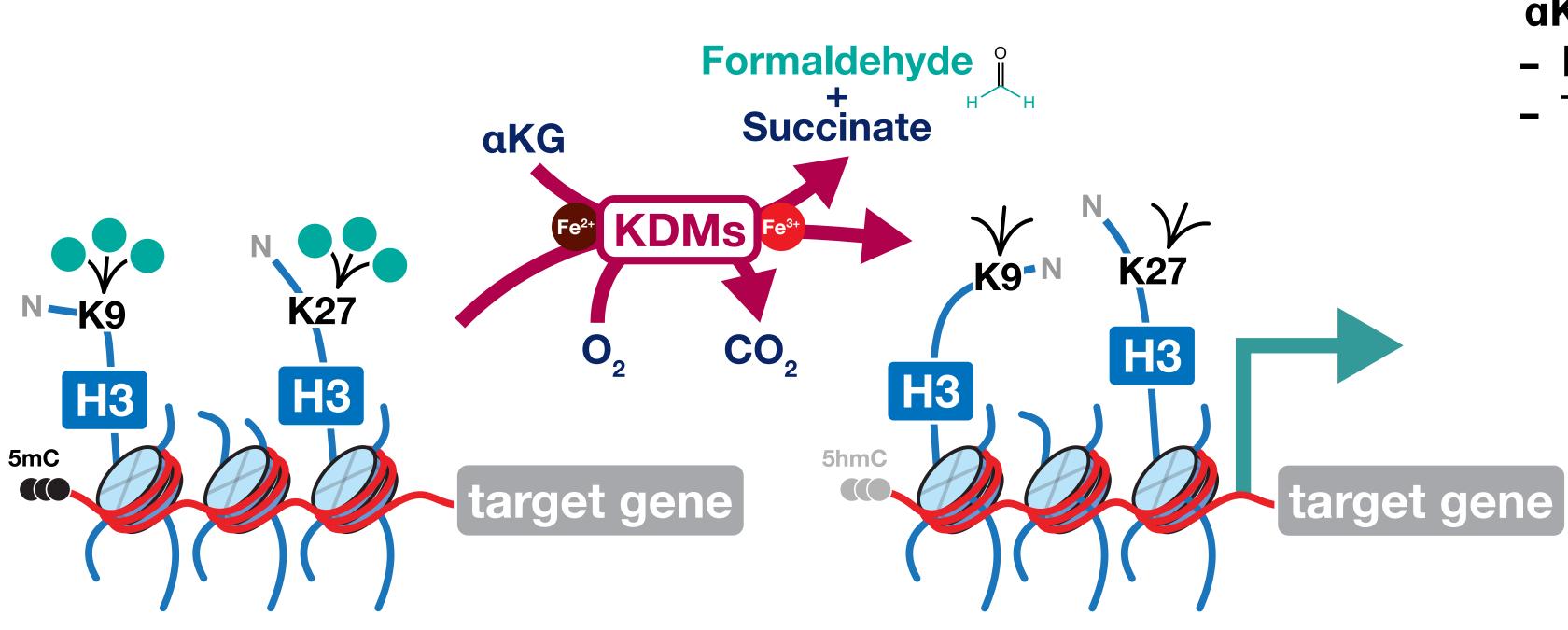
### aKG-dependent dioxygenases remove methyl marks from methylated histones



### Isocitrate Dehydrogenase 1/2 (IDH1/2) neomorphic mutations are found in a diverse set of malignancies derived from all germ layers



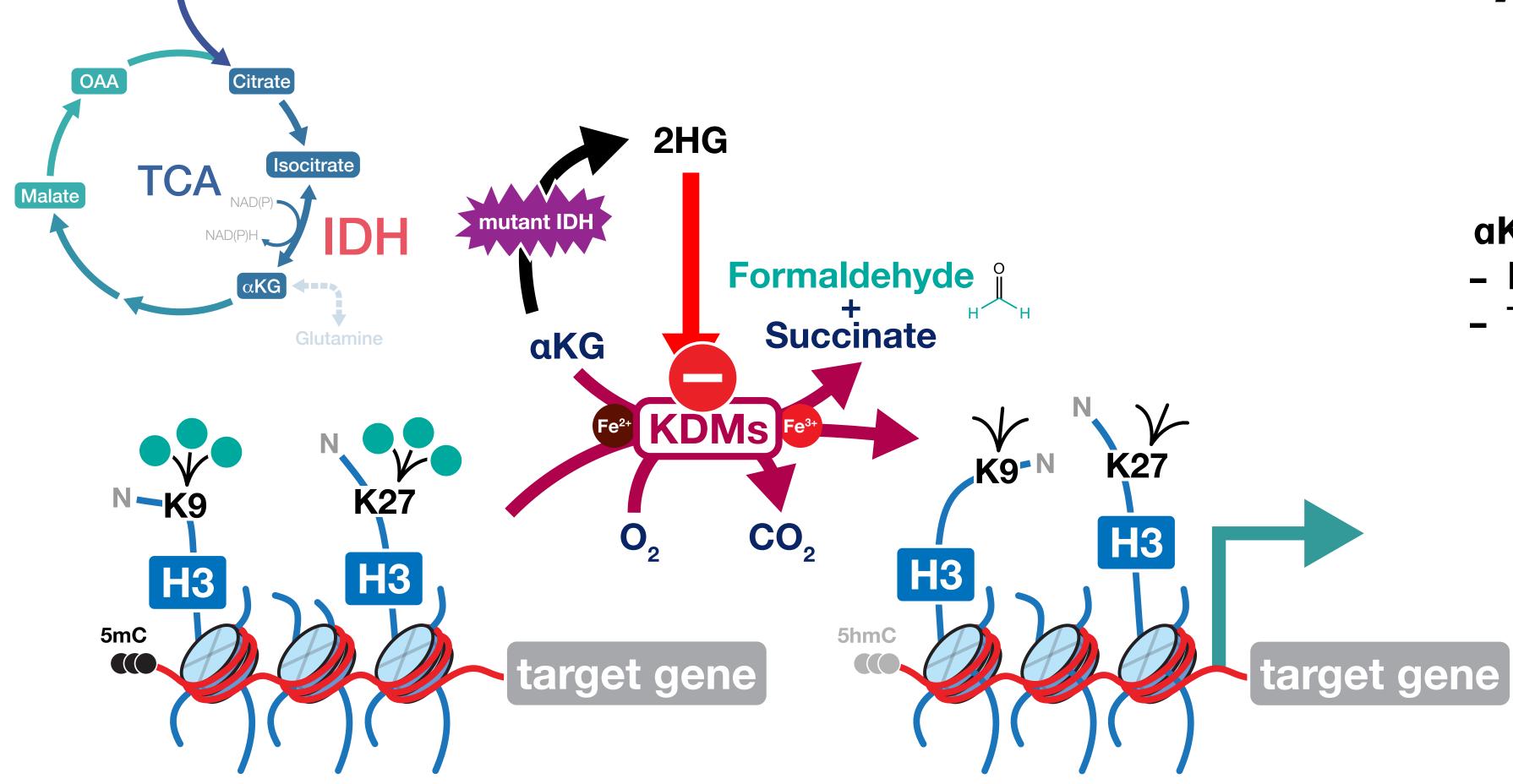
# IDH mutations block differentiation by impairing the removal of heterochromatic H3K9me2/3 methyl marks



#### aKG-dependent dioxygenases

- Family of ~80 Enzymes
- Target diverse substrates:
  - Methylated Histones/DNA
  - HIF1a
  - Proline/Collagen
  - mRNA
  - rRNA

# IDH mutations block differentiation by impairing the removal of heterochromatic H3K9me2/3 methyl marks

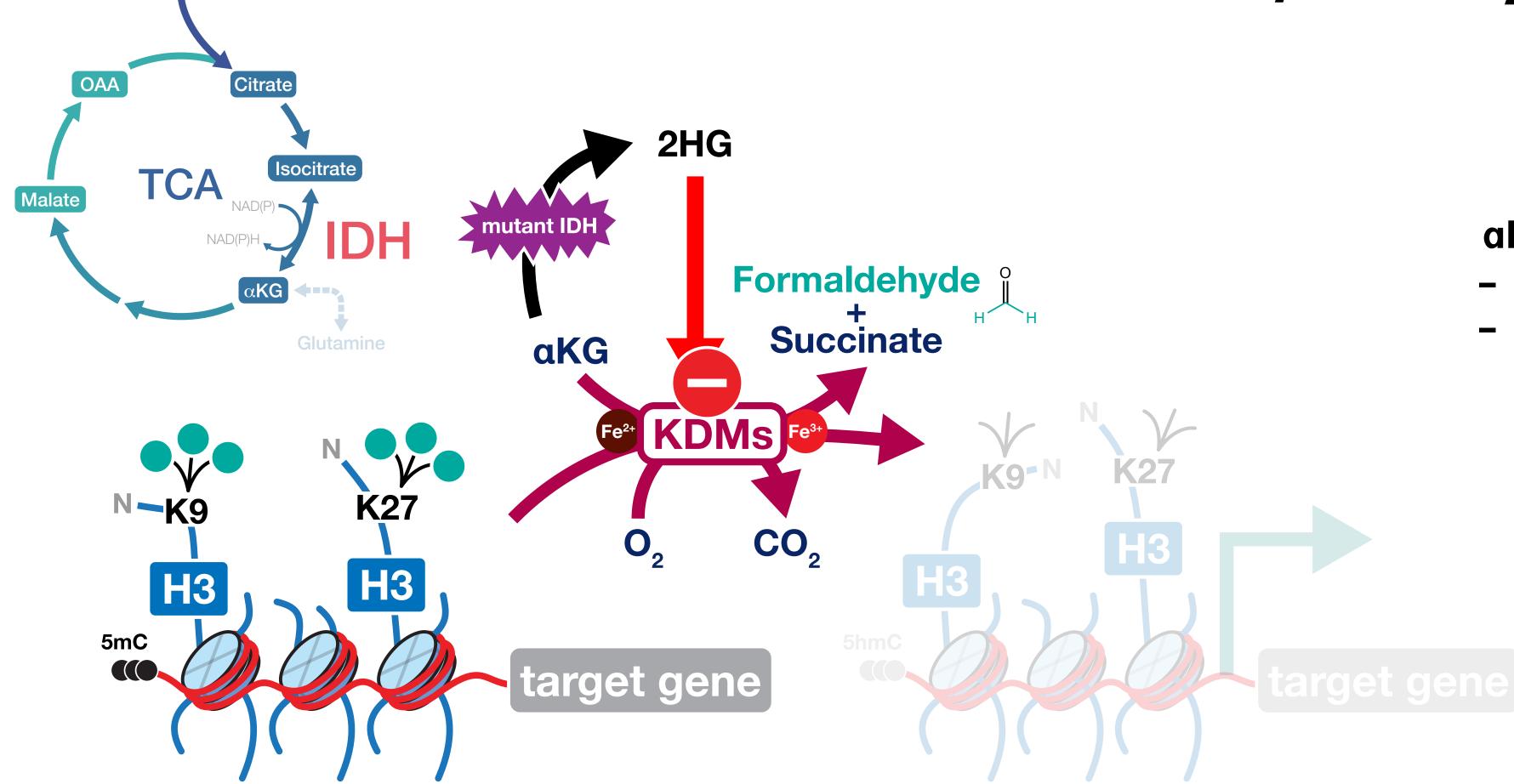


#### aKG-dependent dioxygenases

- Family of ~80 Enzymes
- Target diverse substrates:
  - Methylated Histones/DNA
  - HIF1a
  - Proline/Collagen
  - mRNA
  - rRNA

