



Maple Syrup Urine Disease and the DBT Gene

Tessa Bachinski

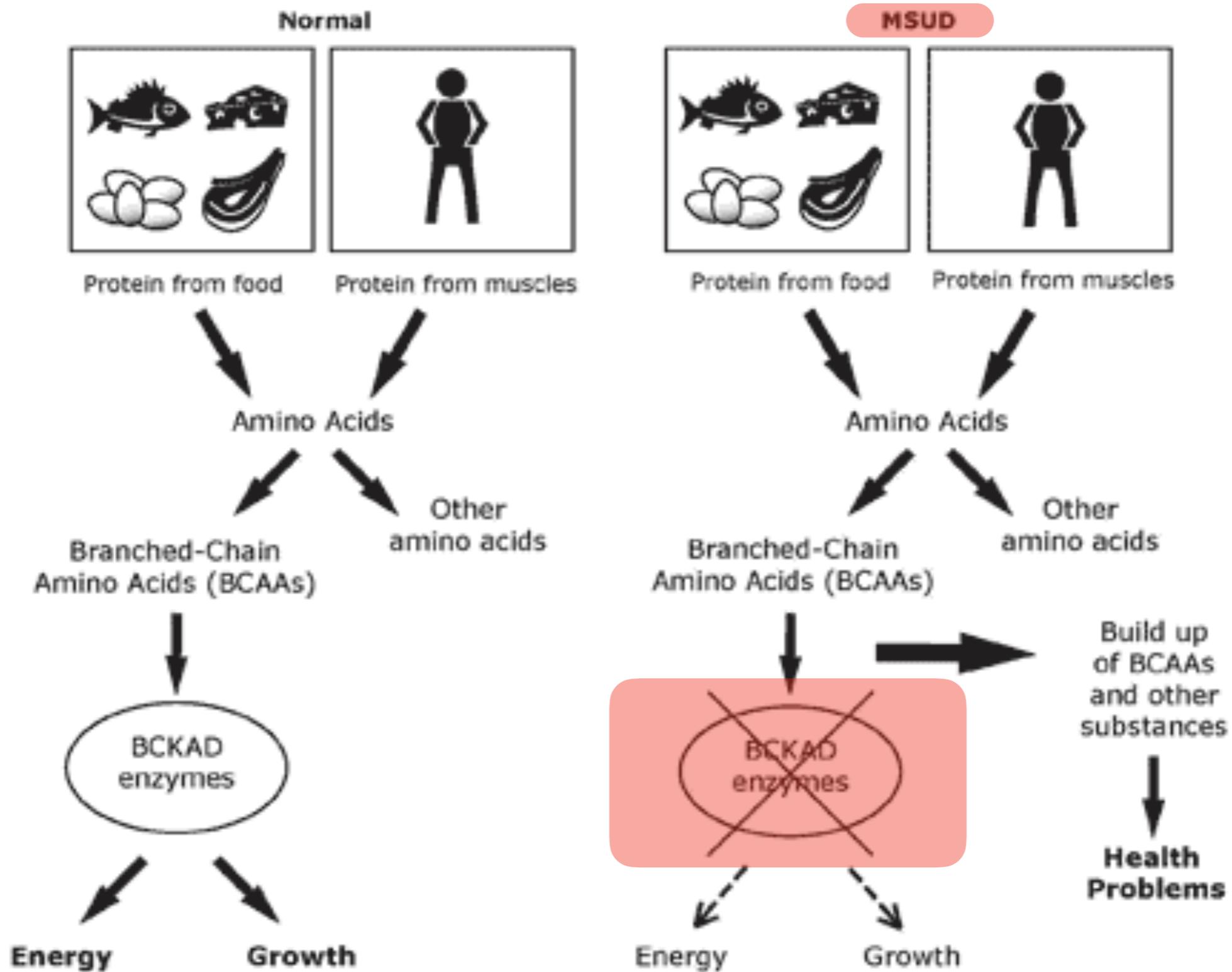
What is Maple Syrup Urine Disease (MSUD)?



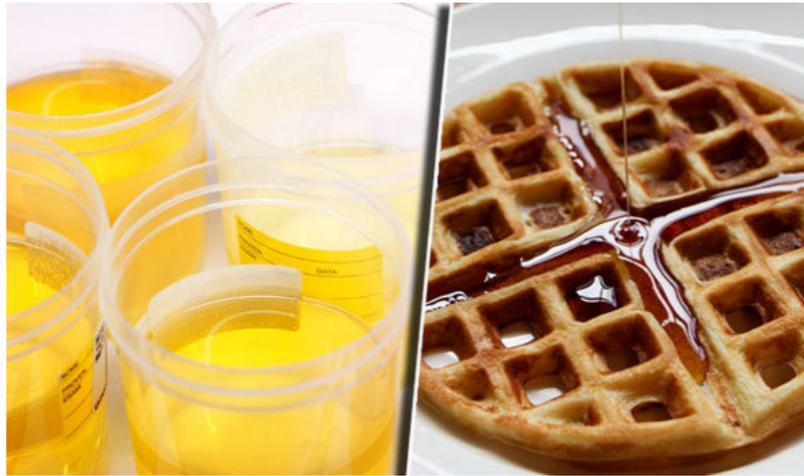
Recommended Uniform Screening Panel
Core Conditions
(As of July 2018)

Core Condition	Metabolic Disorder			Endocrine Disorder	Hemoglobin Disorder	Other Disorder
	Organic acid condition	Fatty acid oxidation disorder	Amino acid disorder			
Propionic Acidemia	X					
Methylmalonic Acidemia (methylmalonyl-CoA mutase)	X					
Methylmalonic Acidemia (Cobalamin disorders)	X					
Isovaleric Acidemia	X					
3-Methylcrotonyl-CoA Carboxylase Deficiency	X					
3-Hydroxy-3-Methylglutaric Aciduria	X					
Holocarboxylase Synthase Deficiency	X					
β-Ketothiolase Deficiency	X					
Glutaric Acidemia Type I	X					
Carnitine Uptake Defect/Carnitine Transport Defect		X				
Medium-chain Acyl-CoA Dehydrogenase Deficiency		X				
Very Long-chain Acyl-CoA Dehydrogenase Deficiency		X				
Long-chain L-3 Hydroxyacyl-CoA Dehydrogenase Deficiency		X				
Trifunctional Protein Deficiency		X				
Argininosuccinic Aciduria			X			
Citrullinemia, Type I			X			
Maple Syrup Urine Disease			X			
Homocystinuria			X			
Classic Phenylketonuria			X			
Tyrosinemia, Type I			X			
Primary Congenital Hypothyroidism				X		
Congenital adrenal hyperplasia				X		
S,S Disease (Sickle Cell Anemia)					X	
S, β-Thalassemia					X	
S,C Disease					X	
Biotinidase Deficiency						X
Critical Congenital Heart Disease						X
Cystic Fibrosis						X
Classic Galactosemia						X
Glycogen Storage Disease Type II (Pompe)						X
Hearing Loss						X
Severe Combined Immunodeficiencies						X
Mucopolysaccharidosis Type 1						X
X-linked Adrenoleukodystrophy						X
Spinal Muscular Atrophy due to homozygous deletion of exon 7 in SMN1						X

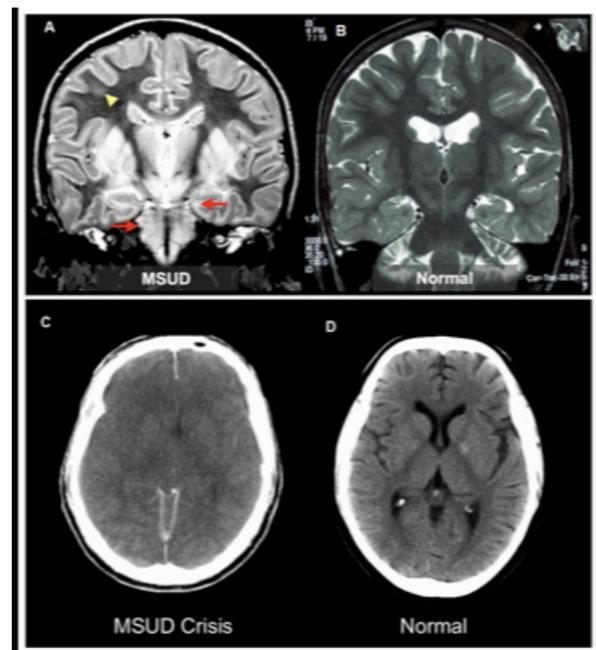
What Causes Maple Syrup Urine Disease?



What are the symptoms of Maple Syrup Urine Disease?



Sweet Smelling Urine



**Brain Damage, Seizures,
and Other Neurological
Symptoms**



Poor Feeding, Lethargy

What is the DBT Gene?

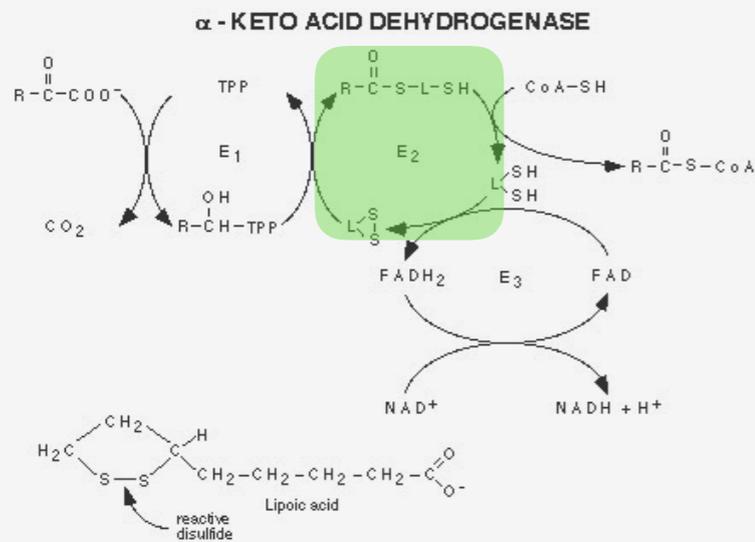
Biotin lipoyl

E3
binding

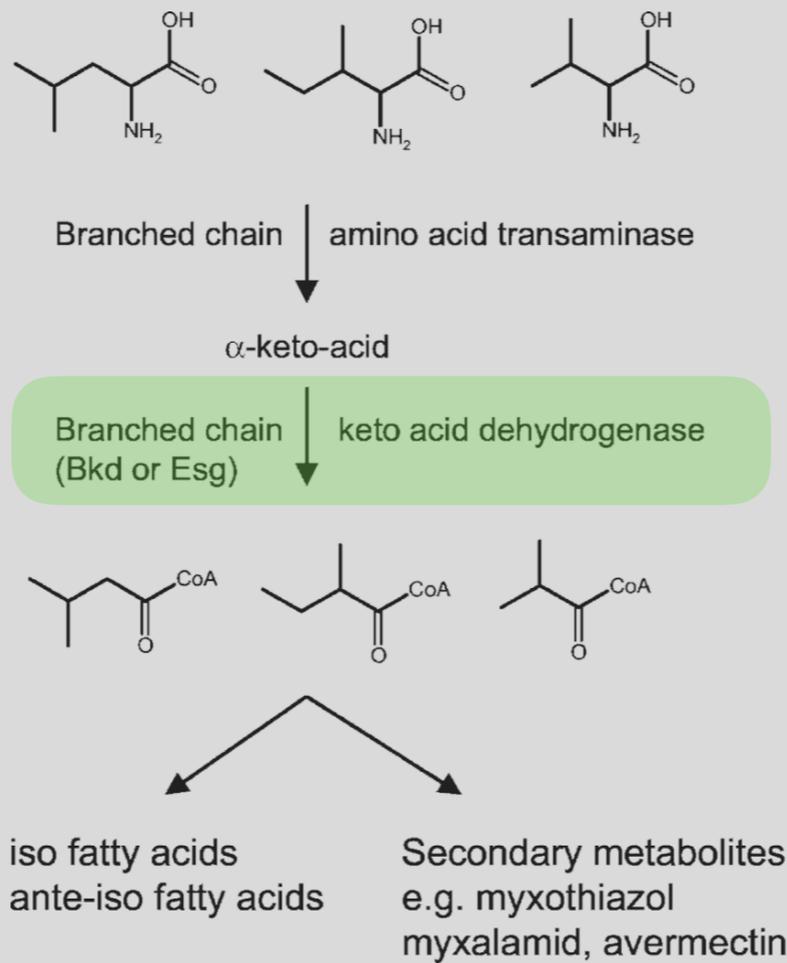
2-oxoacid dehydrogenases
acyltransferase (catalytic domain)

