

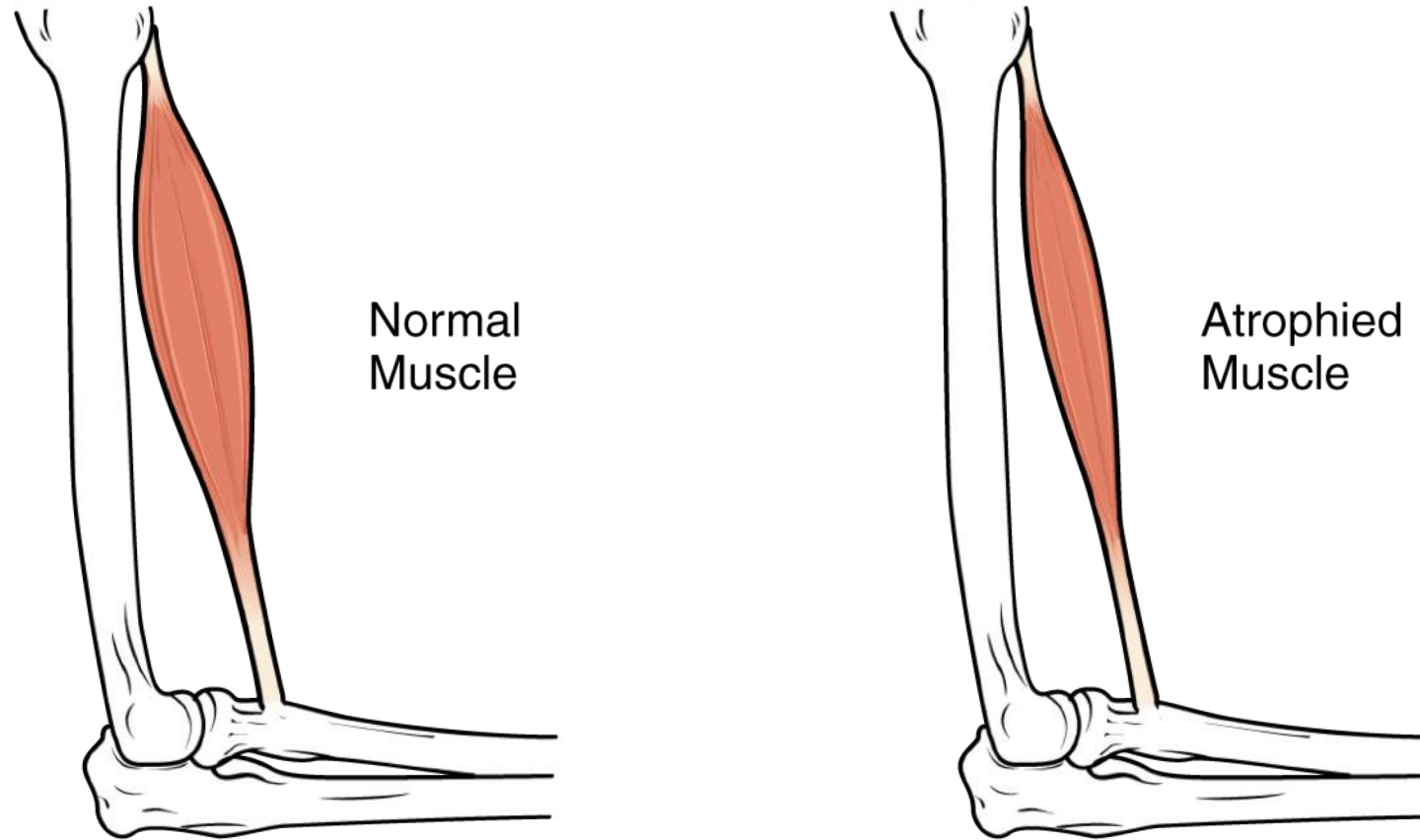
Muscular Dystrophy & *CNBP*



WISCONSIN
UNIVERSITY OF WISCONSIN-MADISON

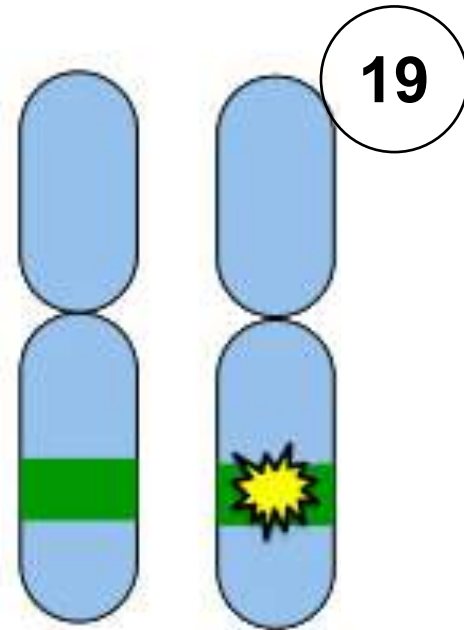
| **Paxton Paine**

Muscular dystrophy (MD) is characterized by muscle loss and weakness