Pyruvate

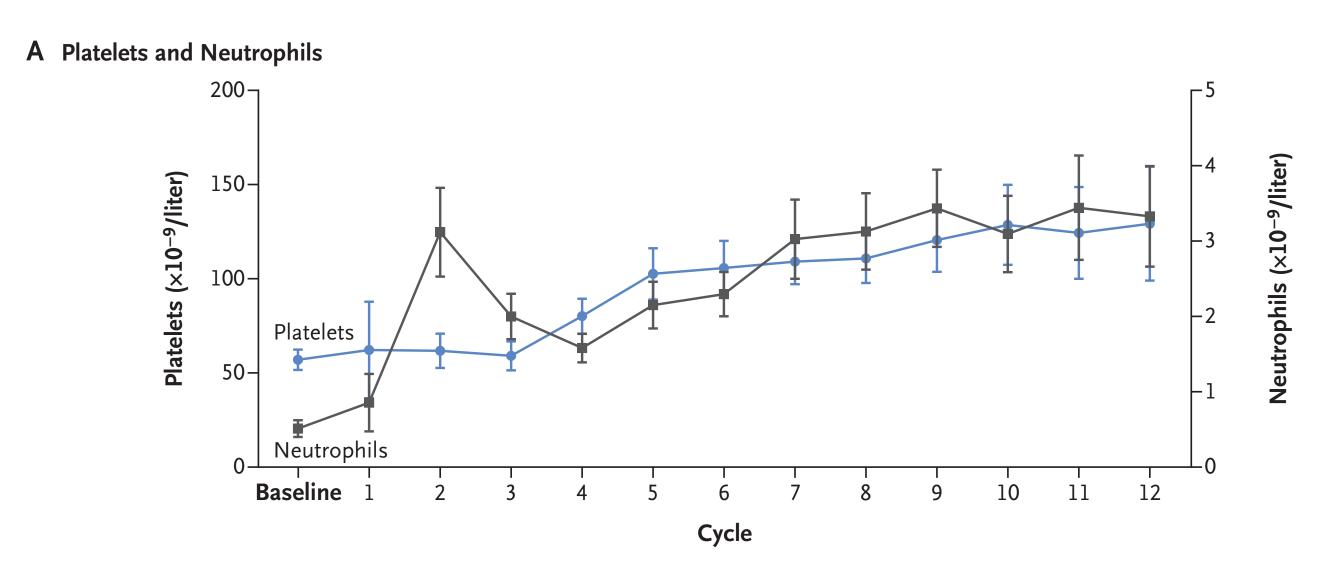
# IDH mutations block differentiation by impairing the removal of heterochromatic H3K9me2/3 methyl marks



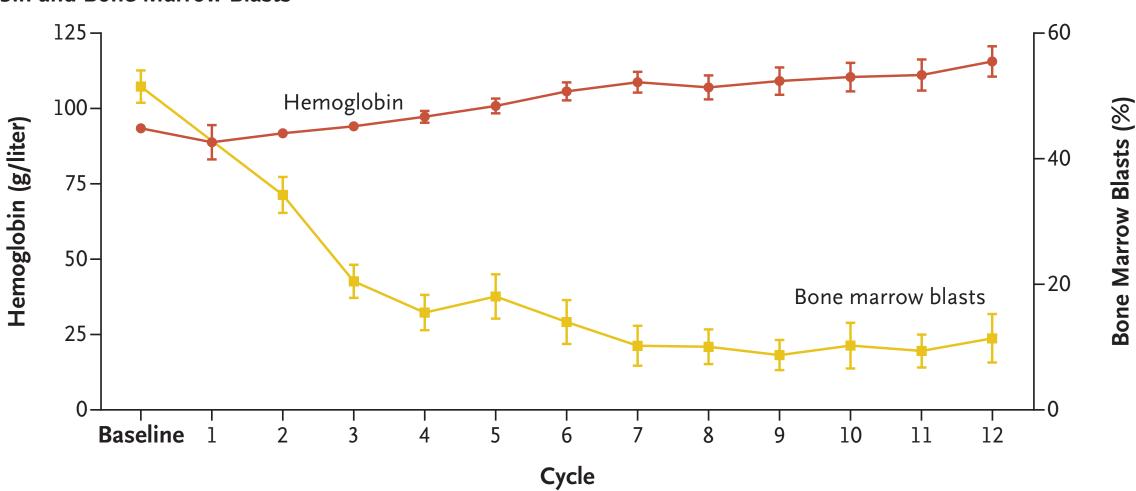
#### aKG-dependent dioxygenases

- Family of ~80 Enzymes
- Target diverse substrates:
  - Methylated Histones/DNA
  - HIF1a
  - Proline/Collagen
  - mRNA
  - rRNA