482AA

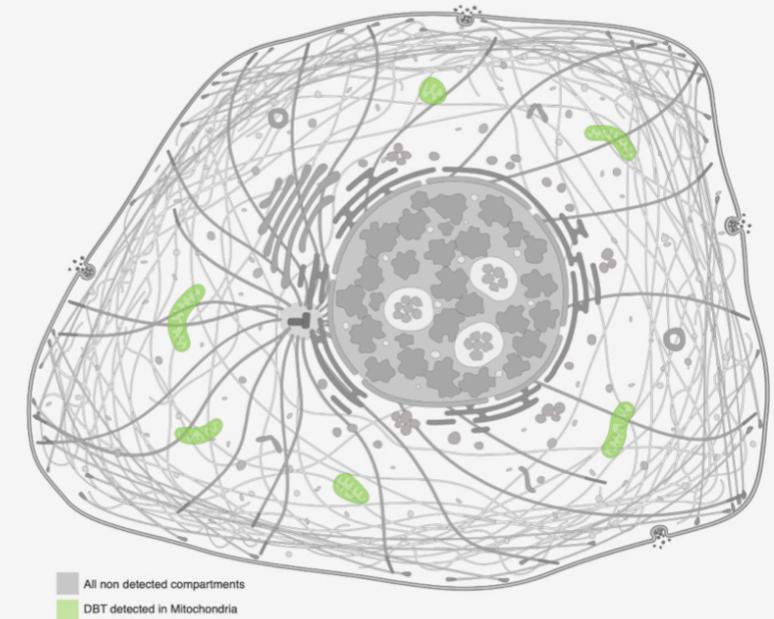
Molecular Function



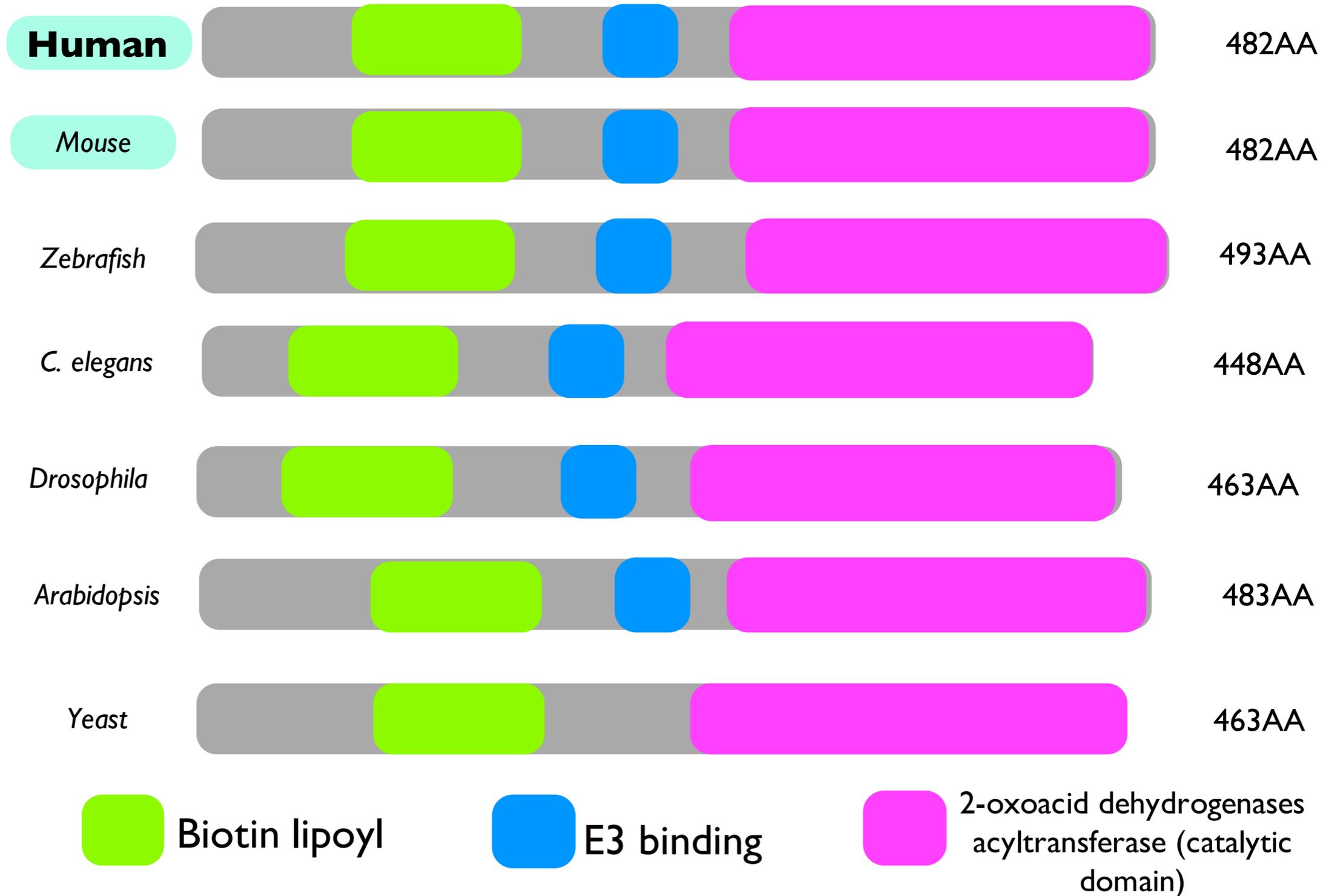
Biological Function



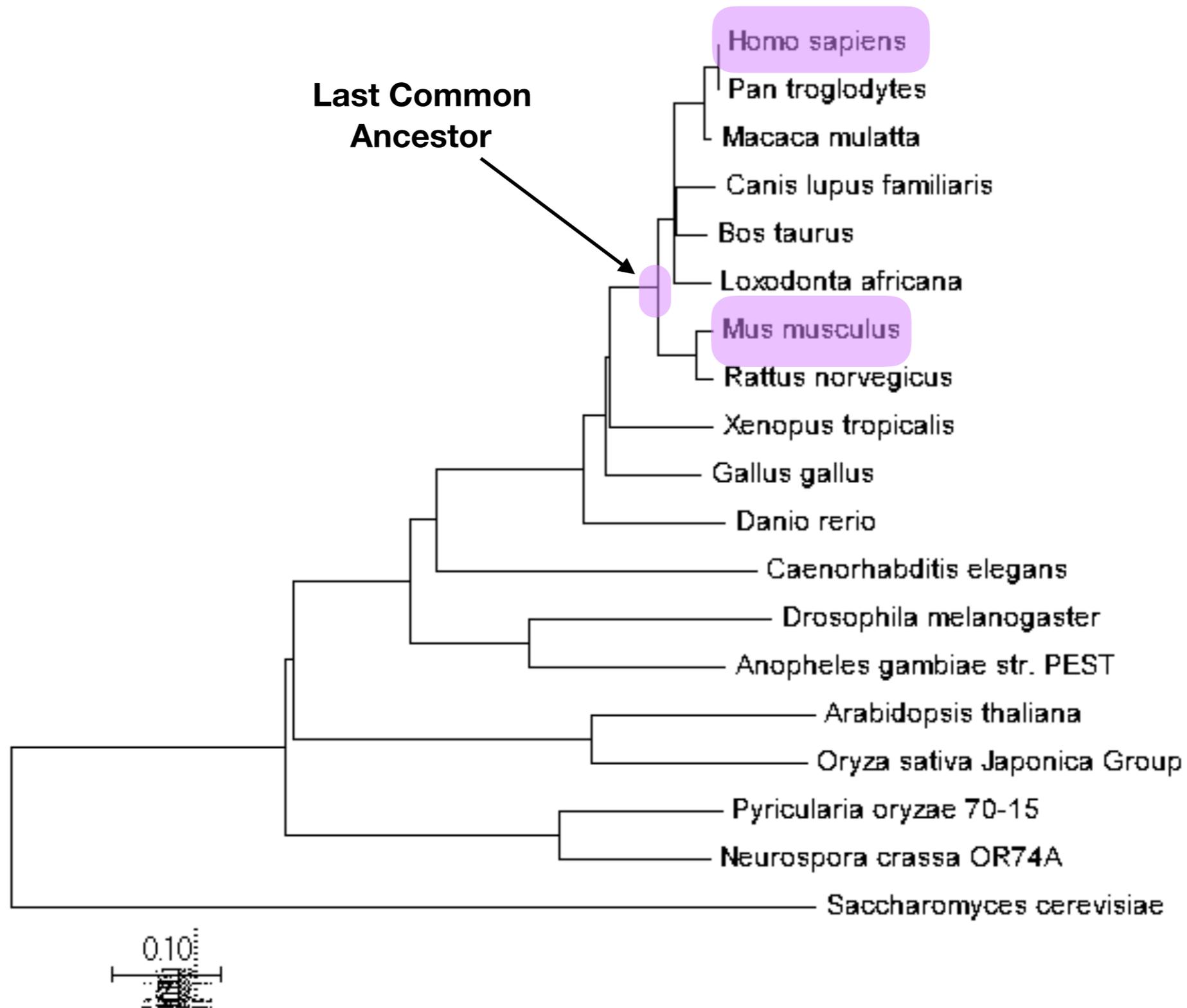
Cellular Function



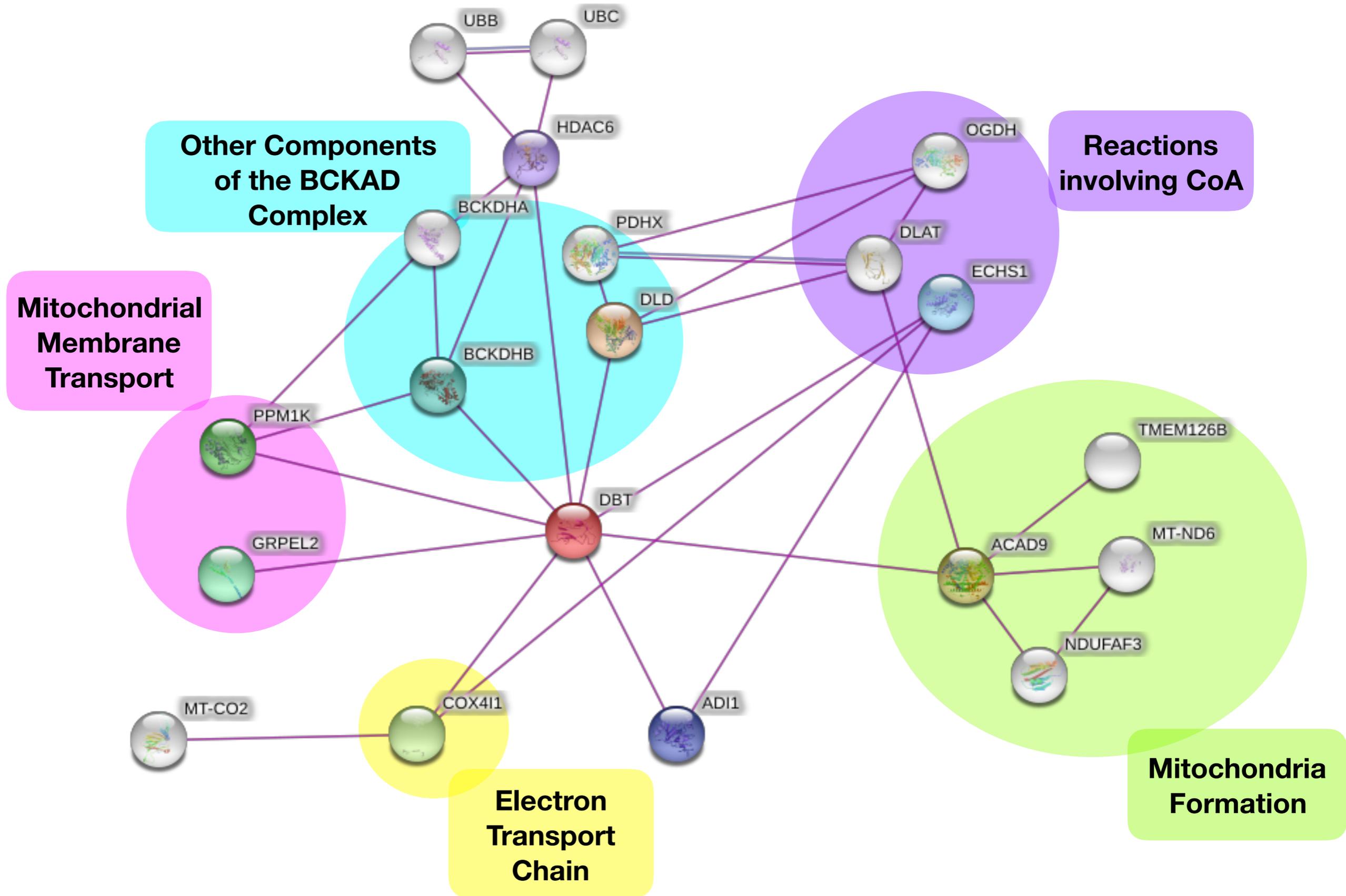
How Well is DBT Conserved?



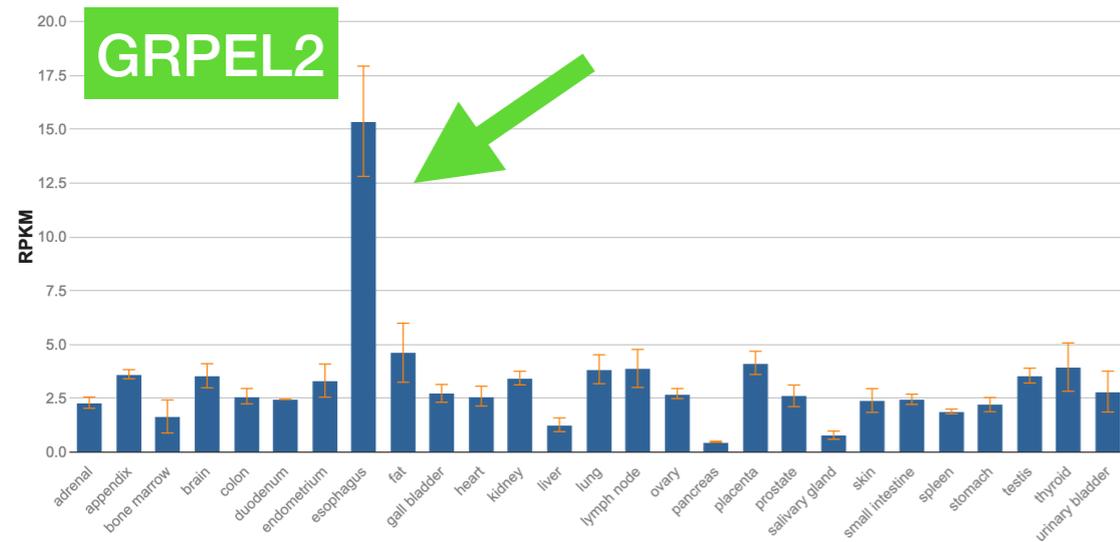
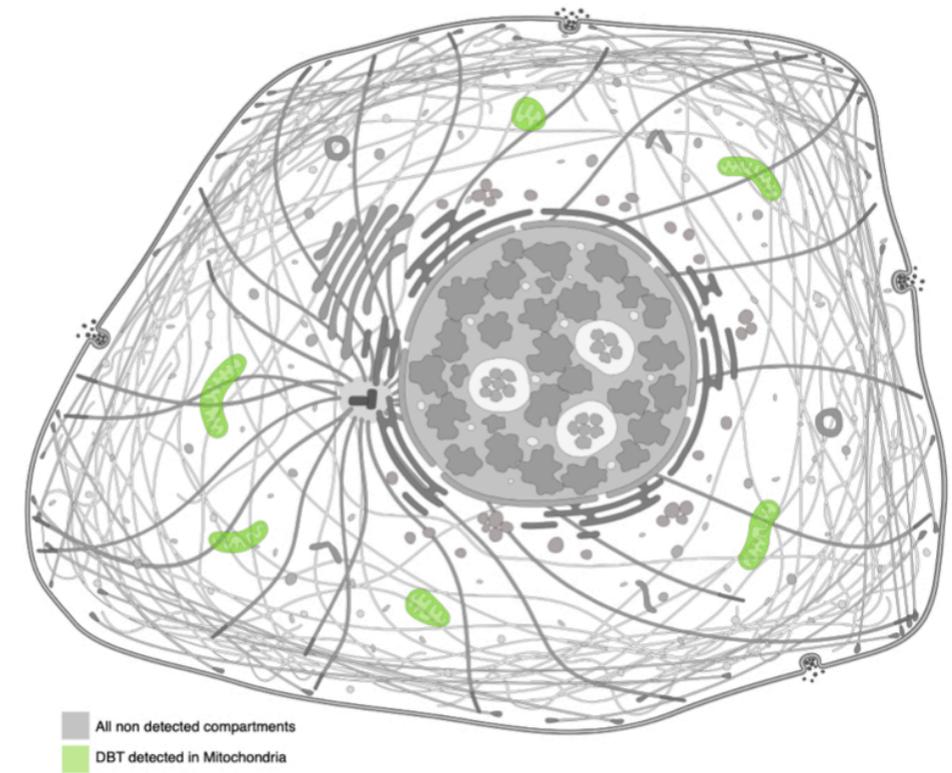
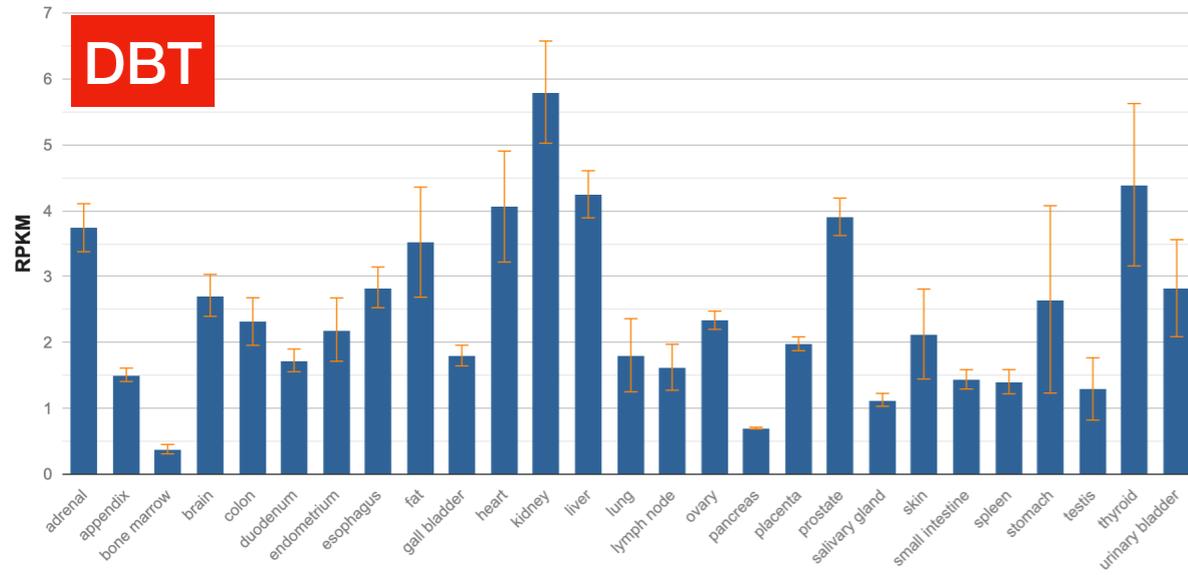
What is the **Phylogenetic Relationship**?



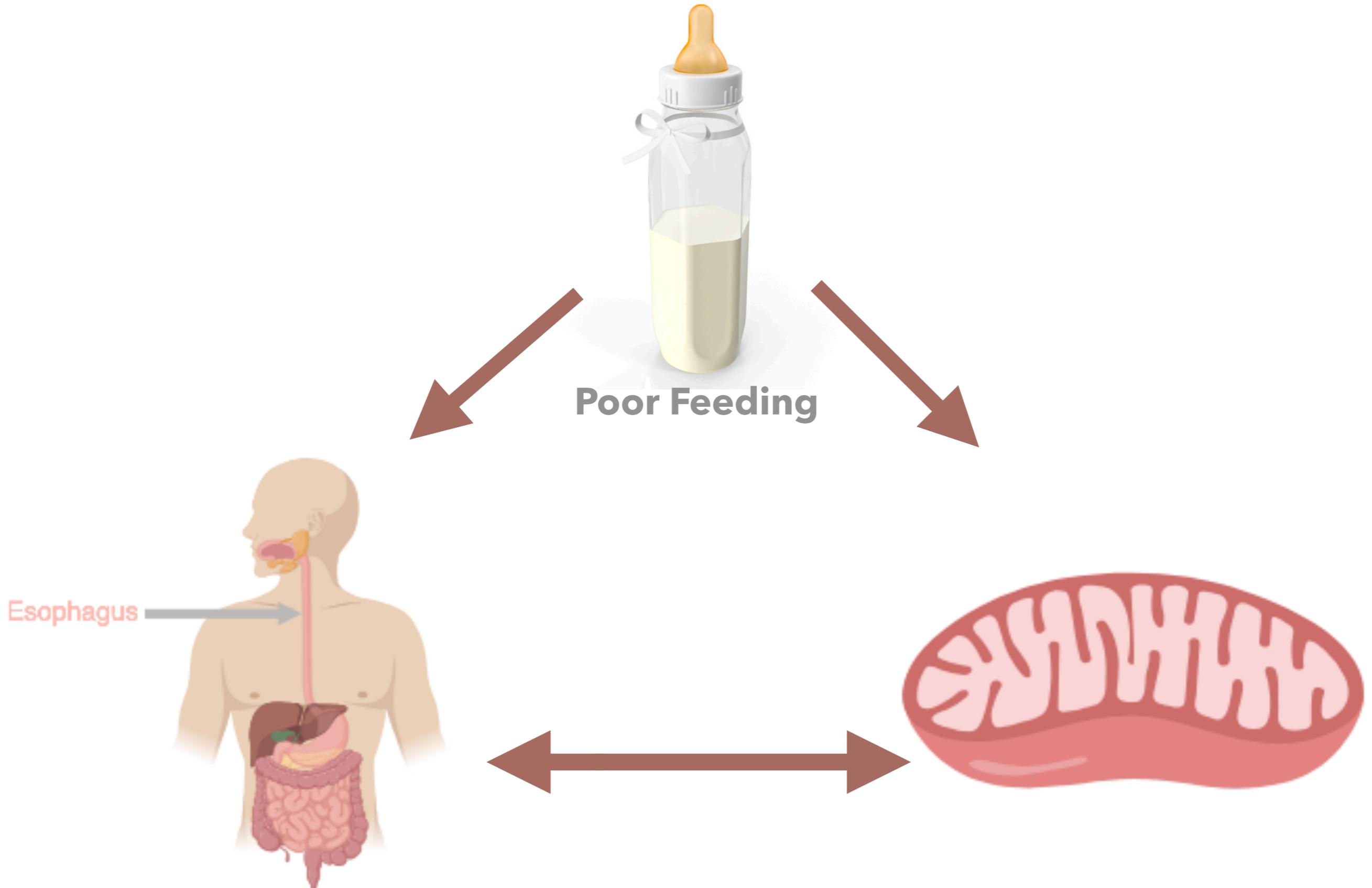
What are the **DBT** protein interactions?



Where do DBT and its interacting proteins localize?



What is the gap in knowledge?



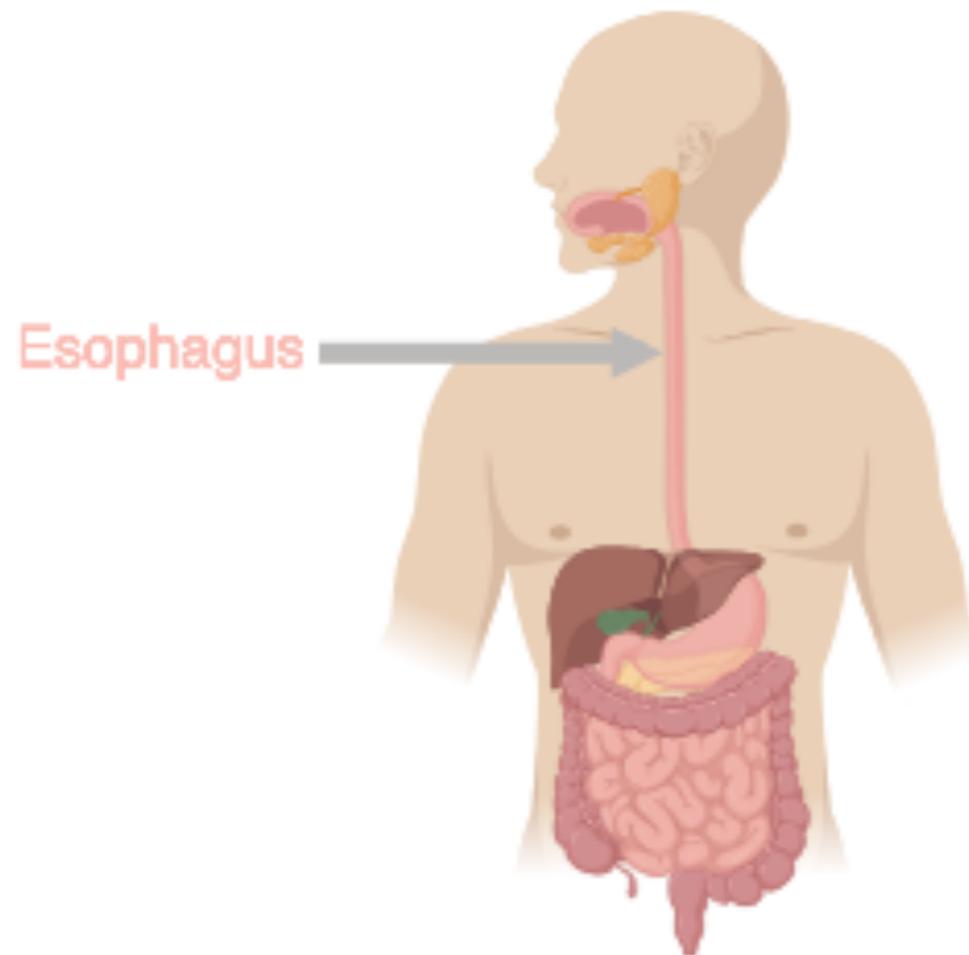
What is the primary goal?

Biotin lipoyl

E3
binding

2-oxoacid dehydrogenases
acyltransferase (catalytic domain)

482AA



Determine how DBT plays a role in esophagus development

What model organism will be used?

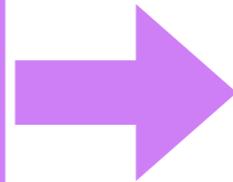
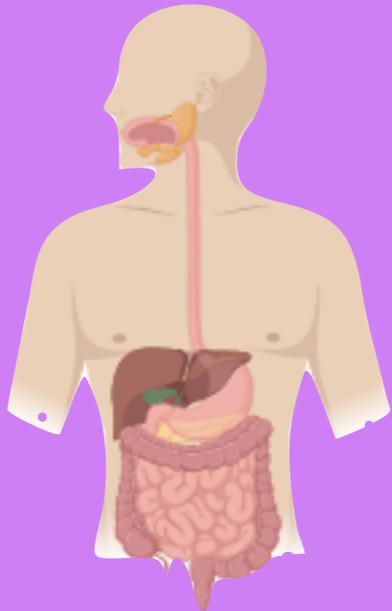


Mus Musculus

How will the primary goal be reached?