# IDH inhibitors are effective in IDH-mutated Leukemia and they work by differentiating leukemic cells

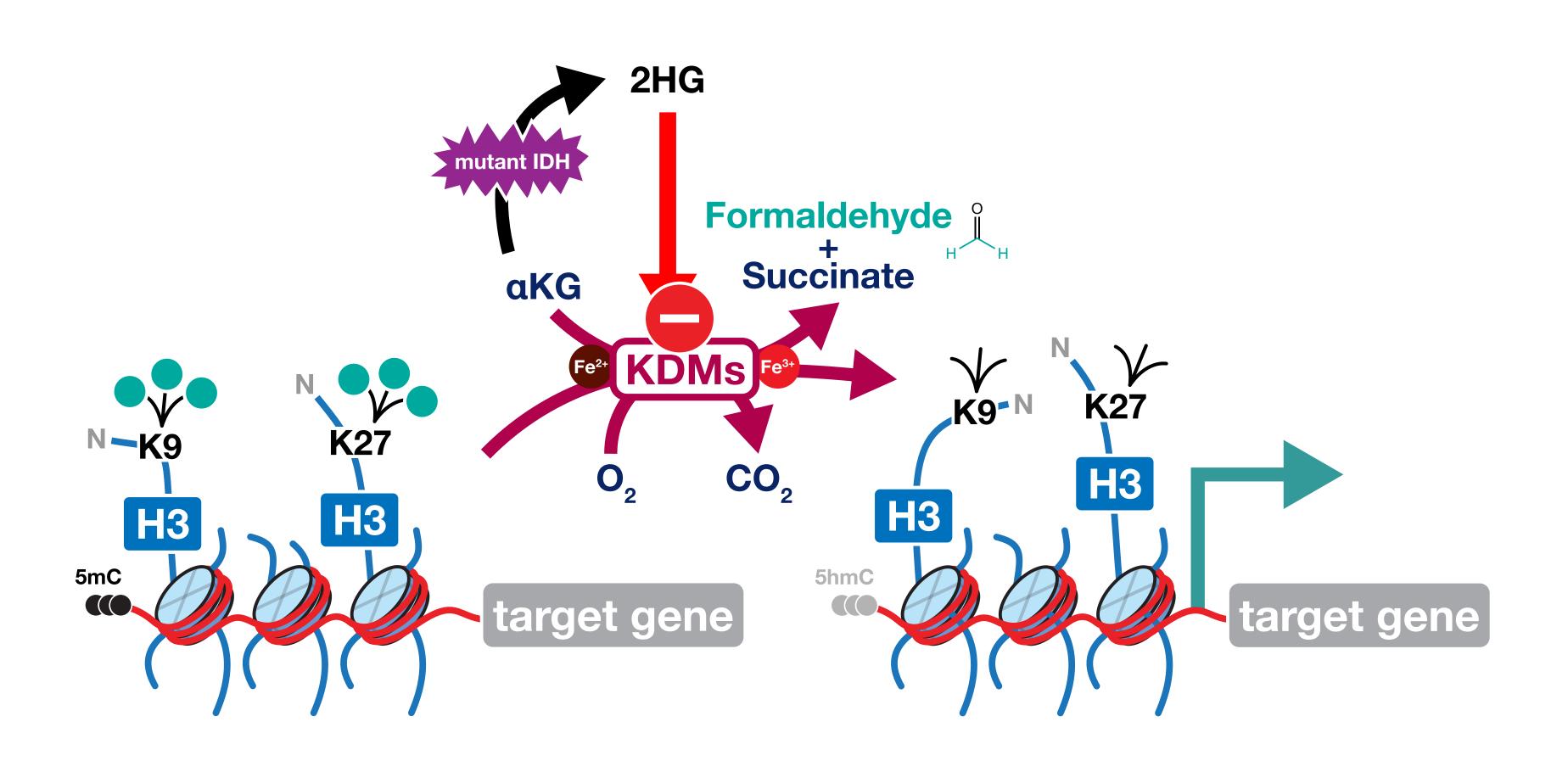


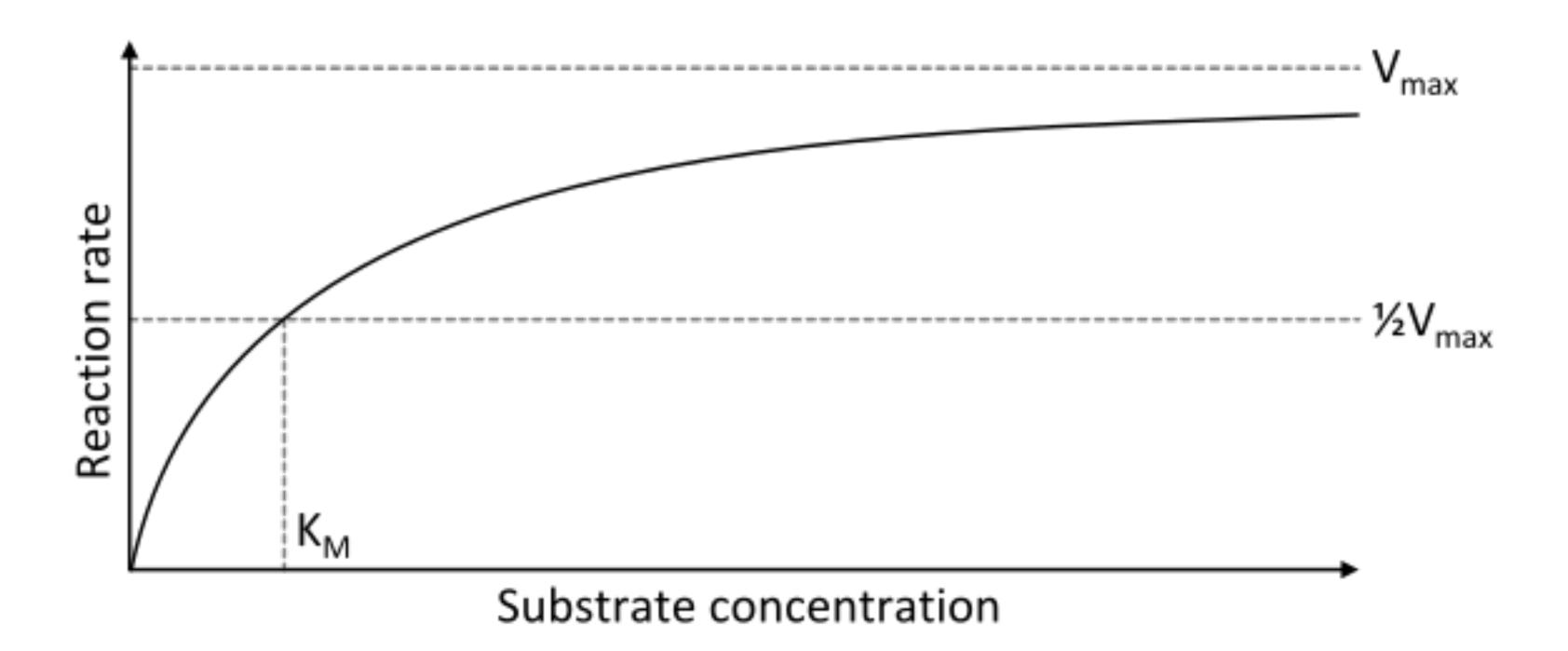




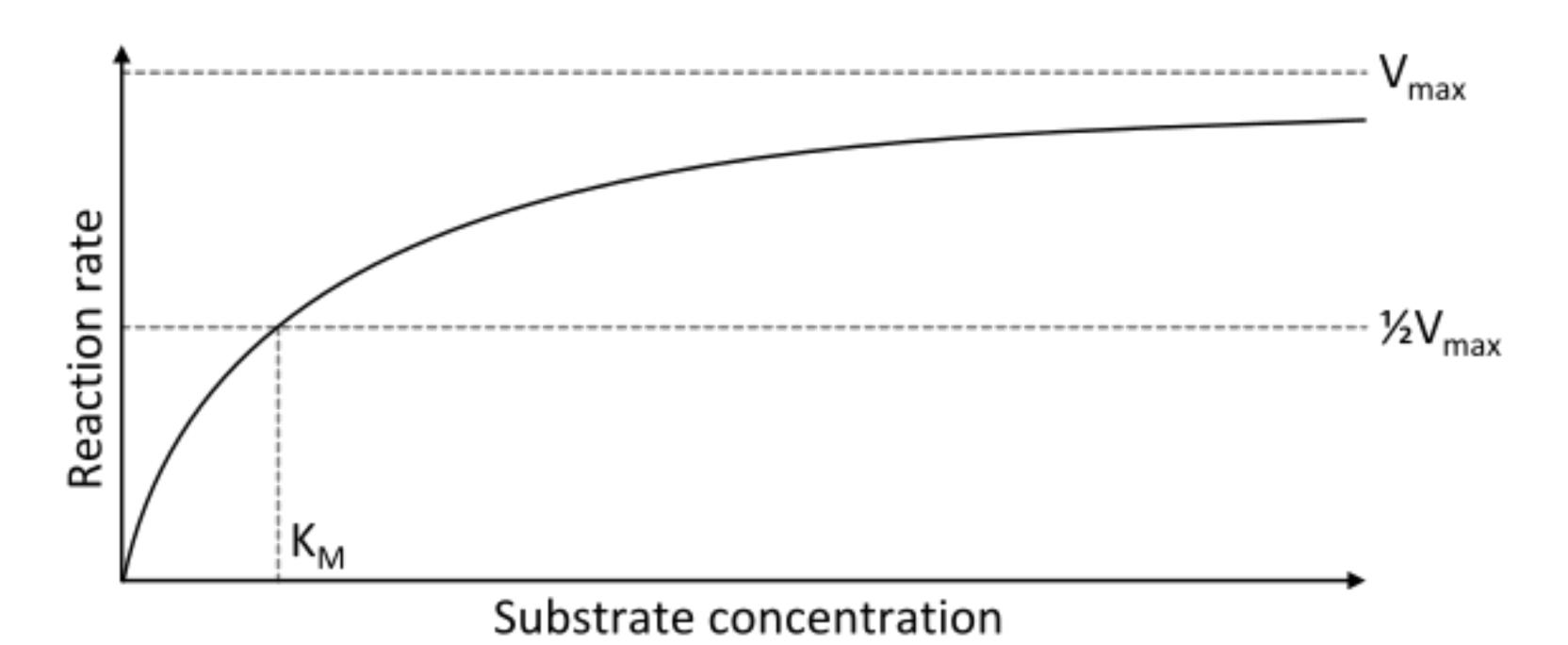
# IDH MUTATIONS OCCUR IN A SMALL FRACTION OF PATIENTS; ARE THESE CONCEPTS GENERALIZABLE?

#### Can dioxygenases act as sensors of metabolism?





#### Michaelis-Menten kinetics:



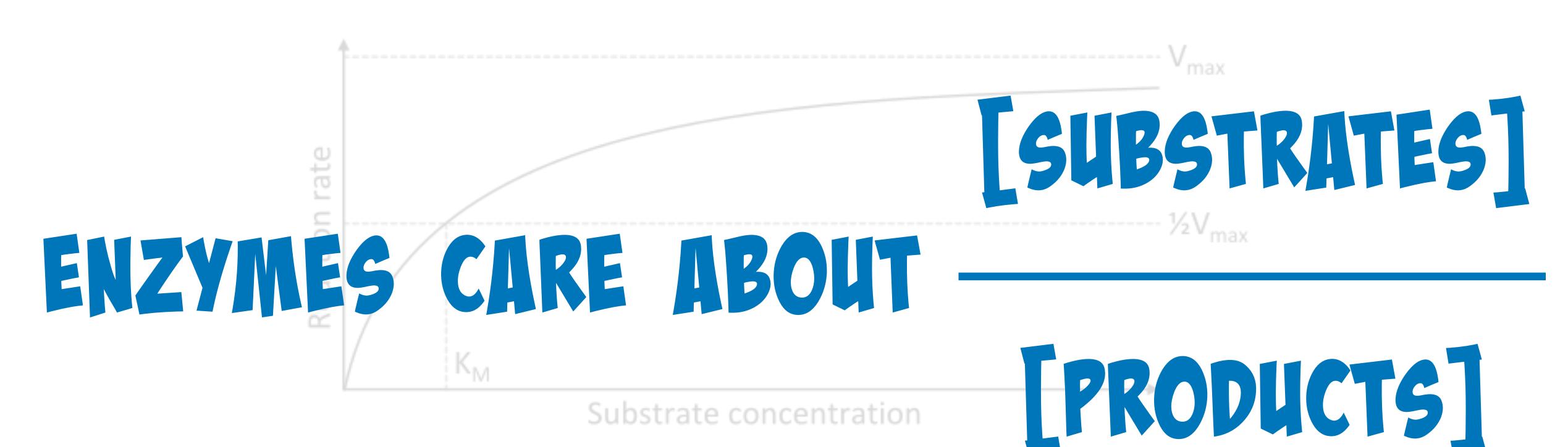
$$egin{array}{c} k_f & k_{
m cat} \ {
m E} + {
m S} & 
ightharpoons & {
m E} + {
m P} \ k_r & \end{array}$$

$$v = rac{ ext{d}[ ext{P}]}{ ext{d}t} = V_{ ext{max}} rac{[ ext{S}]}{K_{ ext{M}} + [ ext{S}]}$$





Leonor Michaelis Maud Menten

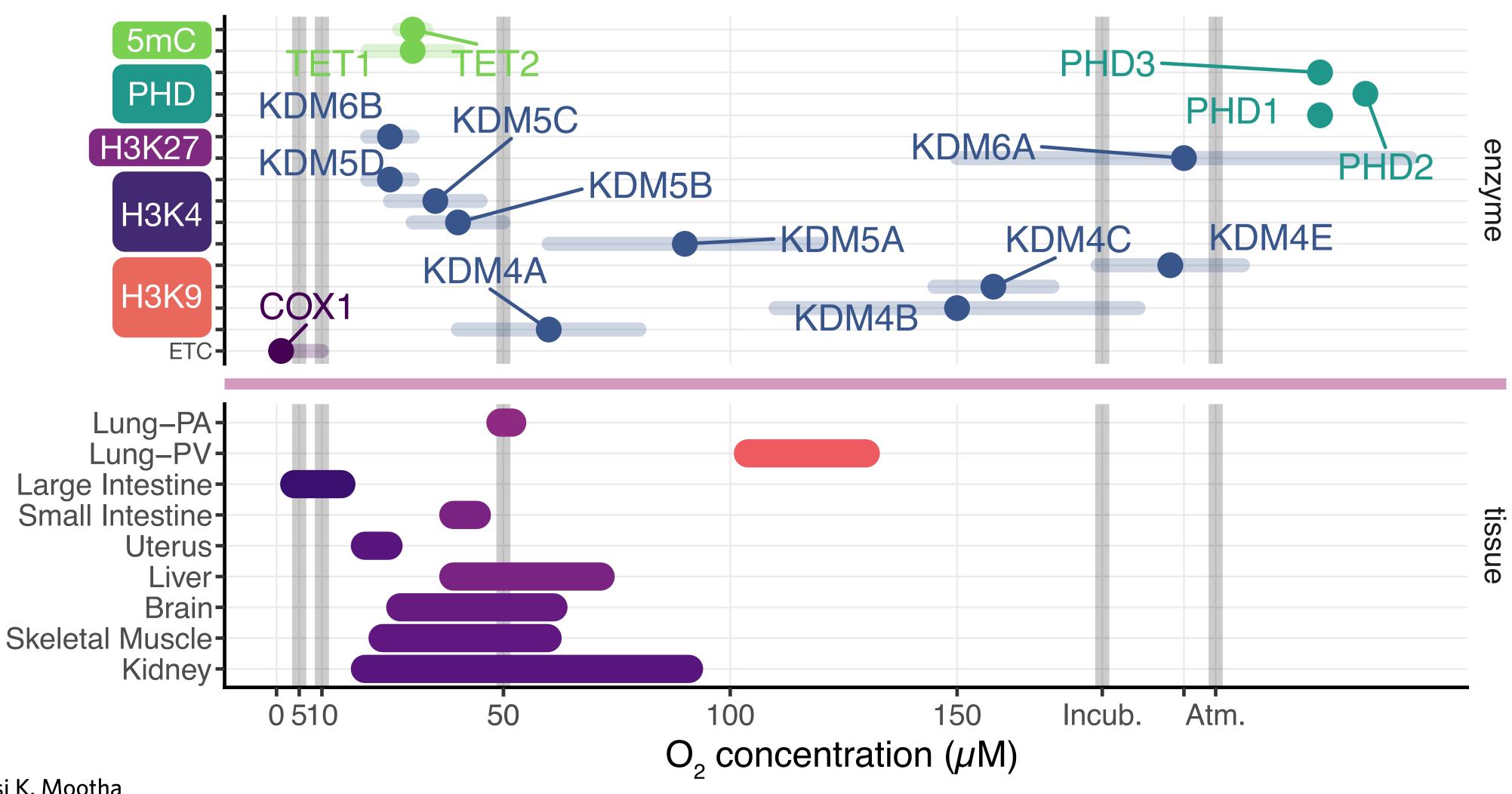




Leonor Michaelis Maud Menten

Substrate concentration

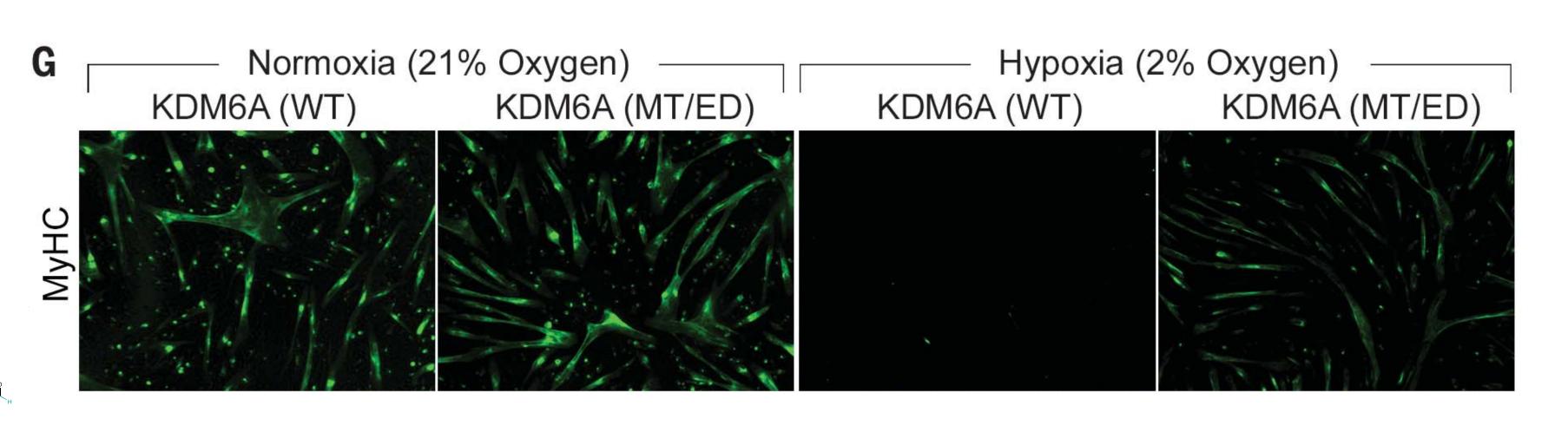
#### The 'regulatable' space of chromatin metabolism



Tslil Ast and Vamsi K. Mootha
Oxygen and mammalian cell culture: are we repeating the experiment of Dr. Ox?