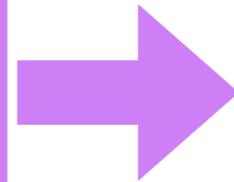
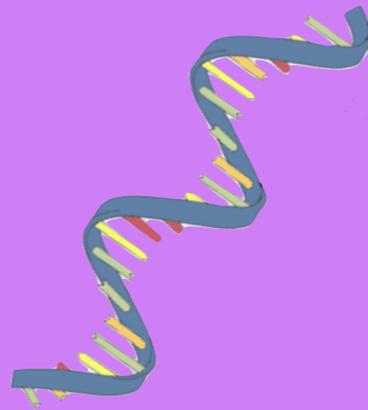
Aim 1:

Identify amino acids that are important for feeding behavior.



Aim 2:

Determine if mitochondrial genes are highly or lowly expressed between wild type and DBT mutant that are important for feeding



Aim 3:

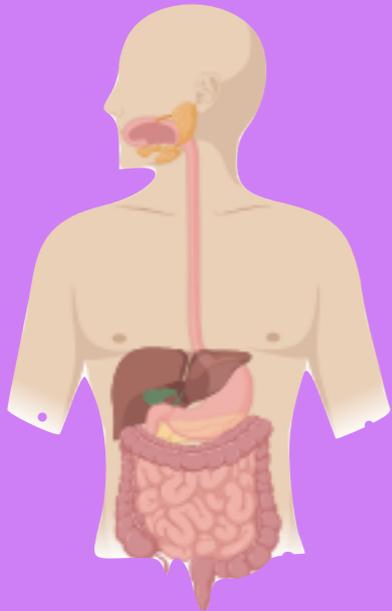
Identify proteins necessary for mitochondrial function and feeding



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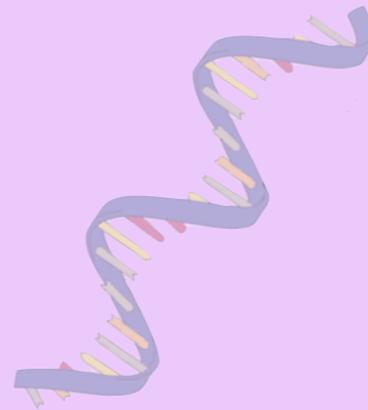
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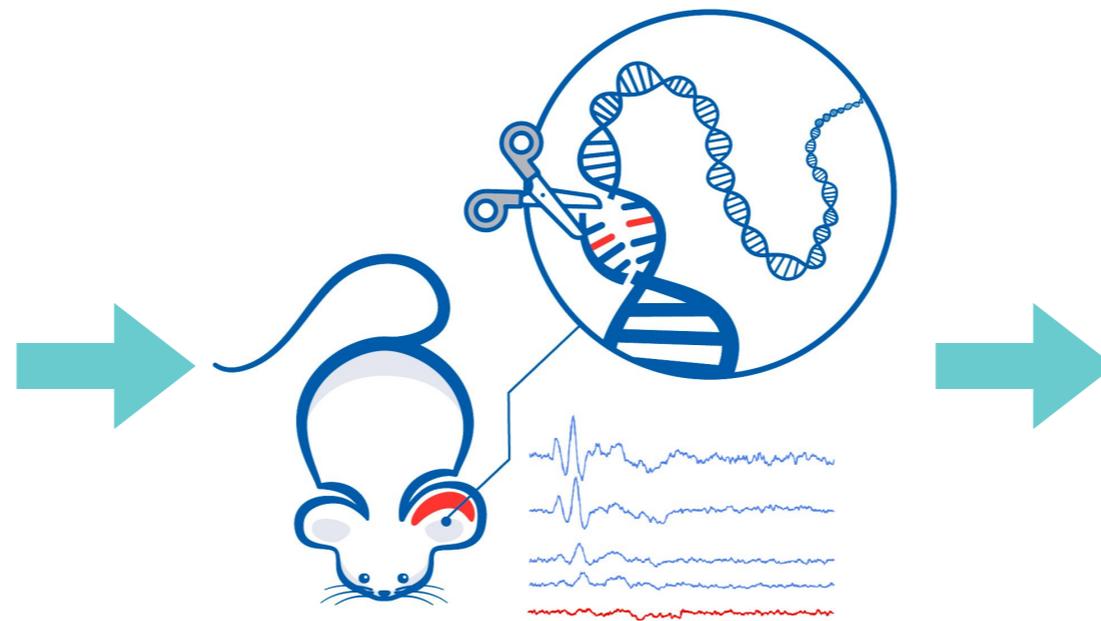
Identify proteins necessary for mitochondrial function and feeding



Aim 1: Identify mutations in the DBT protein that result in poor feeding phenotype

Clustal
Omega

Jalview



CRISPR/Cas9 Editing



Phenotype Assay

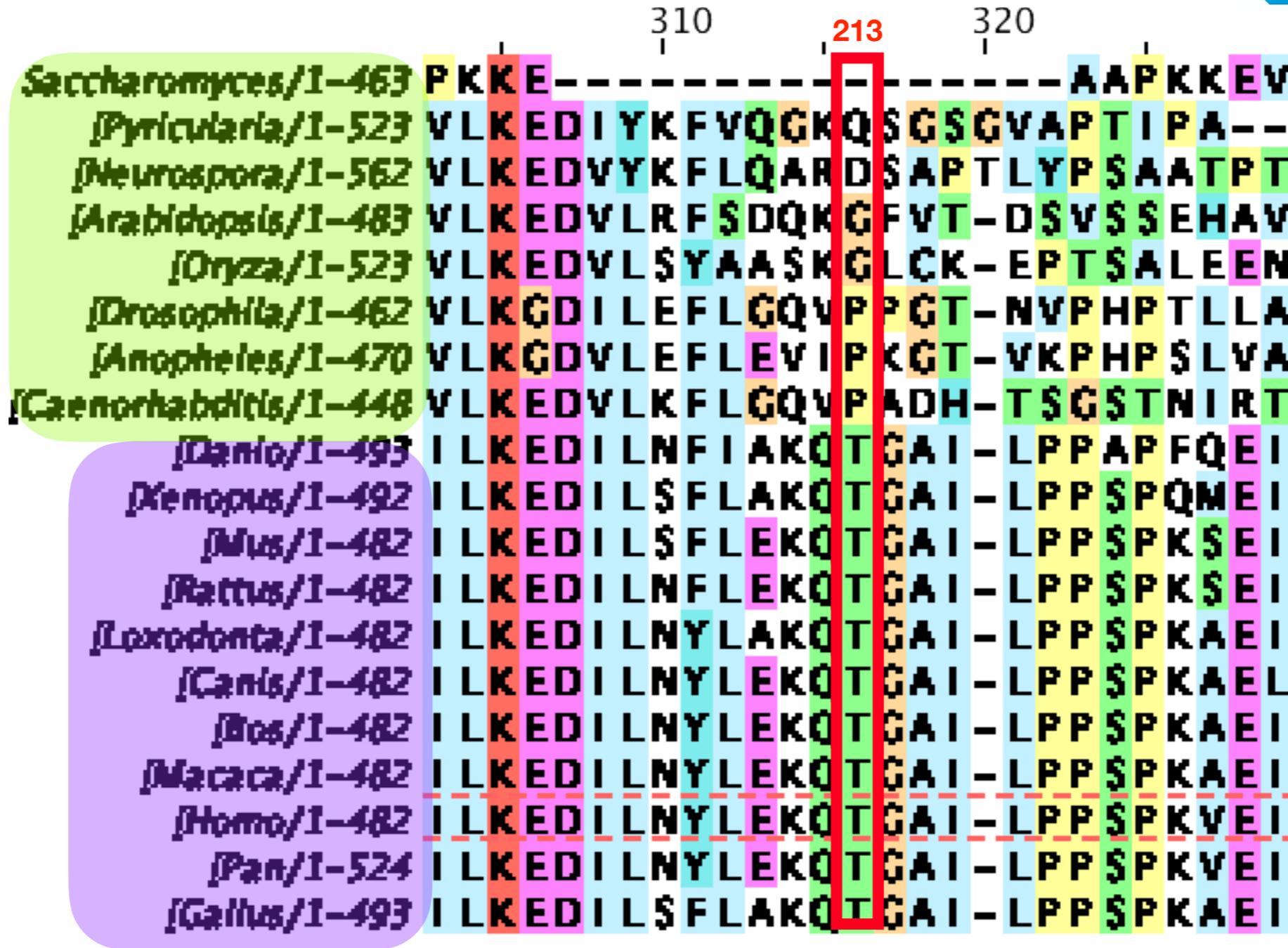
Hypothesis: Amino acids that are conserved in complex digestive organisms, but not simple digestive organisms and are known to be phosphorylated, when mutated will result in a phenotype that exhibits poor feeding and high BCAA levels in urine.

Aim 1: Identify mutations in the DBT protein that result in poor feeding phenotype

Clustal Omega



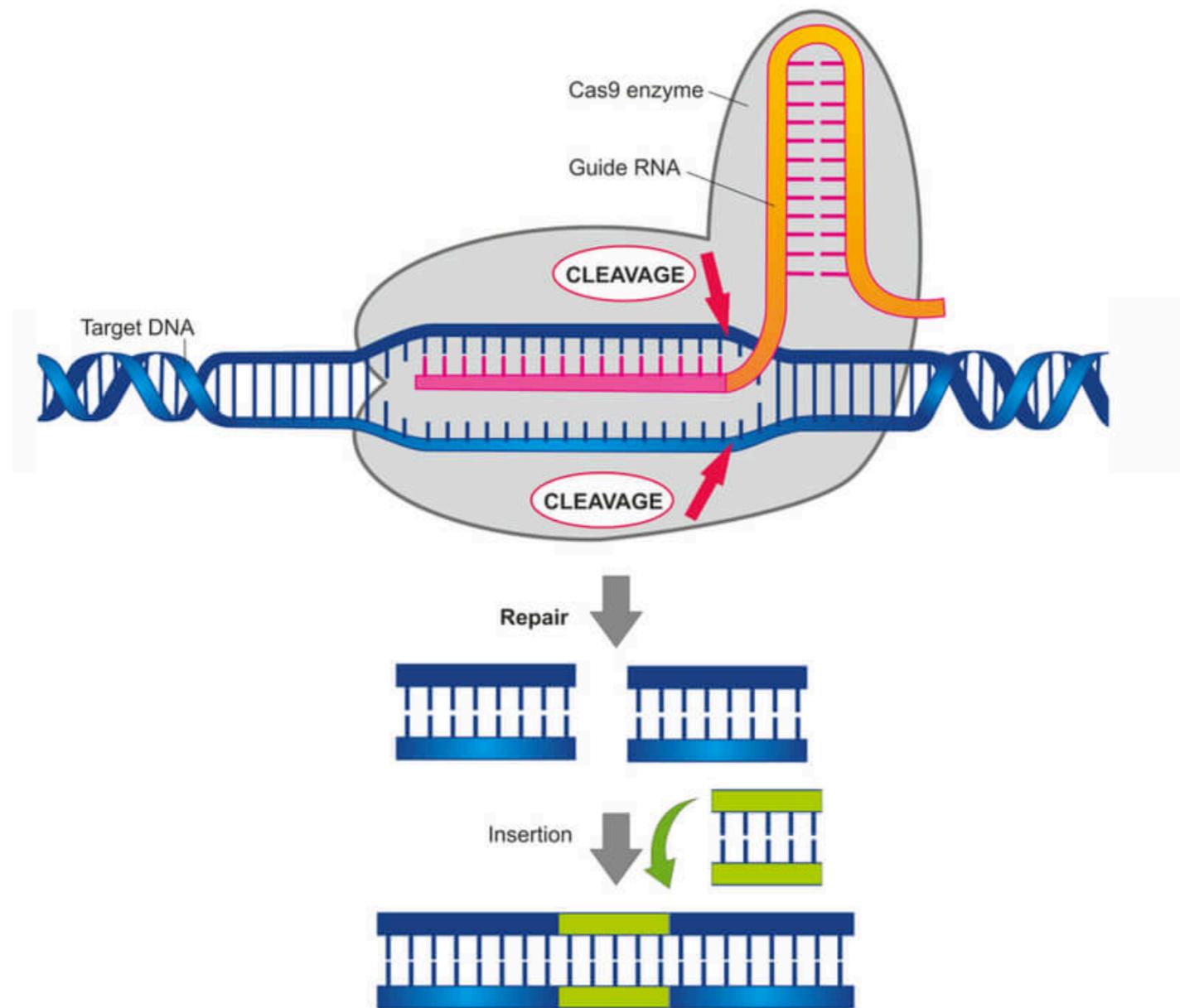
Simple
→
Complex



Sequence 213 T LEKQTGAIL 0.540 PKA YES

Align the sequences to identify amino acids that are **different** between organisms with **simple** and **complex** digestive system