### Are dioxygenases oxygen sensors?



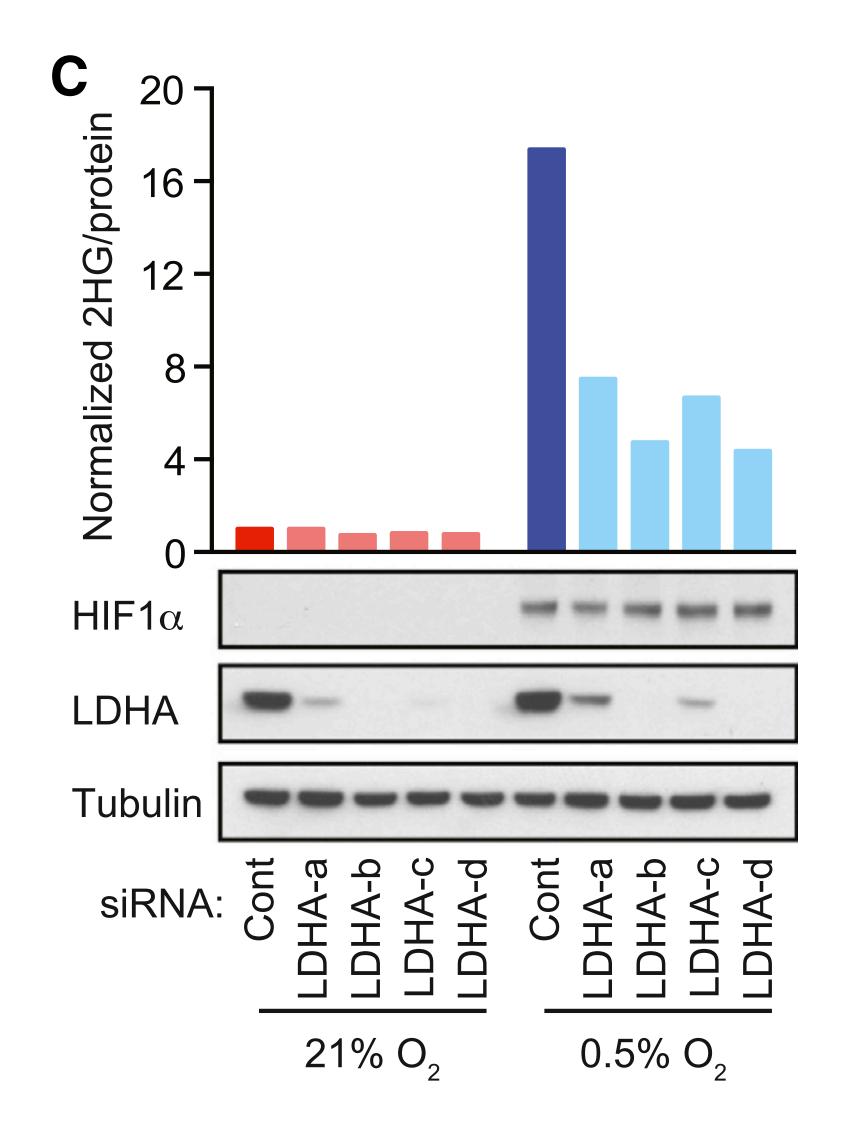


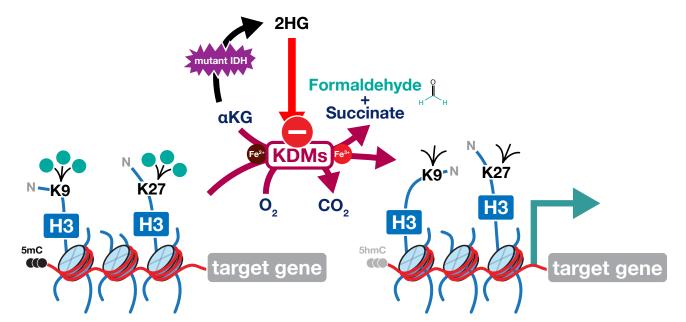


William Kaelin

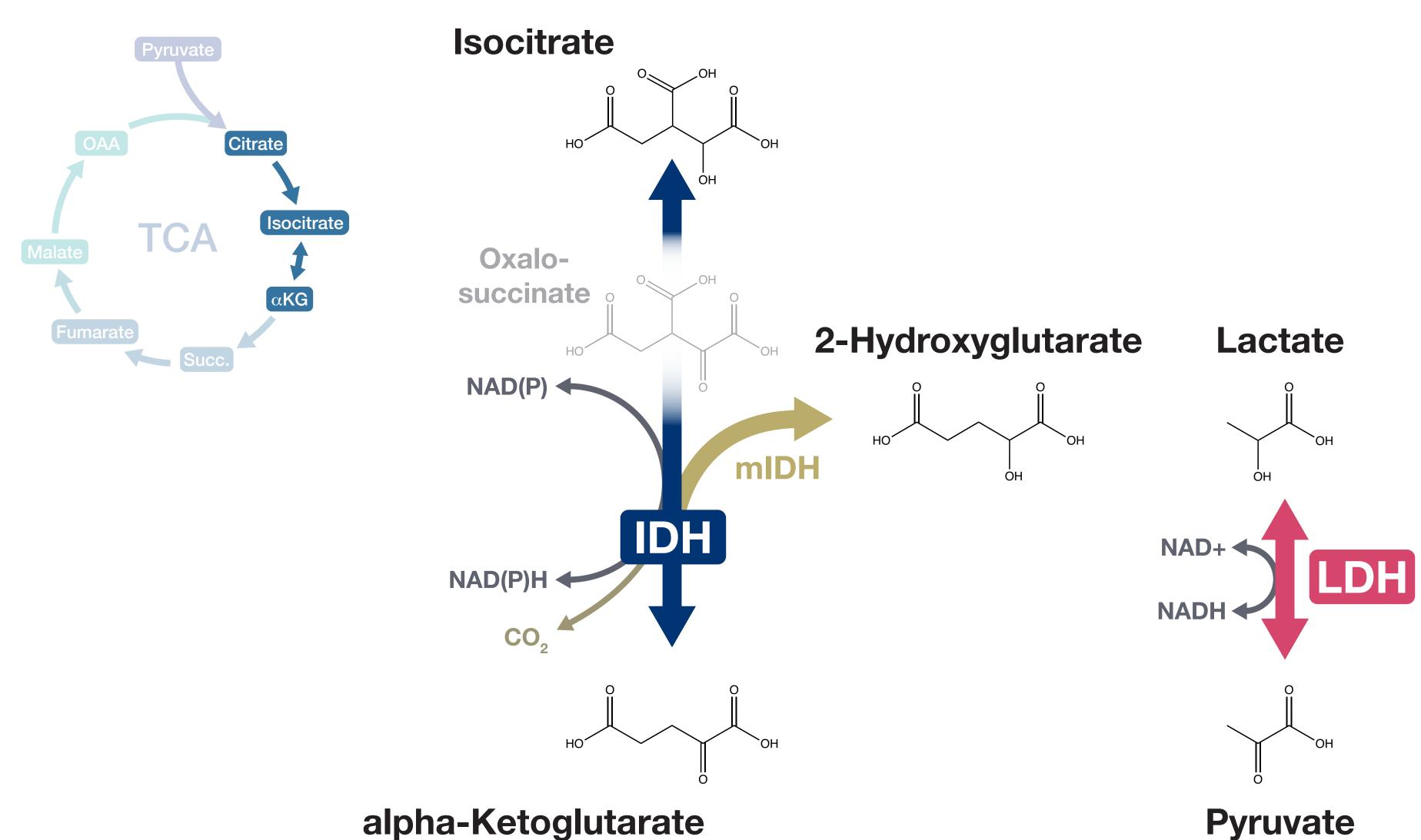
### ENZYMES DON'T READ TEXTBOOKS EITHER

# 2HG can be produced in hypoxia and acidosis independently of IDH mutations

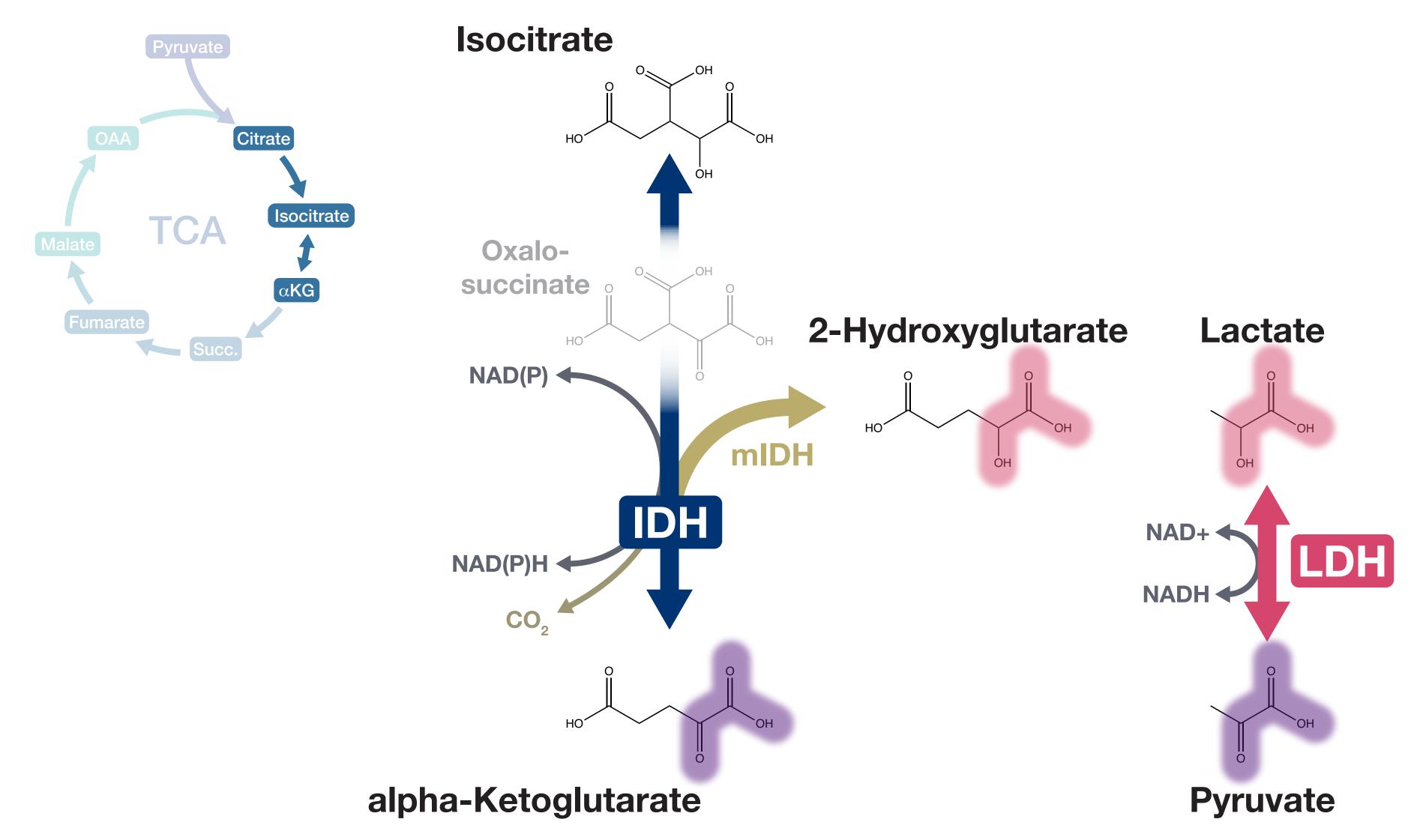




# 2HG can be produced in hypoxia and acidosis independently of IDH mutations

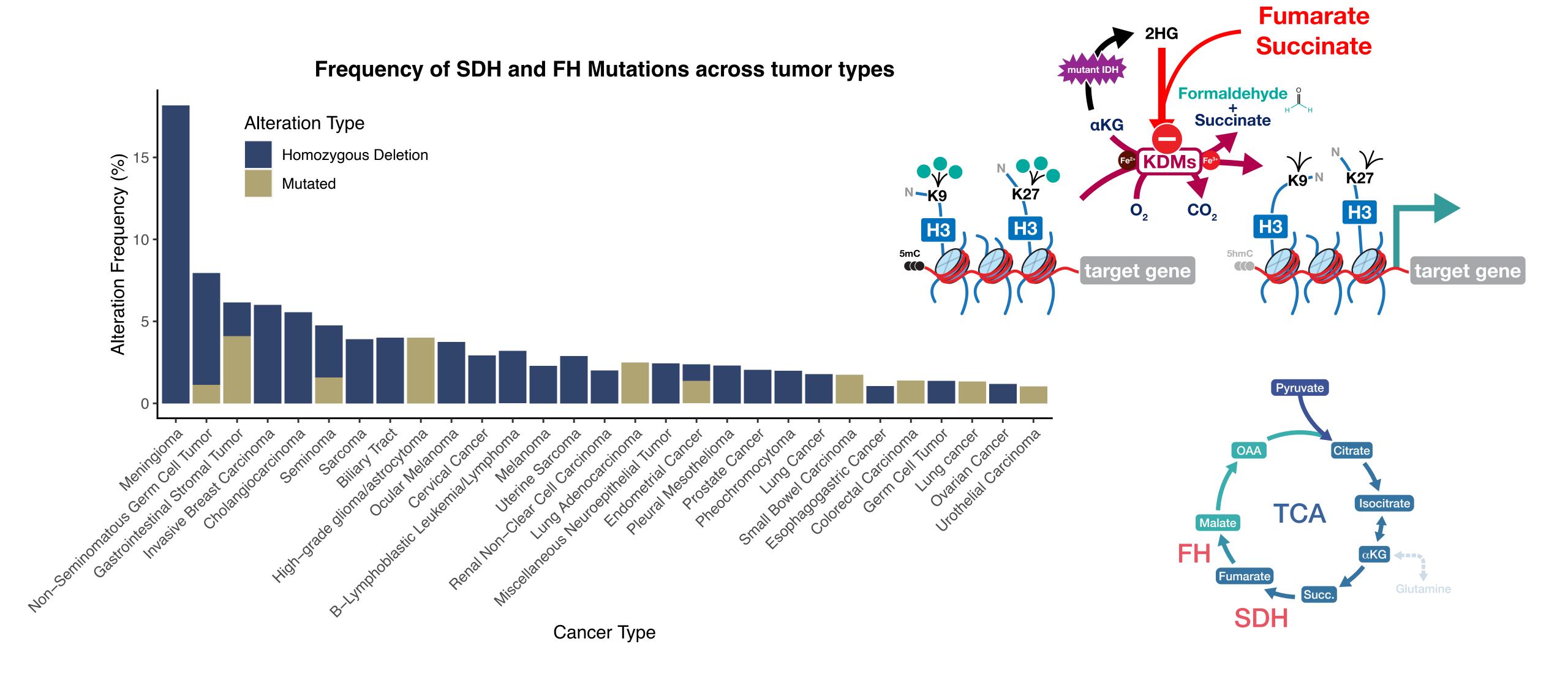


# 2HG can be produced in hypoxia and acidosis independently of IDH mutations



### PRODUCTS MATTER TOO!!!

### Succinate (and Fumarate) can also inhibit dioxygenase function

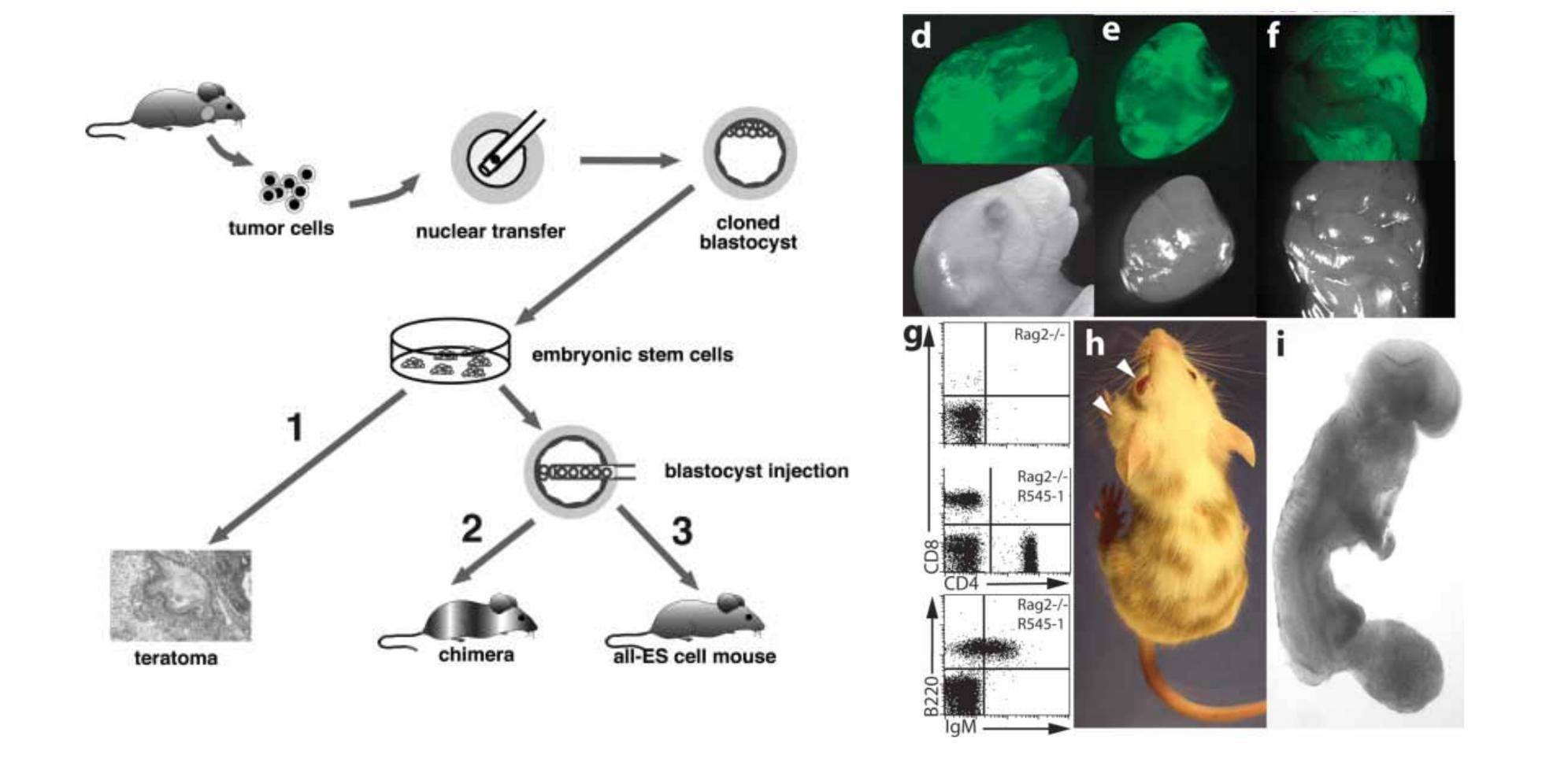