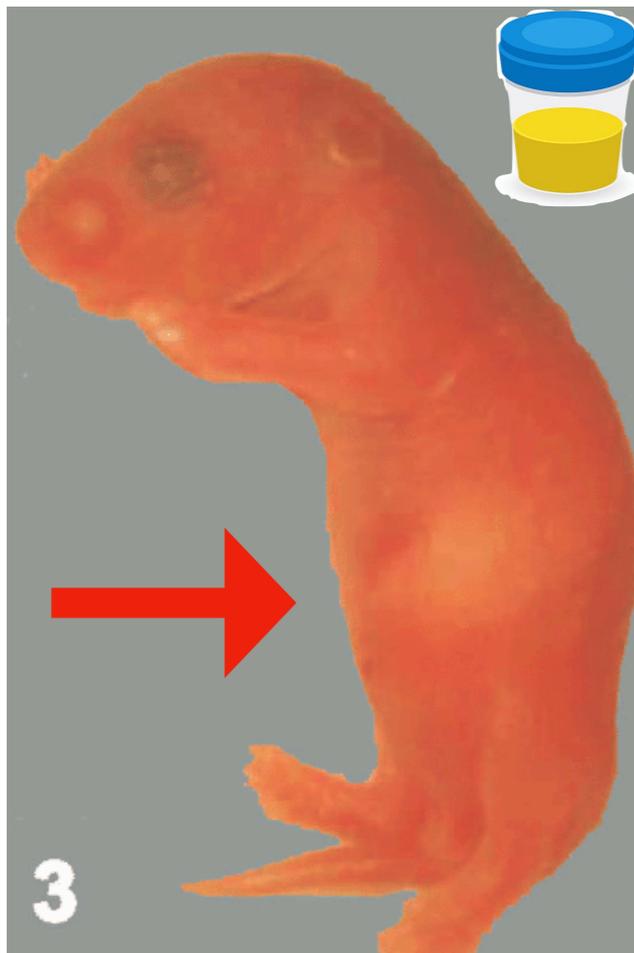
Aim 1: Identify mutations in the DBT protein that result in poor feeding phenotype



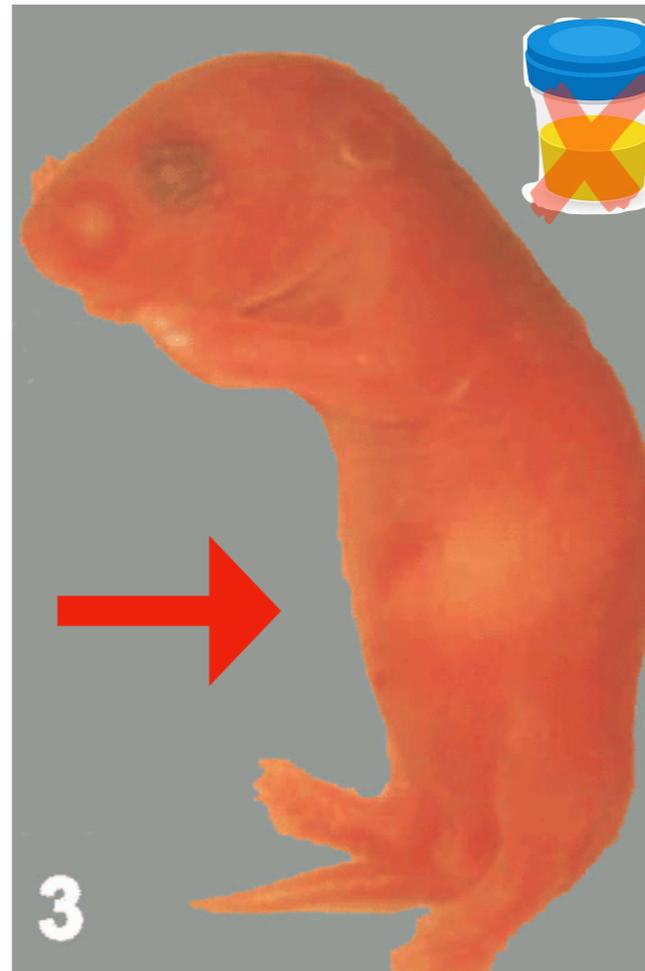
Use CRISPR/Cas9 technology to change amino acids at those sites to an amino acid found in lower level organisms

Aim 1: Identify mutations in the DBT protein that result in poor feeding phenotype

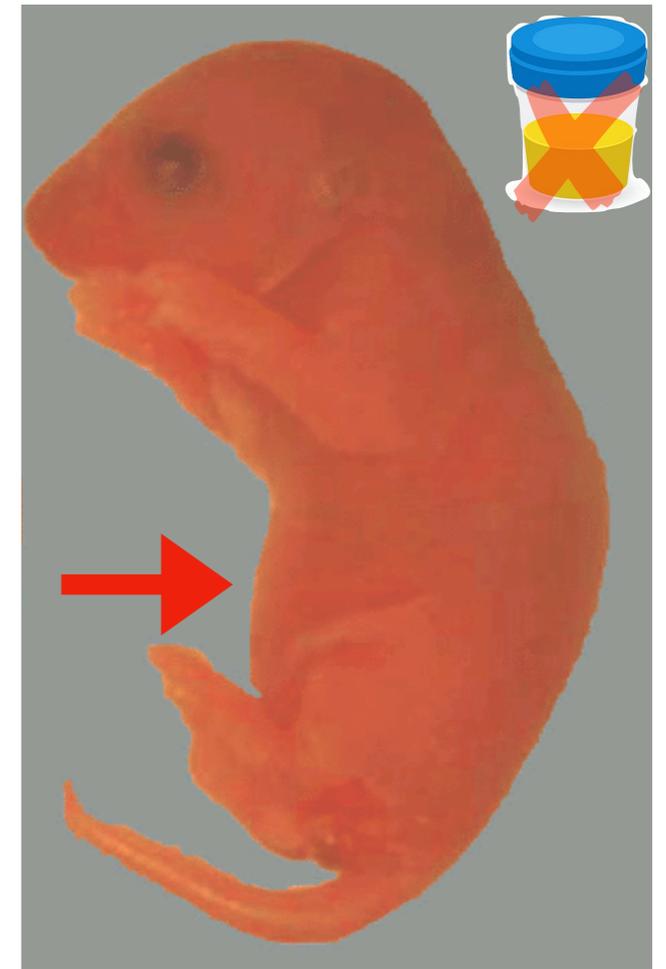
Negative Control



MSUD Mutant



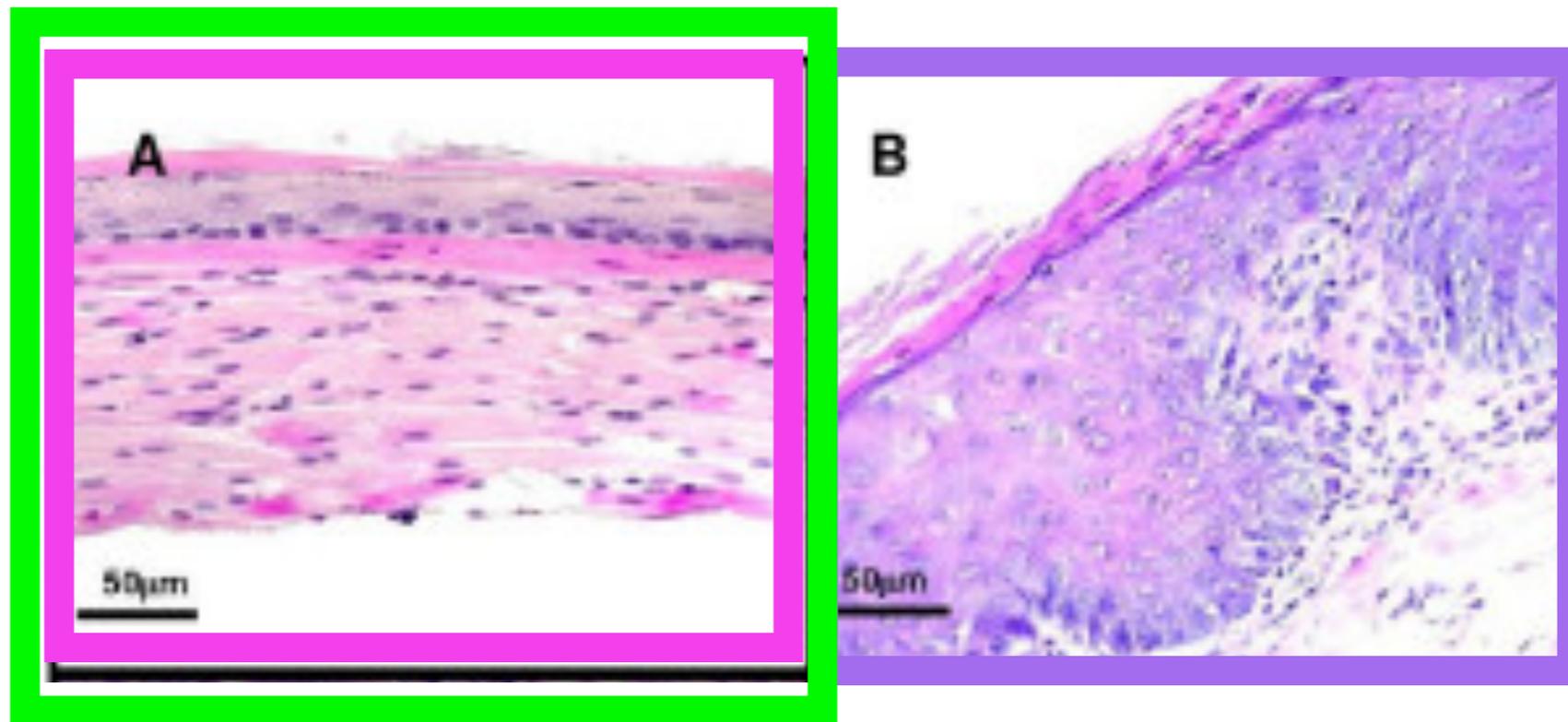
MSUD Mutant +
Poor Feeding



Screen for mice that both exhibit **poor feeding** and have the **urine** phenotype

Aim 1: Identify mutations in the DBT protein that result in poor feeding phenotype

Extraction of Esophageal Tissue:



Hypothesis: Esophageal tissue will be distorted in mice who display the poor feeding phenotype (B) and will not be distorted in mice that do not (A)

How will the primary goal be reached?

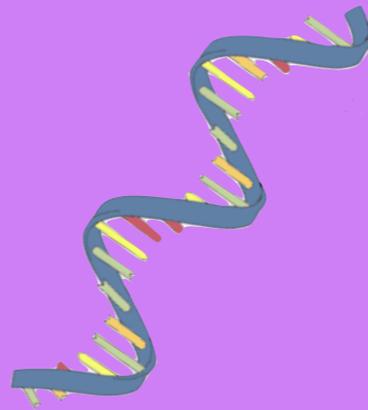
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Identify amino acids that are important for feeding behavior.



Aim 2:

Determine if mitochondrial genes are highly or lowly expressed between wild type and DBT mutant that are important for feeding

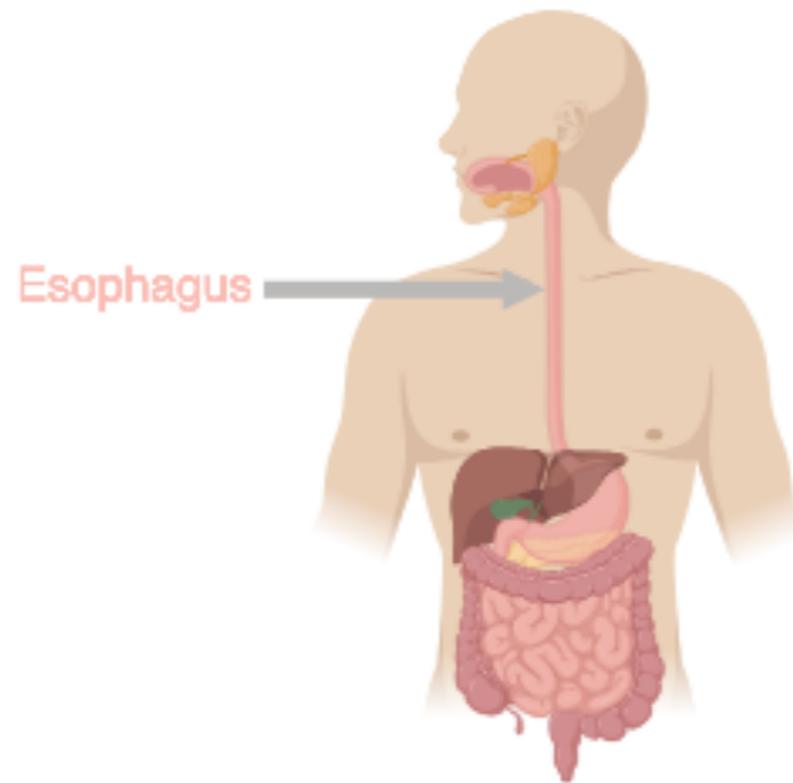


Aim 3:

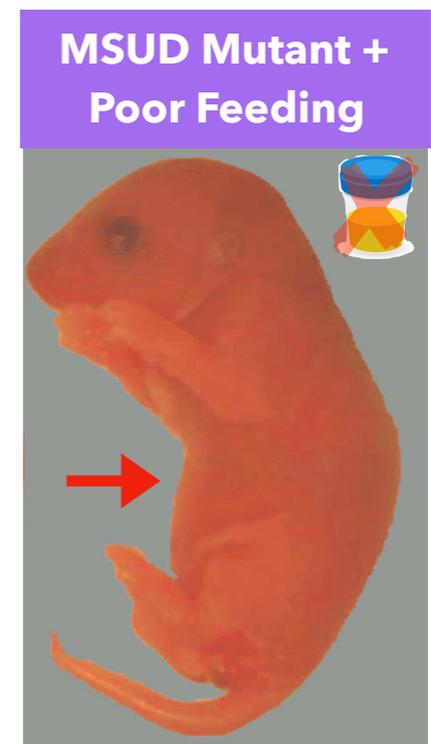
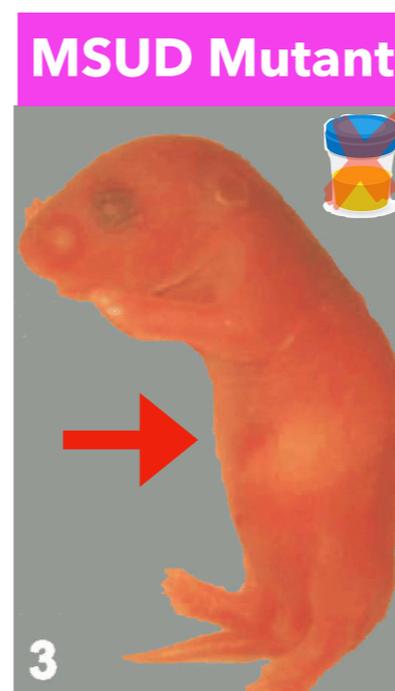
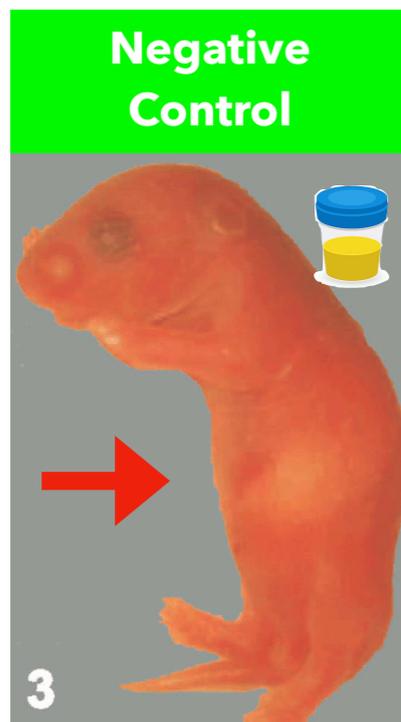
Identify proteins necessary for mitochondrial function and feeding



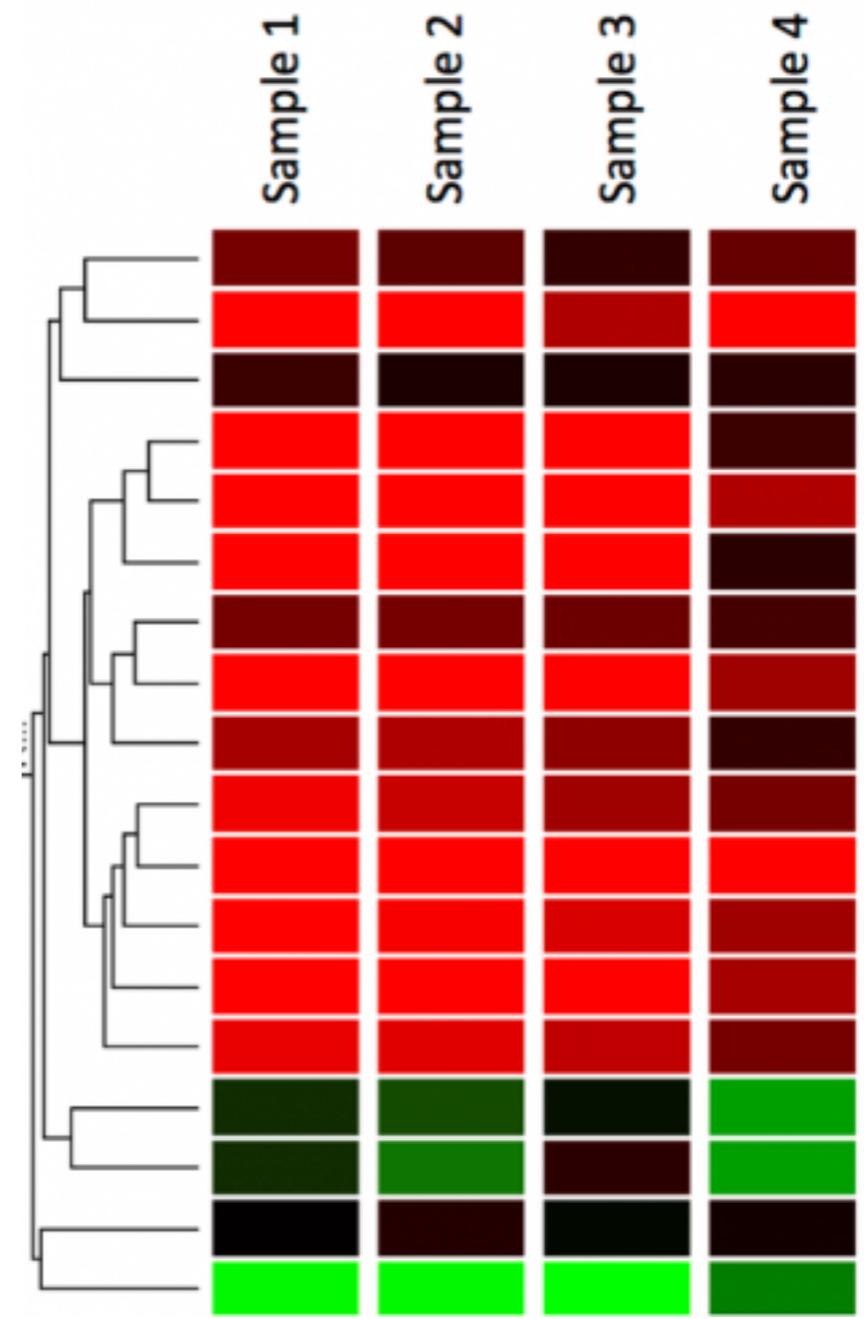
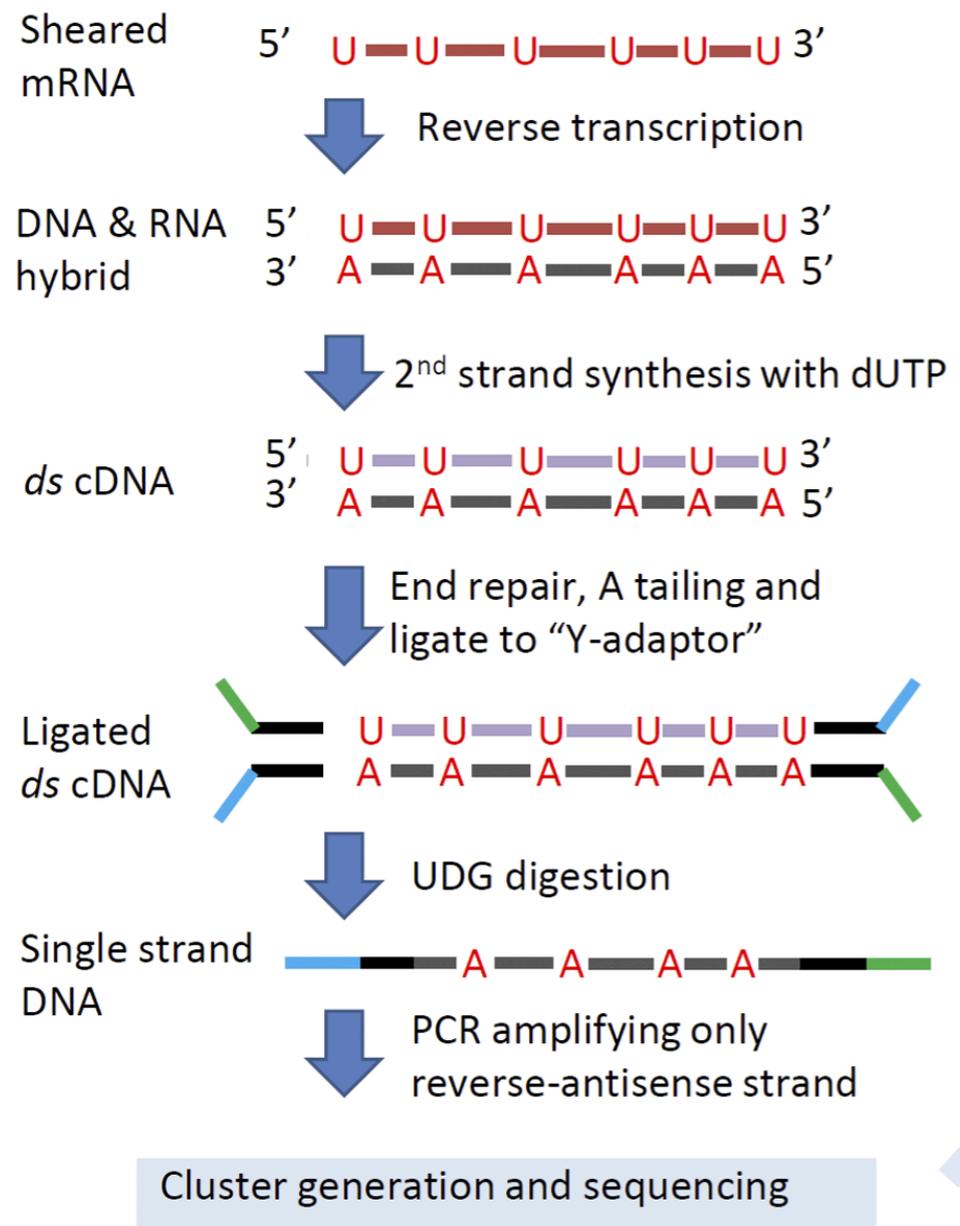
Aim 2: Determine if mitochondrial genes are highly or lowly expressed between wild type and DBT mutant that are important for feeding



Extract **esophageal** tissue from all three test groups

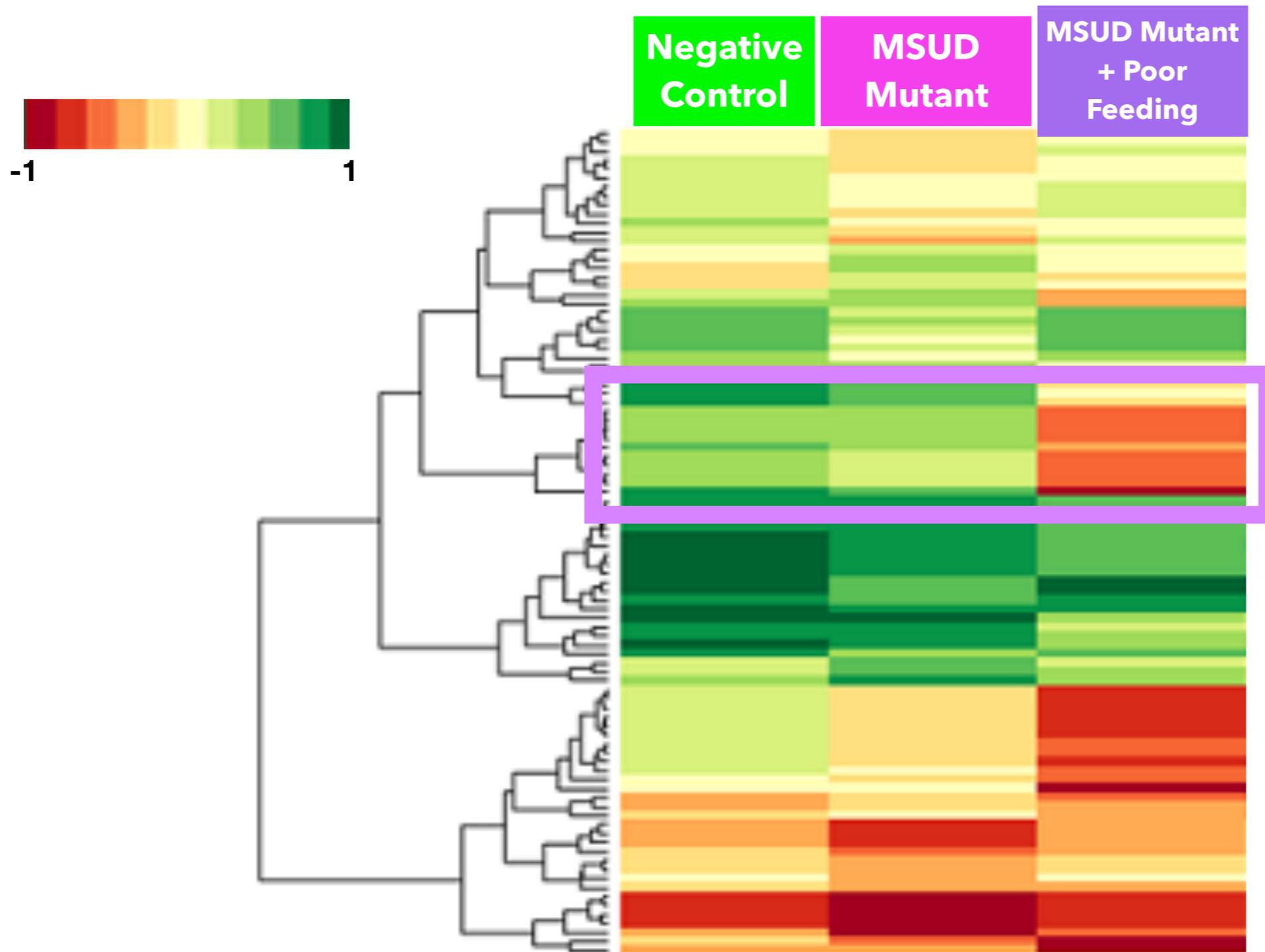


Aim 2: Determine if mitochondrial genes are highly or lowly expressed between wild type and DBT mutant that are important for feeding



RNA-Seq

Aim 2: Determine if mitochondrial genes are highly or lowly expressed between wild type and DBT mutant that are important for feeding

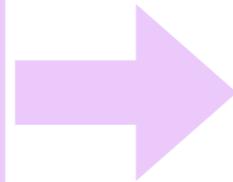


Hypothesis: Transcripts that interact with DBT and are highly localized in the esophagus will be **under expressed** in MSUD mutants with poor feeding

How will the primary goal be reached?