### Takehome points 3

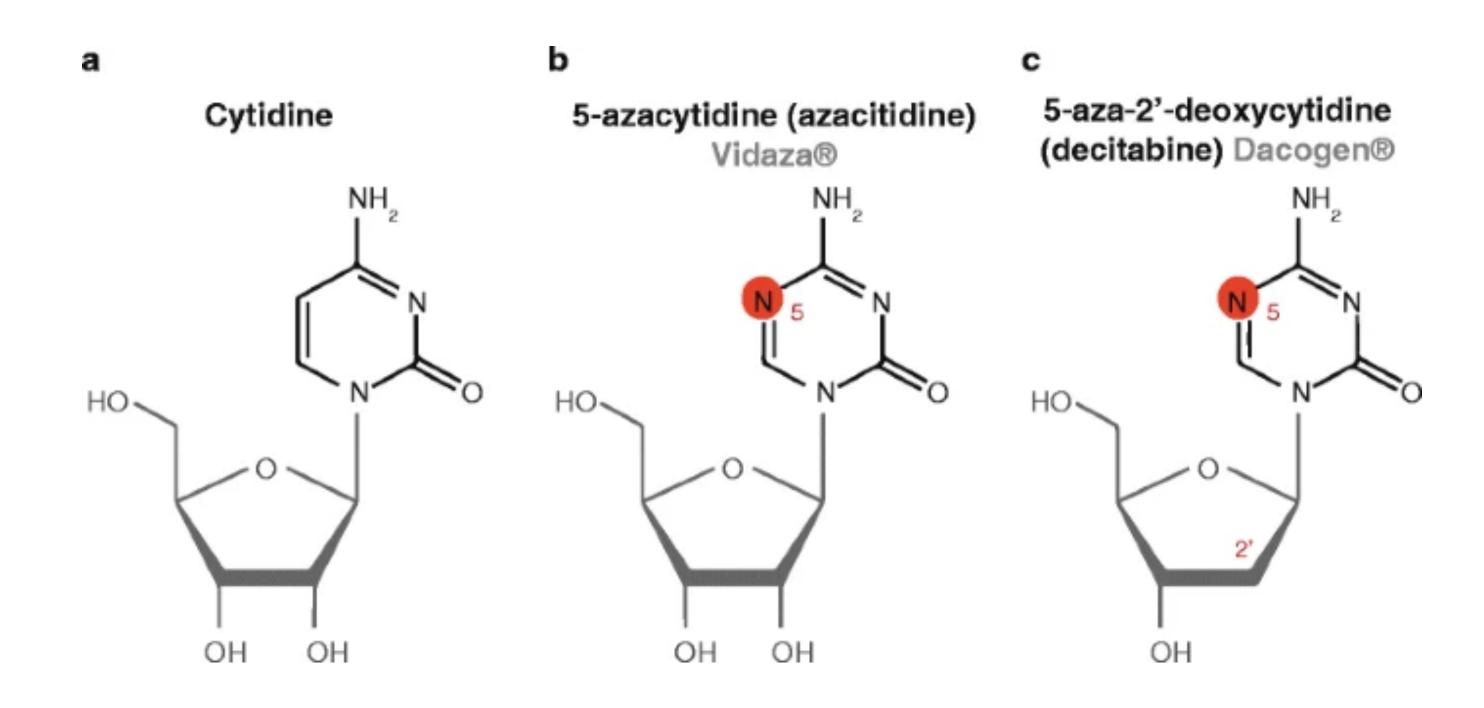
- 1. Chromatin is a substrate for many metabolic enzymes
- 2. Chromatin modifications have meaning/impart function
- 3. Substrates Matter: altering the levels of metabolites can affect chromatin function
- 4. Mutations can alter metabolism → alter chromatin / cell states
- 5. Remember the KM!!!
- 6. Enzymes also care about products
- 7. Enzymes don't read textbooks



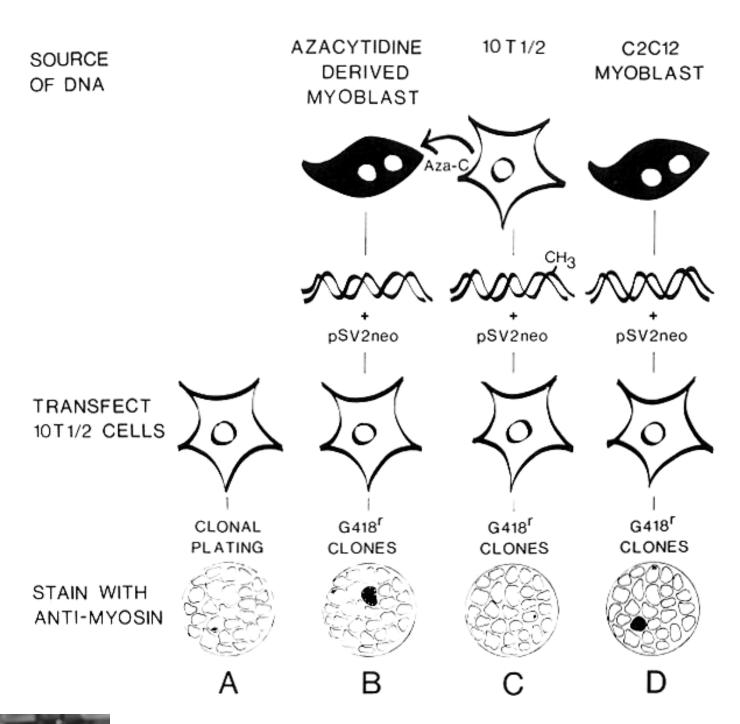
### Even a cancer cell line (melanoma) can be reprogrammed