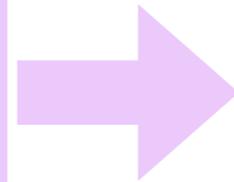
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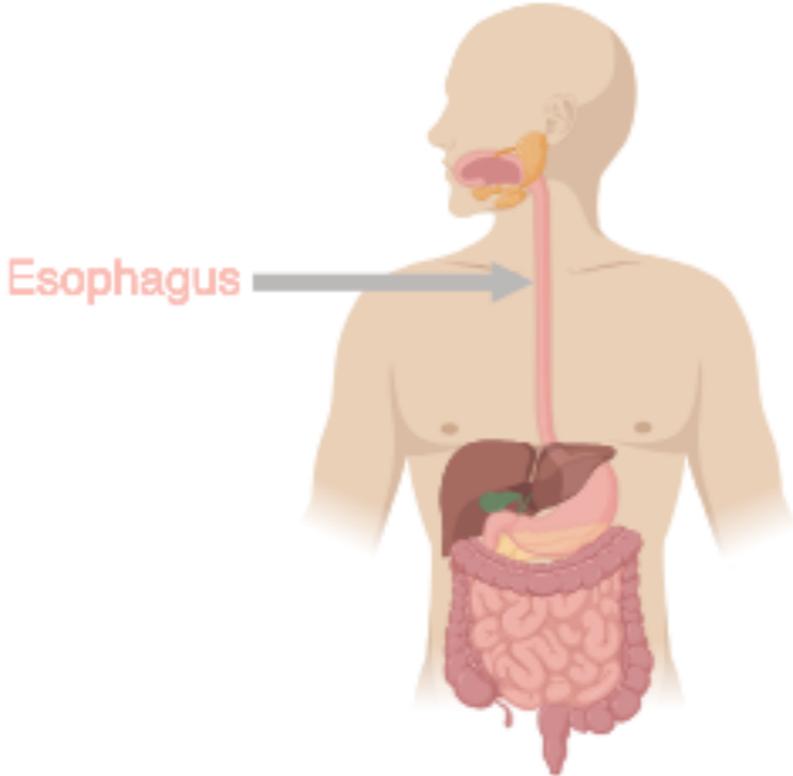


Aim 3:

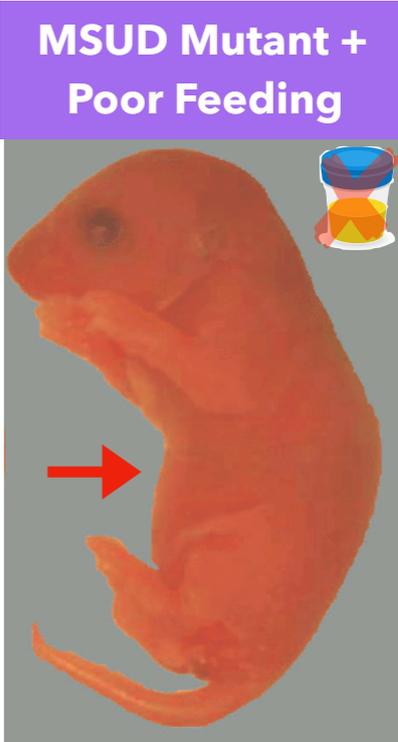
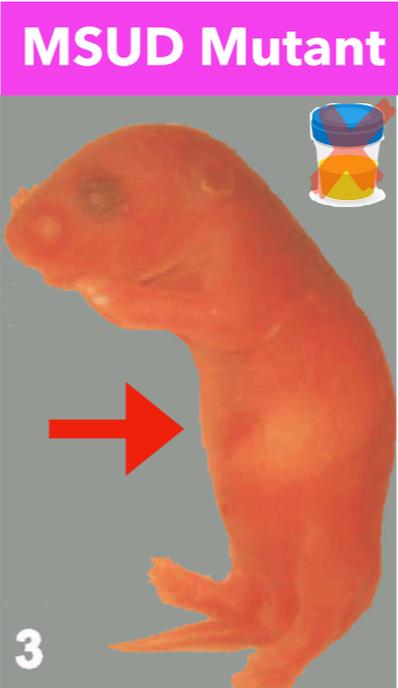
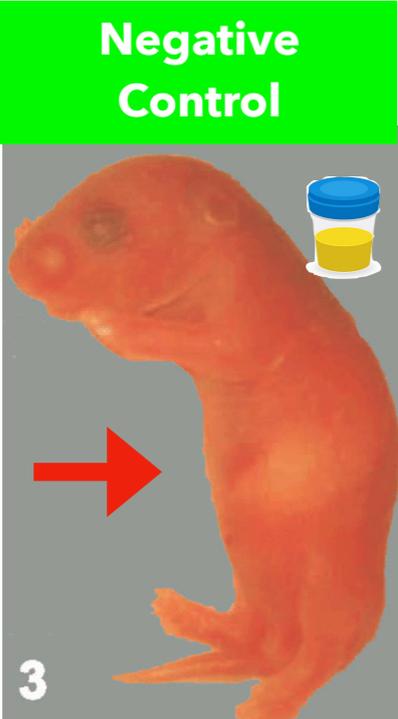
Identify proteins necessary for mitochondrial function and feeding



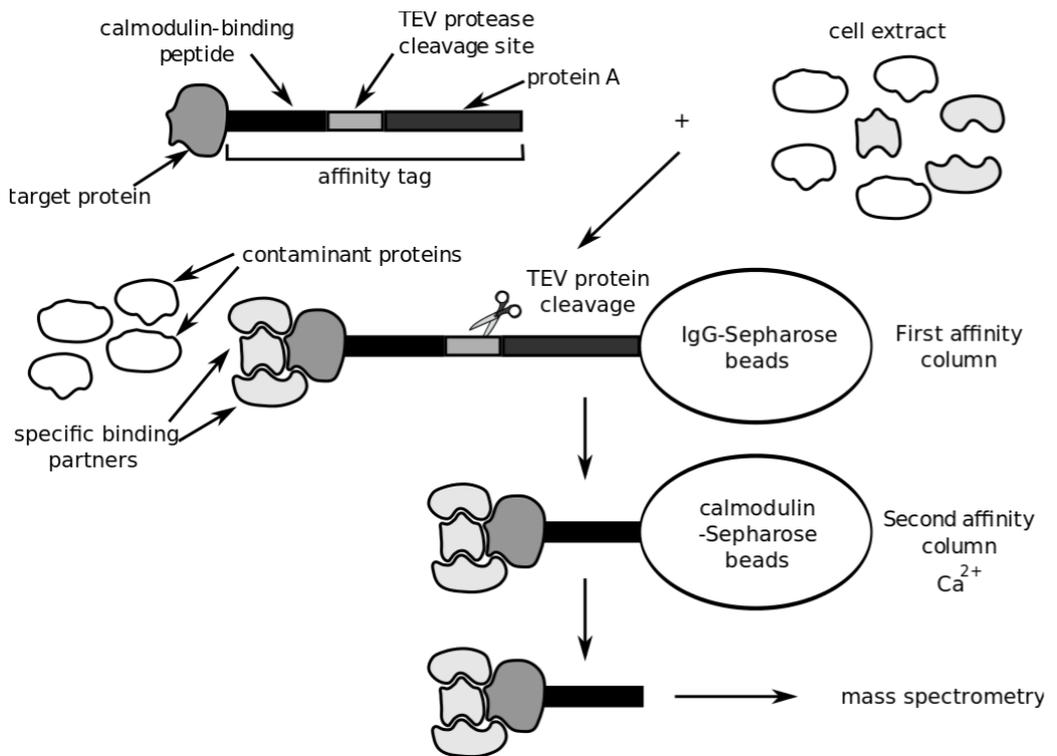
Aim 3: Identify proteins necessary for mitochondrial function and feeding



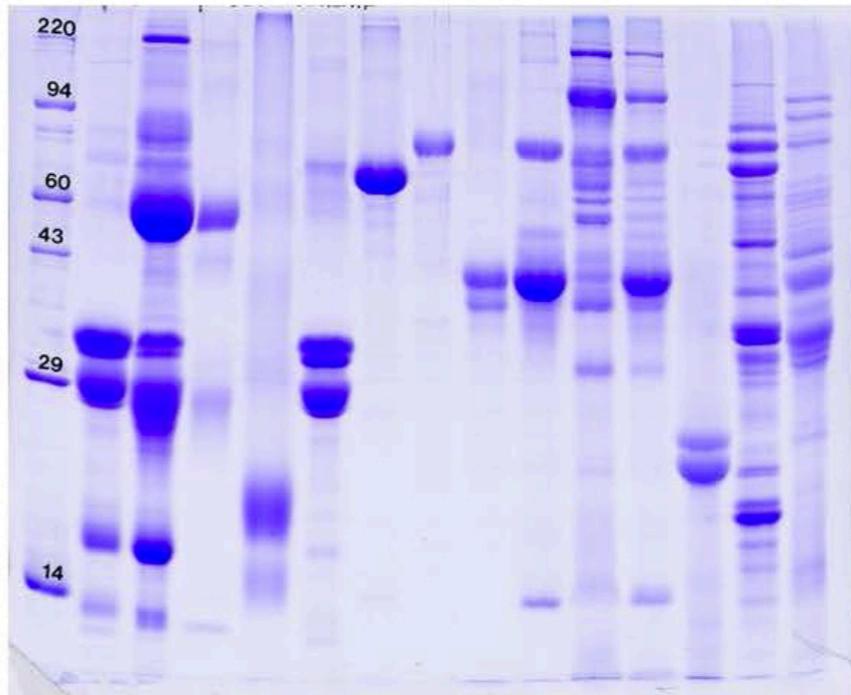
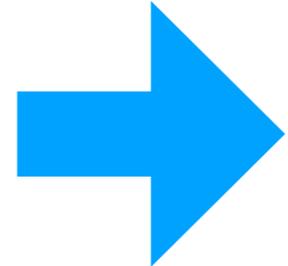
Extract **esophageal** tissue from all three test groups



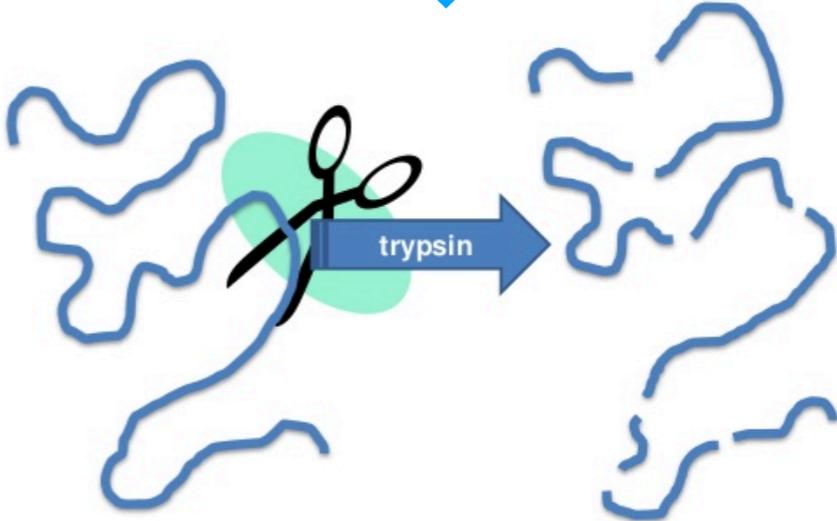
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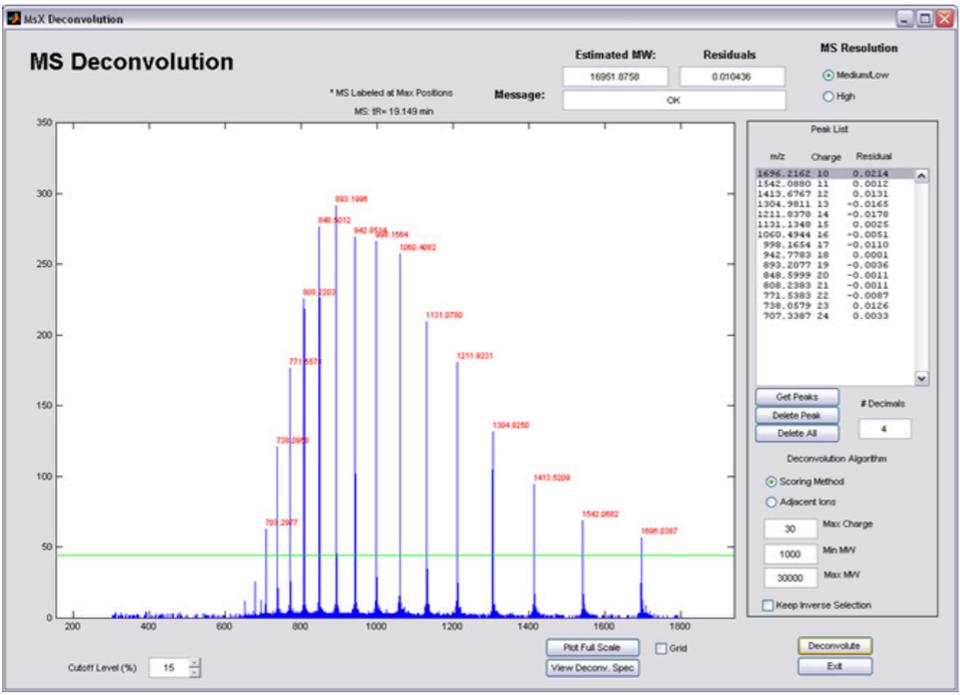
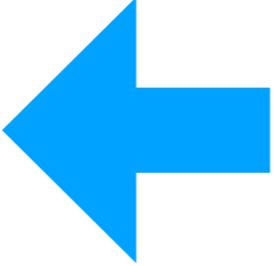
Tandem Affinity Purification