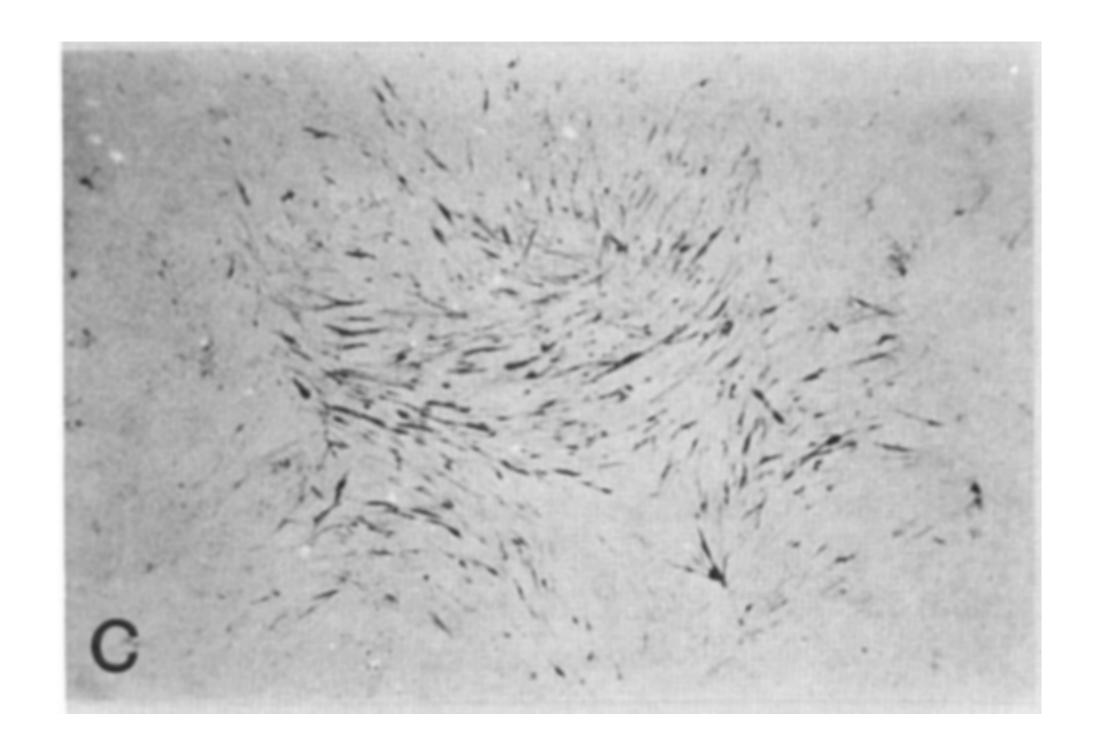


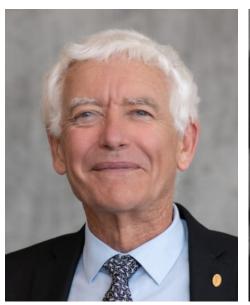
# Inhibition of DNA methylation can lead to differentiation through derepression of lineage-specifying transcription factors



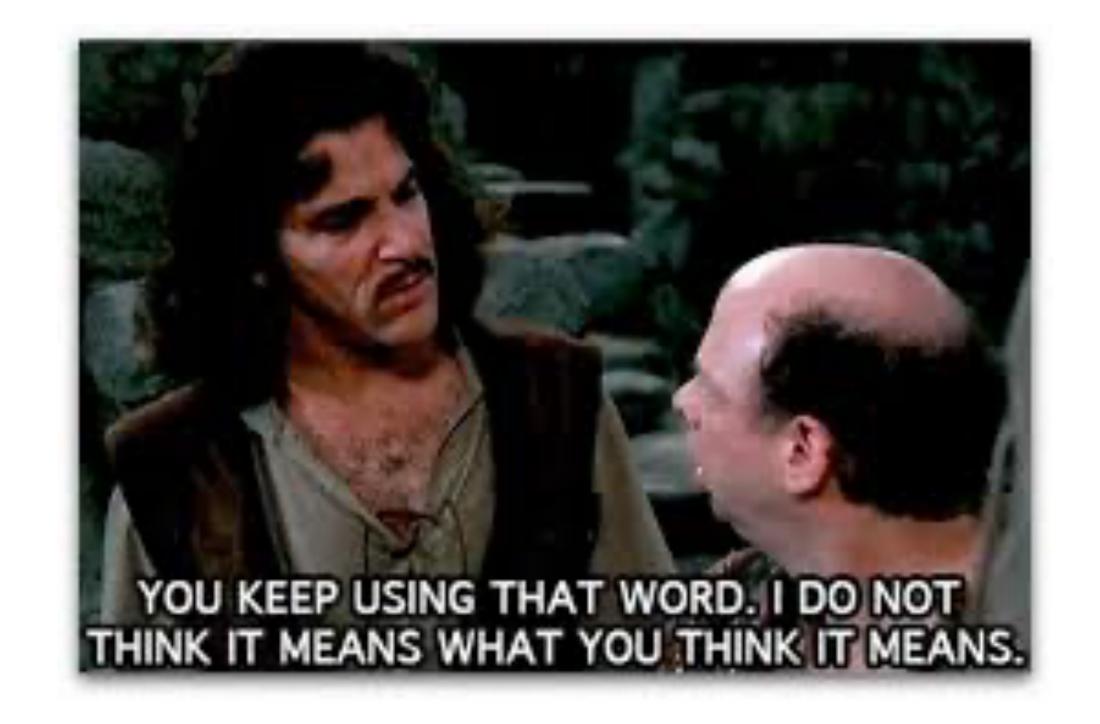
# Inhibition of DNA methylation can lead to differentiation through derepression of lineage-specifying transcription factors





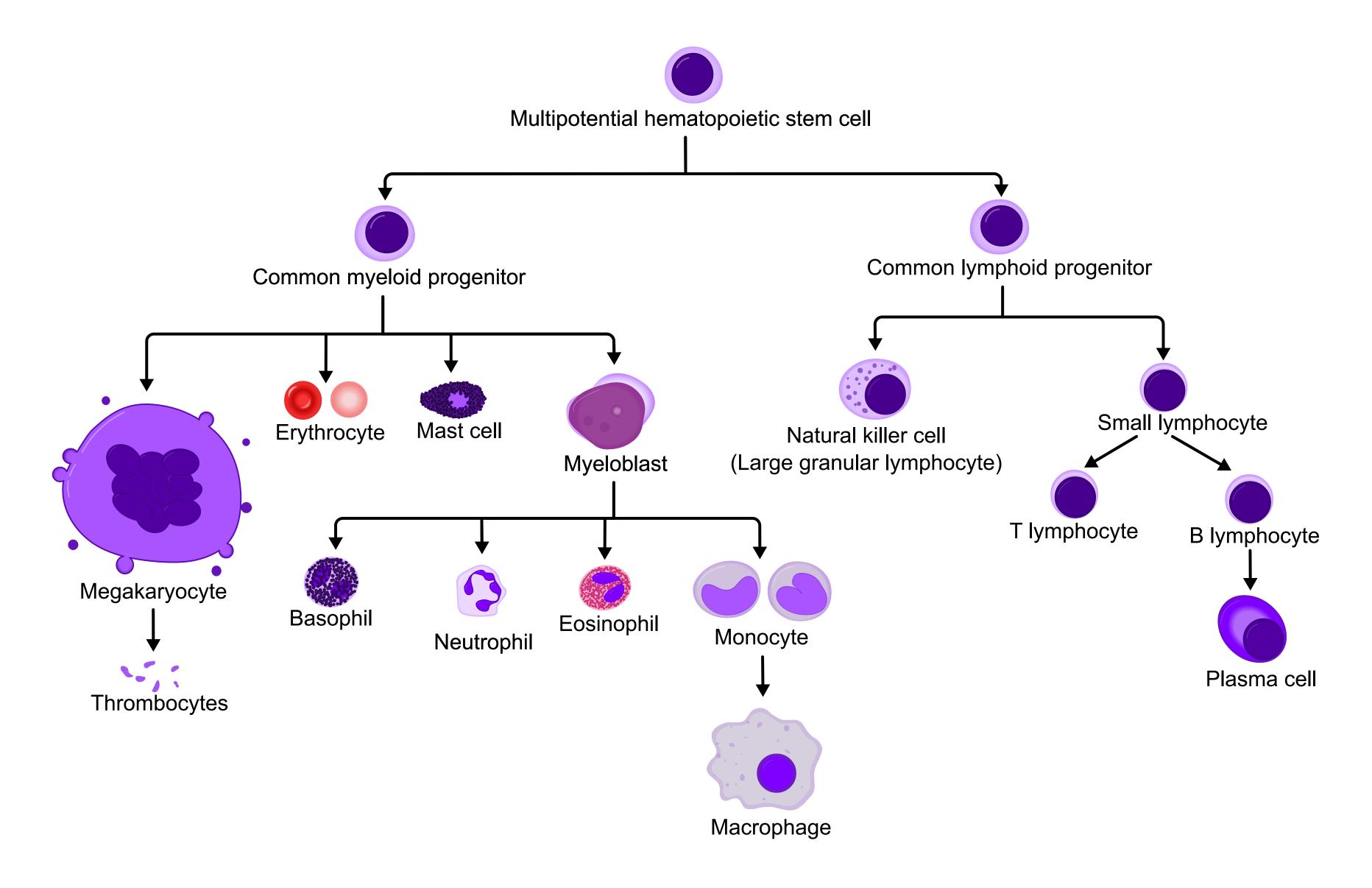


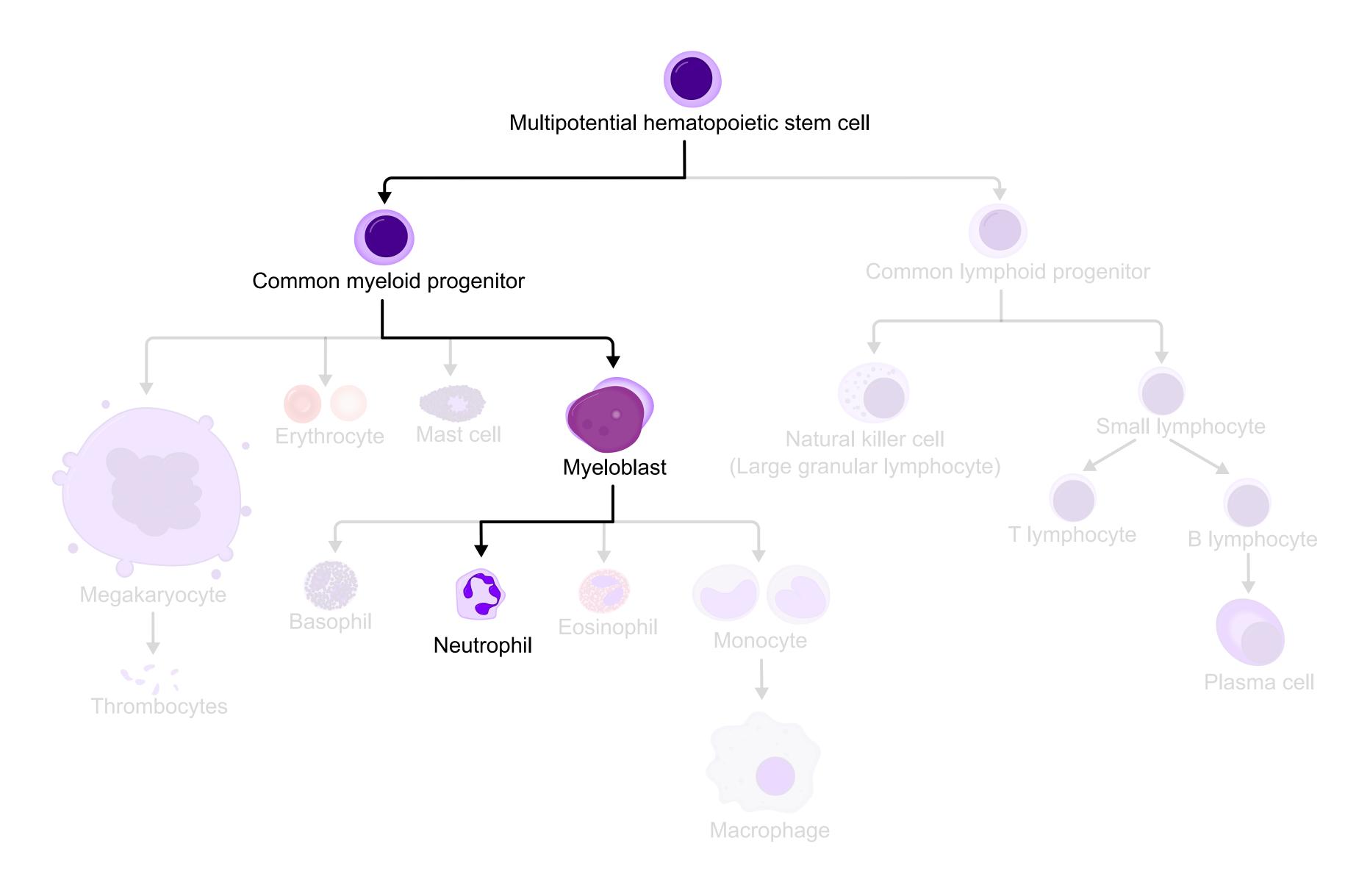
Peter Jones Harold Weintraub

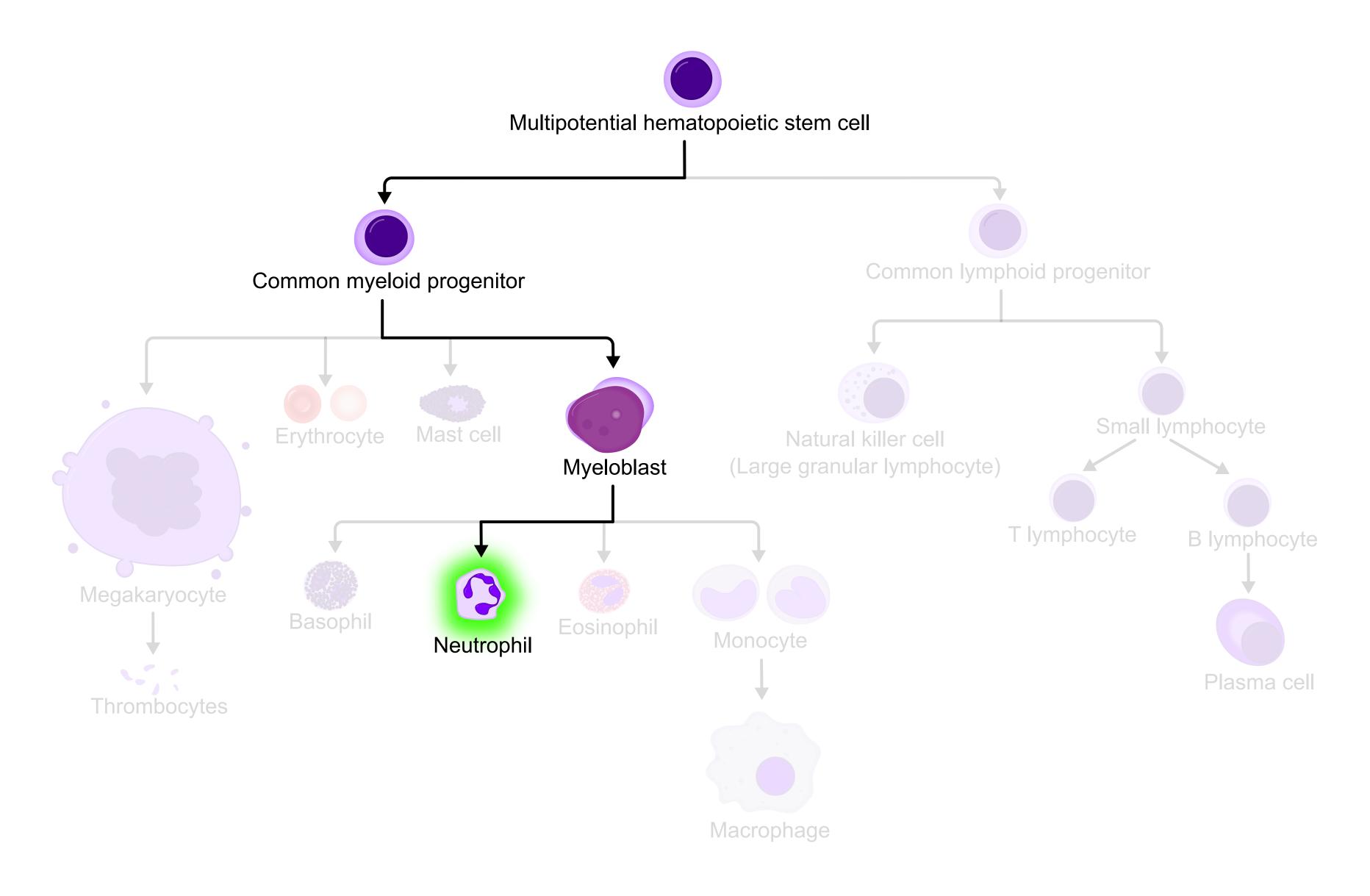


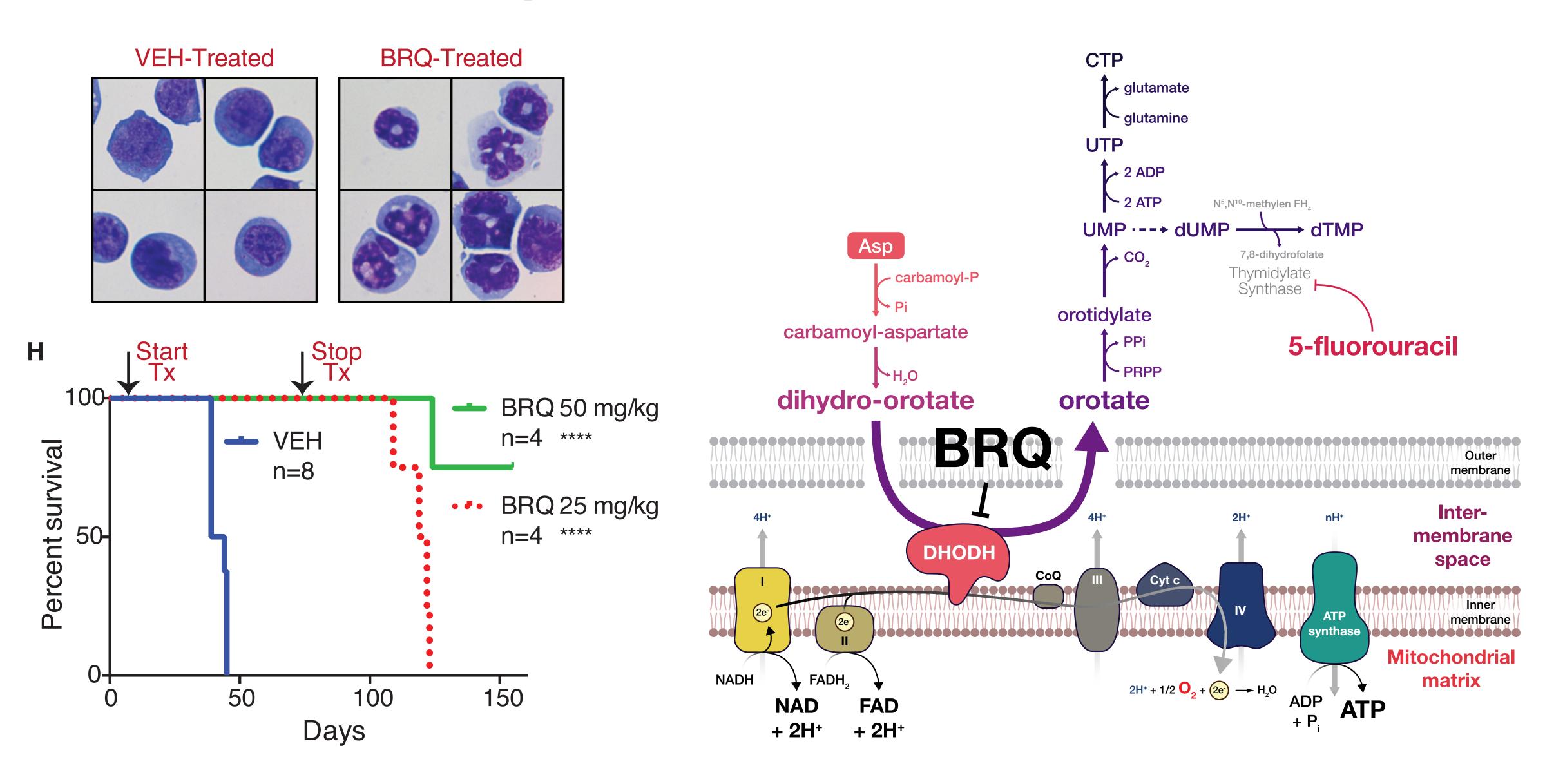
REVISITING ANTIMETABOLITES

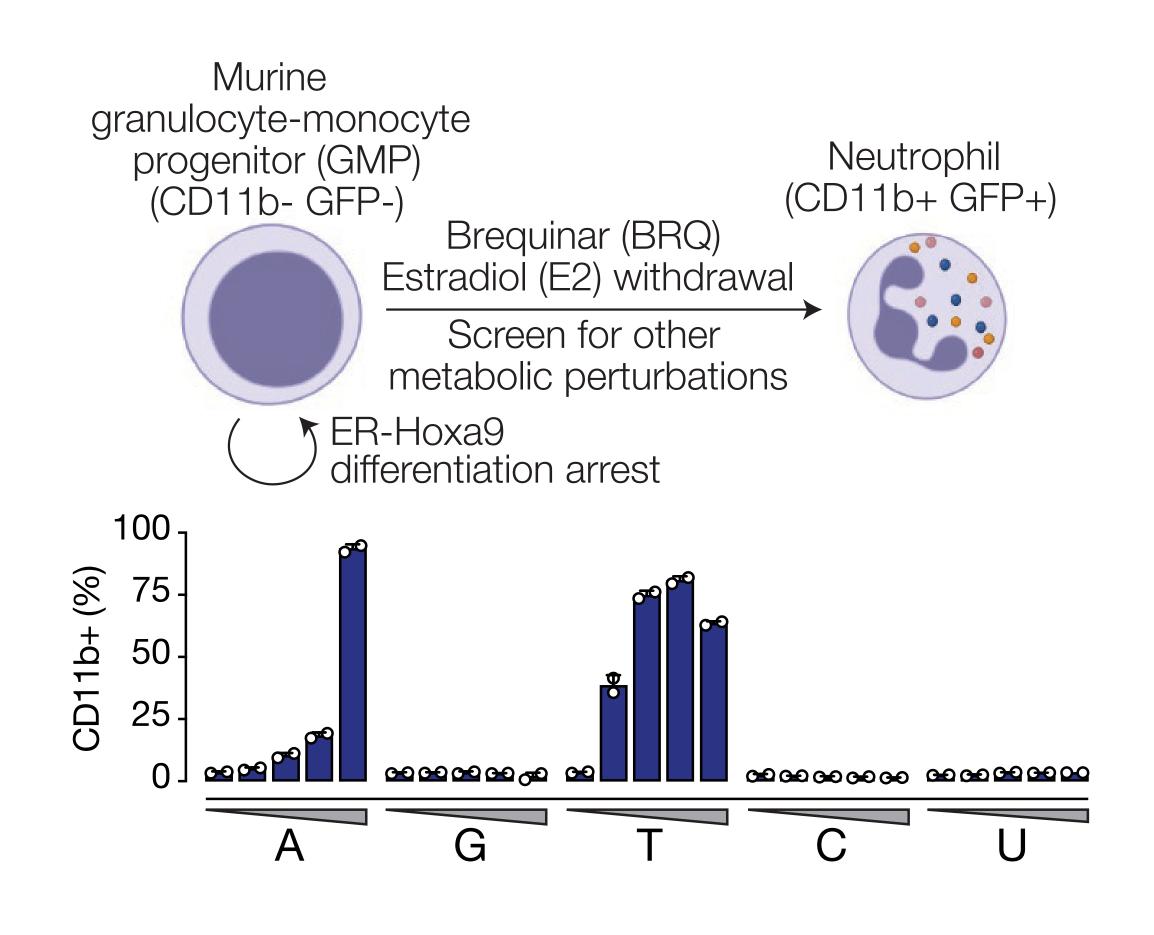
5-azacytidine
Decitabine
5-fluorouracil
Capecitabine
Floxuridine
Fludarabine
Cytarabine
Gemcitabine
Azathioprine
Mercaptopurine
Methotrexate
Pemetrexed
(Hydroxyurea NB RNRi)