SDS Page Separation

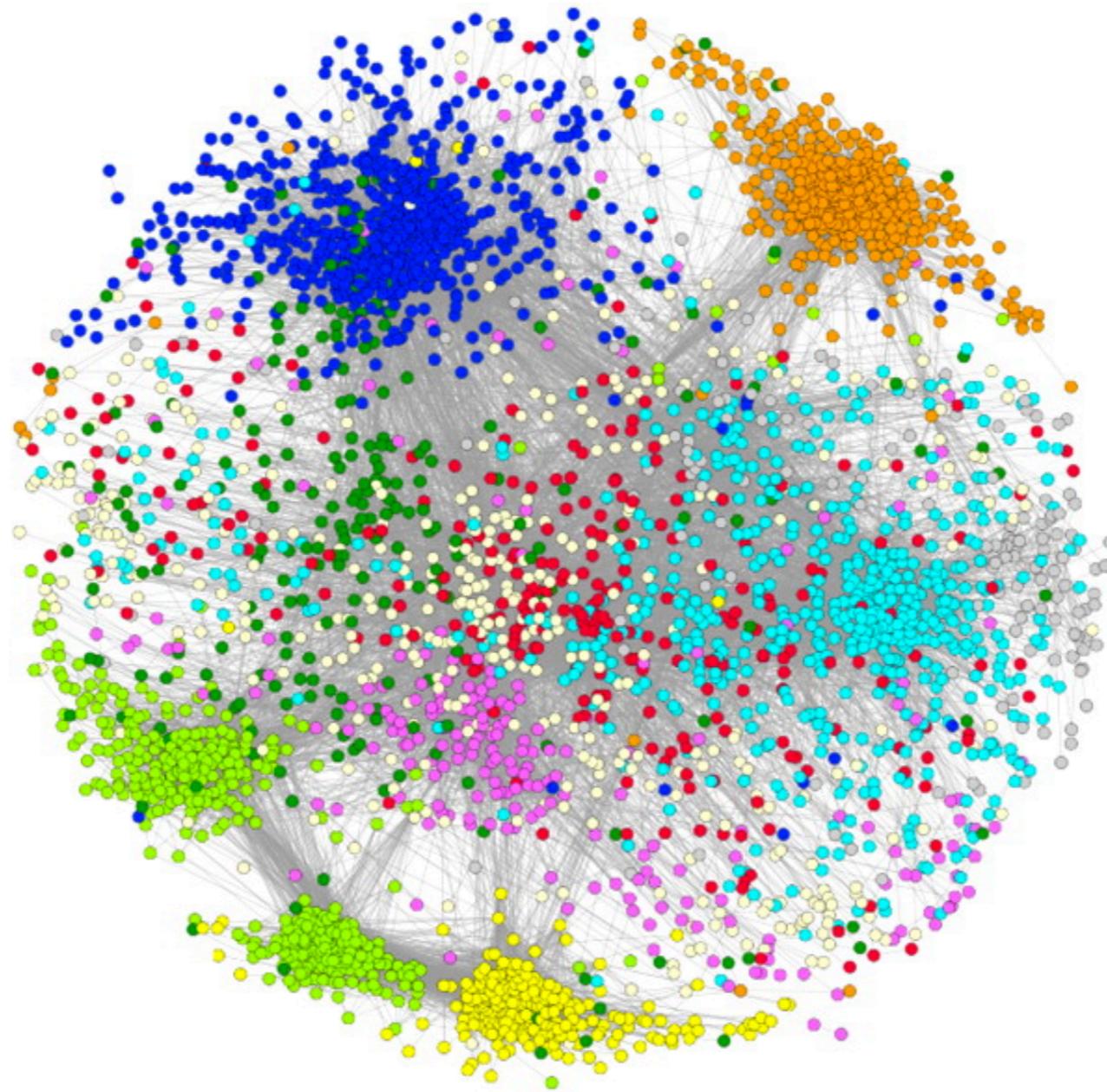


Trypsin Digestion



Mass Spectrometry

Aim 3: Identify proteins necessary for mitochondrial function and feeding

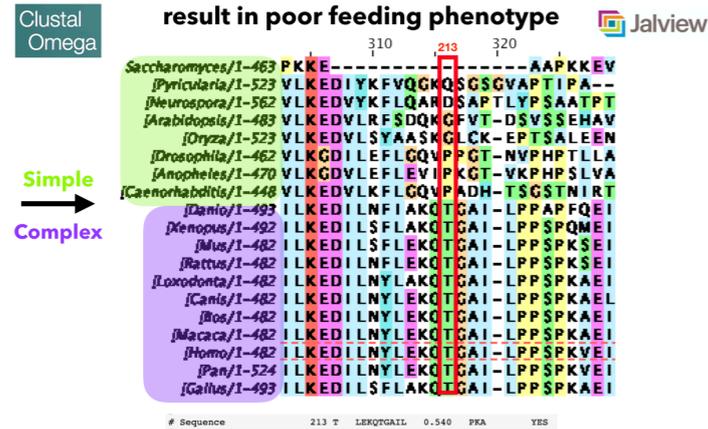


Organize protein interaction network based on pathways and function

Hypothesis: The protein interactions in the mutant DBT with poor feeding will have loss of protein interaction with mitochondrial transport proteins

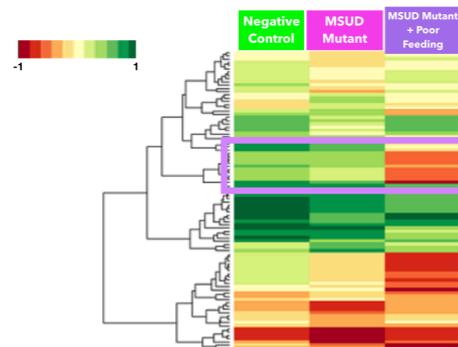
Conclusions

Aim 1: Identify mutations in the DBT protein that result in poor feeding phenotype



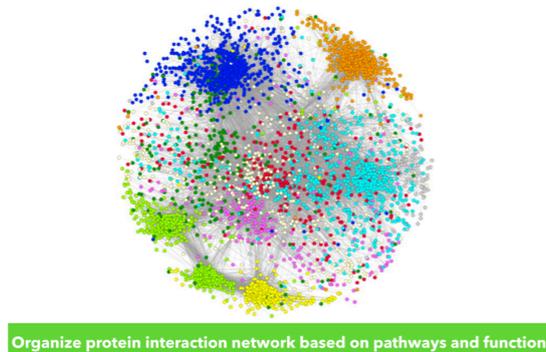
Sequence alignment of homologous proteins allows identification of conserved regions among organisms with complex digestive systems

Aim 2: Determine if mitochondrial genes are highly or lowly expressed between wild type and DBT mutant that are important for feeding



RNAseq identifies transcript level changes in esophageal cells when mutant for DBT

Aim 3: Identify proteins necessary for mitochondrial function and feeding



Proteomic analysis allows for identification of protein interaction changes when mutant for DBT

Image References

Title Slide: <https://cbmaplefarm.com/shop/category/vermont-maple-syrup/>

Slide 1: <https://diyhealthacademy.com/newborn-screening-tests-state-requires/> <https://www.hrsa.gov/sites/default/files/hrsa/advisory-committees/heritable-disorders/rusp/rusp-uniform-screening-panel.pdf>

Slide 2: <https://www.newbornscreening.info/Parents/aminoaciddisorders/Images/MSUD.gif>

Slide 3: <https://cdn.images.express.co.uk/img/dynamic/11/590x/Urine-smell-841566.jpg> <https://dovemed-prod-k8s.s3.amazonaws.com/media/images/msud-Image004.width-750.png> https://www.healthline.com/hlcmsresource/images/imce/feeding-tube-infants_thumb.jpg

Slide 4: http://education.med.nyu.edu/courses/molecular/AminoAcids06/questions/aa_answers/ans23a.html https://www.researchgate.net/profile/Rolf_Mueller4/publication/11290128/figure/fig1/AS:394580399804421@1471086706637/Degradation-of-leucine-isoleucine-and-valine-via-the-branched-chain-keto-acid.png <https://www.proteinatlas.org/ENSG00000137992-DBT/cell>

Slide 6: Generated by Mega Software

Slide 7: <https://string-db.org/cgi/network.pl?taskId=ldKnOD6d1YVA>

Slide 8: <https://www.ncbi.nlm.nih.gov/gene/1629> <https://www.ncbi.nlm.nih.gov/gene/134266> <https://www.proteinatlas.org/ENSG00000137992-DBT/cell>

Slide 9: <http://atlas-content-cdn.pixelsquid.com/stock-images/white-baby-bottle-half-full-ENmmZkB-600.jpg> biorender.com

Slide 10: biorender.com

Slide 11: <https://encrypted-tbn0.gstatic.com/images?q=tbn:ANd9GcTQt1xRRpXmWYootdNbnWVYW17Qnde1hSG0vTFqxbRHTr5TnPGs>

Slide 12,13,19, 23: [biorender.com https://www2.le.ac.uk/projects/vgec/diagrams/115-mrna.gif](https://www2.le.ac.uk/projects/vgec/diagrams/115-mrna.gif)

Slide 14: <https://lw-static-files.s3.amazonaws.com/public/logos/2997.png> http://www.jalview.org/Web_Installers/InstData/com/zerog/ia/installer/images/Splash.gif

<https://pbs.twimg.com/media/DWgv7eCU8AAact9.jpg> <https://cdn.the-scientist.com/assets/articleNo/32610/ilmg/6309/663848b7-98da-4684-a5ac-5d07d1464c84-lt1.jpg>

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<http://www.cbs.dtu.dk/cgi-bin/webface2.fcgi?jobid=5CA66549000010A4B31756DC&wait=20>

Slide 16: <https://294305267s7hqfks2cfh08ip-wpengine.netdna-ssl.com/wp-content/uploads/2017/05/crispr-cas9-drug-discovery2-1.jpg>

Slide 17, 20, 24: [https://media.istockphoto.com/vectors/urine-test-vector-id495099954?](https://media.istockphoto.com/vectors/urine-test-vector-id495099954?k=6&m=495099954&s=612x612&w=0&h=UCor4LAlkUsFlpNcLpQDL1gB1r46H8u3IPt3pHaJ8ZI=)

https://www.researchgate.net/profile/Brad_Bolon/publication/23679831/figure/fig3/AS:601637286580224@1520452915347/Neonatal-mice-postnatal-day-1-Relative-to-its-Clc-wild-type-littermate-left-the-Clc.png biorender.com

Slide 18: https://www.researchgate.net/figure/Histopathology-of-mouse-esophagus-after-reflux-surgery-A-In-the-non-operated-control_fig3_26690937

Slide 21: [https://www.google.com/search?](https://www.google.com/search?q=transcriptomics+heat+map&source=lnms&tbm=isch&sa=X&ved=0ahUKEwi3zoWr4bfhAhUMw4MKHc7KBbQQ_AUIDigB&biw=1269&bih=623#imgcr=NkG4AM3I7huskM)

[q=transcriptomics+heat+map&source=lnms&tbm=isch&sa=X&ved=0ahUKEwi3zoWr4bfhAhUMw4MKHc7KBbQQ_AUIDigB&biw=1269&bih=623#imgcr=NkG4AM3I7huskM](https://www.google.com/search?q=transcriptomics+heat+map&source=lnms&tbm=isch&sa=X&ved=0ahUKEwi3zoWr4bfhAhUMw4MKHc7KBbQQ_AUIDigB&biw=1269&bih=623#imgcr=NkG4AM3I7huskM): <https://journals.plos.org/plosone/article/file?id=10.1371/journal.pone.0026426.g002&type=large>

Slide 22: https://newonlinecourses.science.psu.edu/stat555/sites/onlinecourses.science.psu.edu.stat555/files/cluster/heatmap_01/index.png

Slide 25: [https://upload.wikimedia.org/wikipedia/commons/thumb/9/96/Principle_of_Tandem_Affinity_Purification.svg/1024px-](https://upload.wikimedia.org/wikipedia/commons/thumb/9/96/Principle_of_Tandem_Affinity_Purification.svg/1024px-Principle_of_Tandem_Affinity_Purification.svg.png)

[Principle_of_Tandem_Affinity_Purification.svg.png](https://upload.wikimedia.org/wikipedia/commons/thumb/9/96/Principle_of_Tandem_Affinity_Purification.svg.png) <https://kendricklabs.com/wp-content/uploads/2017/03/bigger-size-gel-image-nutrition.jpg> [https://](https://image.slidesharecdn.com/1-141219000811-conversion-gate02/95/1proteomics-coursework3-dec2012aky-19-638.jpg?cb=1418949297)

image.slidesharecdn.com/1-141219000811-conversion-gate02/95/1proteomics-coursework3-dec2012aky-19-638.jpg?cb=1418949297 <http://www.msmetrix.com/wp-content/uploads/2013/05/ProteinDeconvolution.jpg>

Slide 26: <http://www.g3journal.org/content/ggg/2/4/453/F5.large.jpg>

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