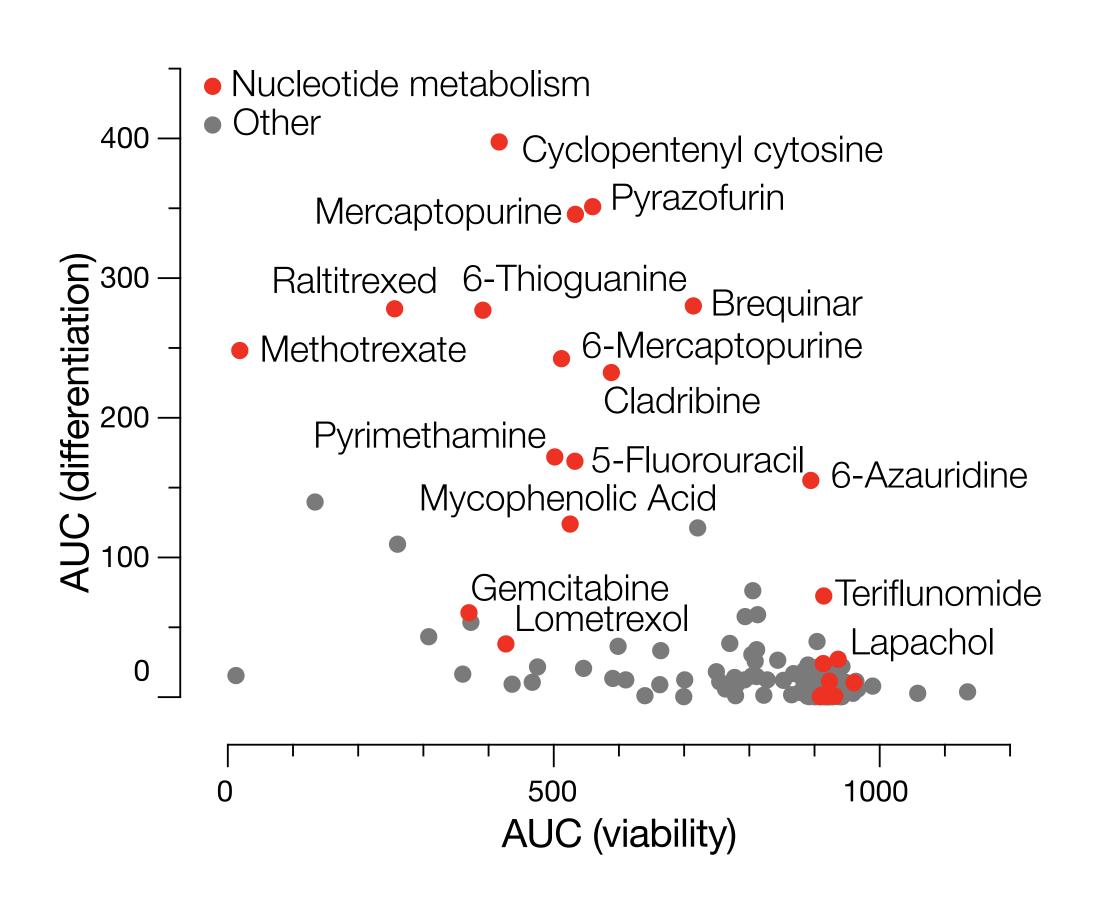








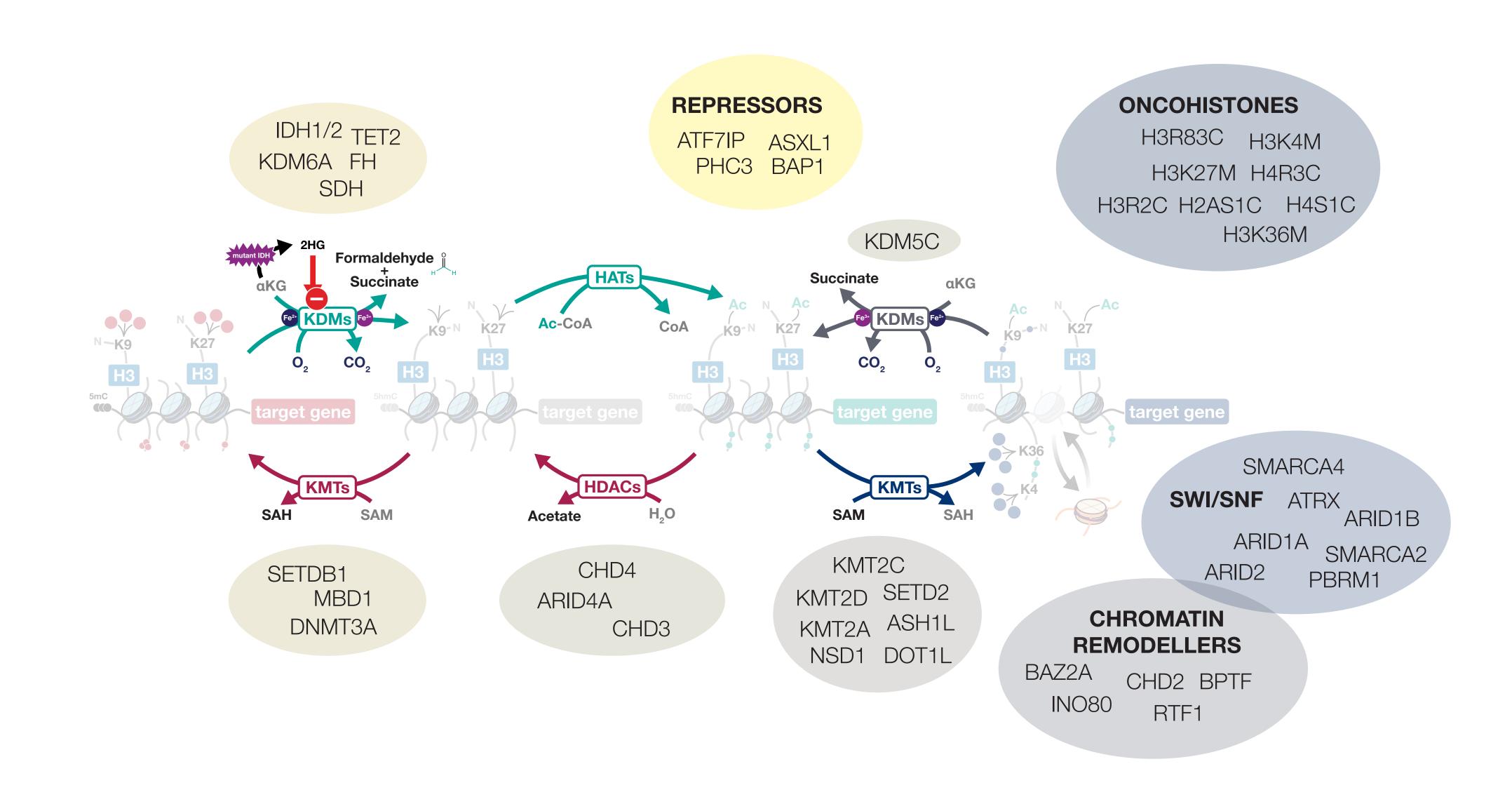




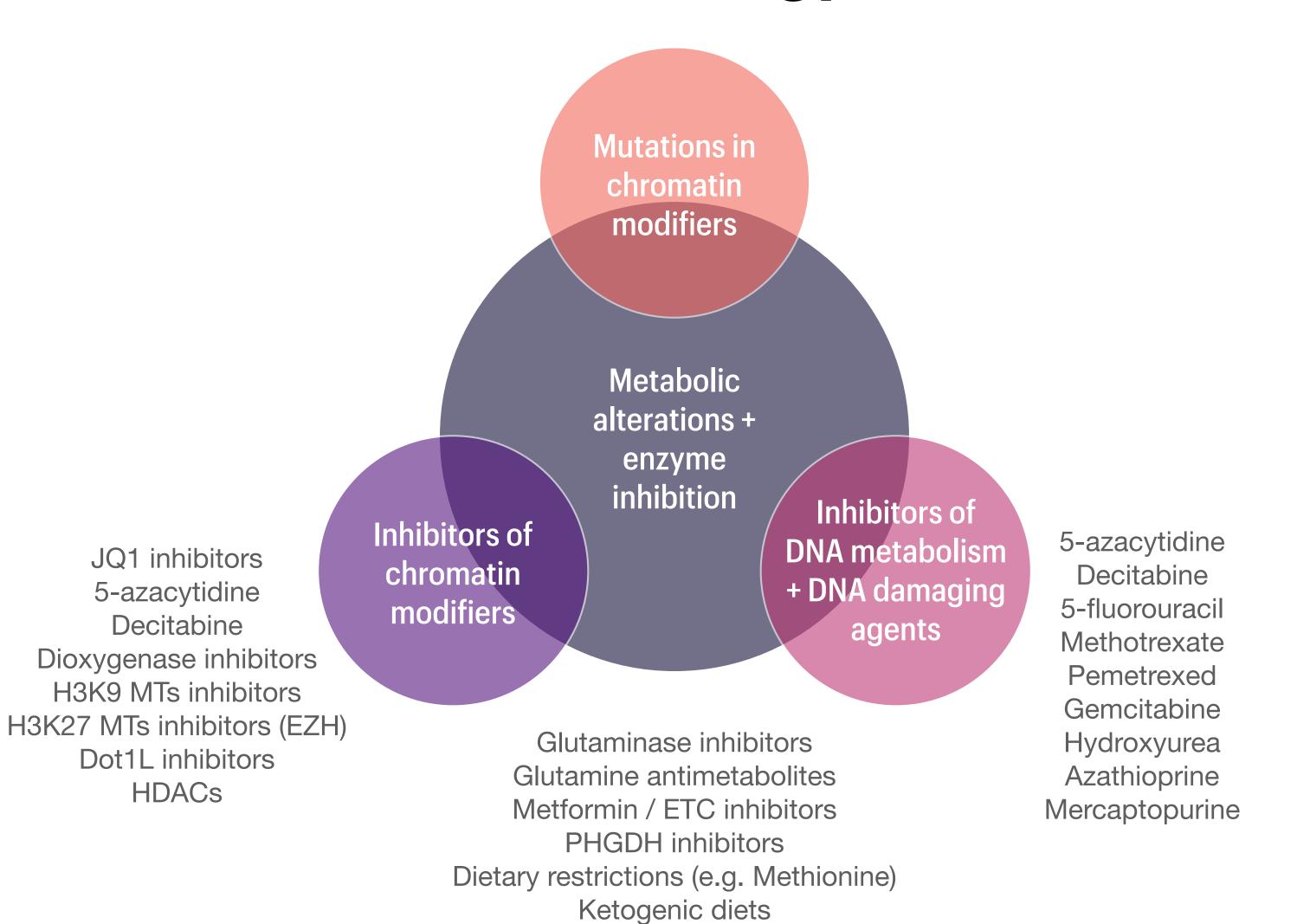
### ALTERING NUCLEOTIDES CAN DIFFERENTIATE CELLS????



### A wide range of chromatin modifying enzymes are mutated in cancers



# Metabolic pathways as a therapeutic handle for chromatin biology in cancer

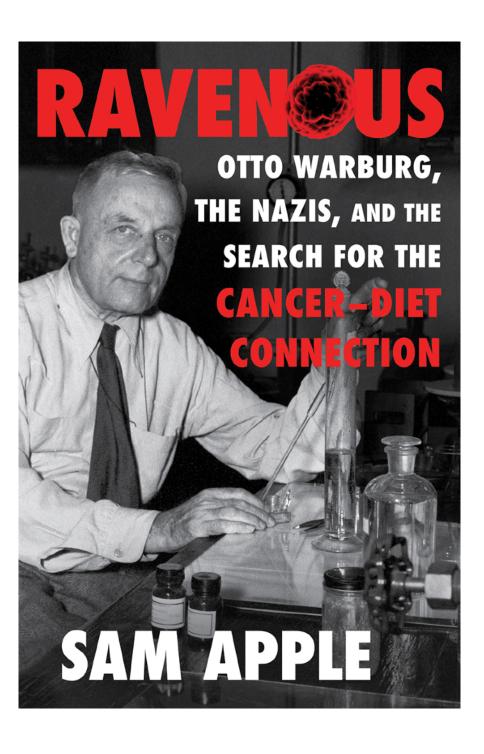


### Final Takehome points

- 1. Cells don't read textbooks
- 2. Metabolism is wired to meet different anabolic/catabolic needs
- 3. Many pathways are redundant → Auxotrophy + forward genetics to understand metabolic pathways
- 4. If you want to look at metabolism, look at metabolism:
  - metabolomics and flux analysis; NOT RNA!!
- 5. Km: Enzymes care about [substrates] / [products]
- 6. All substrates/products matter
- 7. Enzymes also don't read textbooks

### If you want to learn more:









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