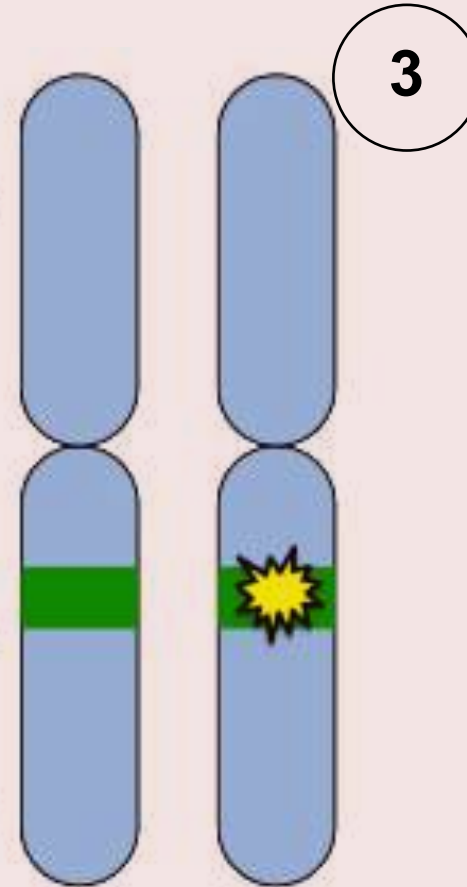


Myotonic dystrophy (DM) will be this talk's focus

Myotonic dystrophy—Most common MD found in adults

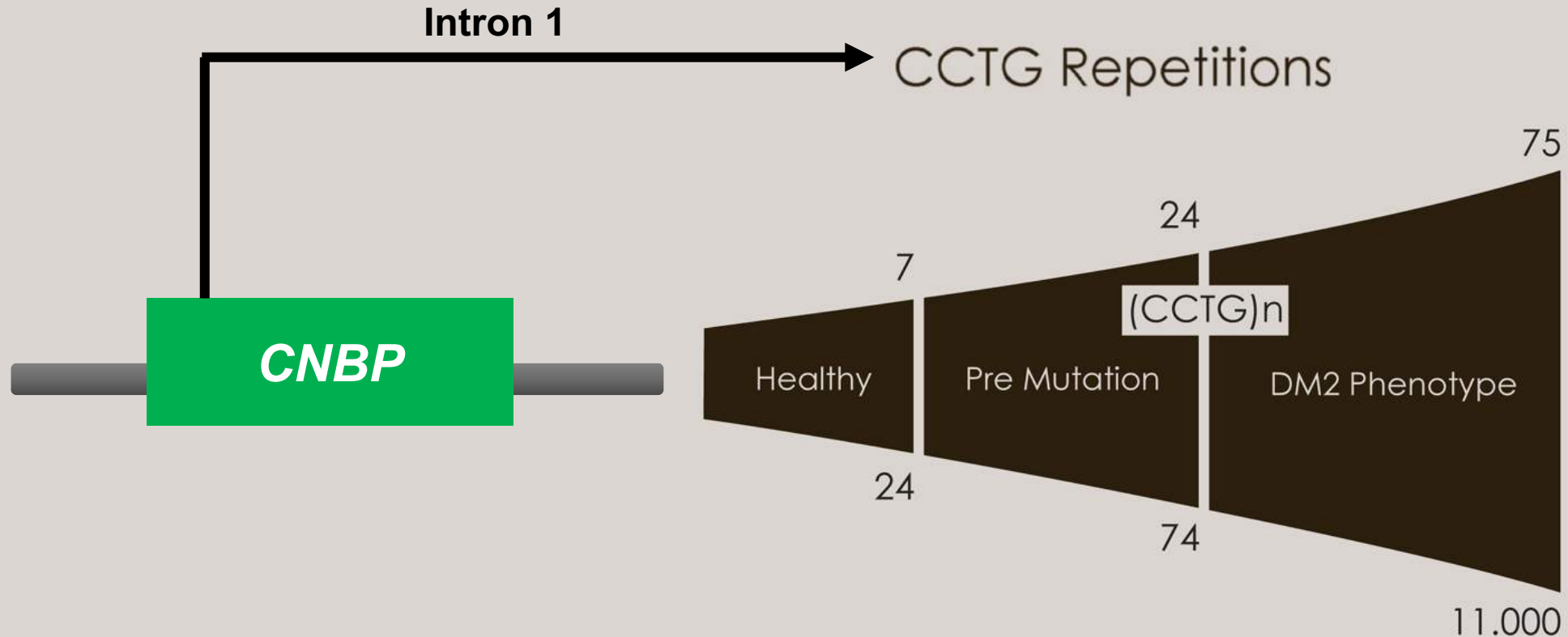


Type 1 (DM1)
DMPK gene mutation



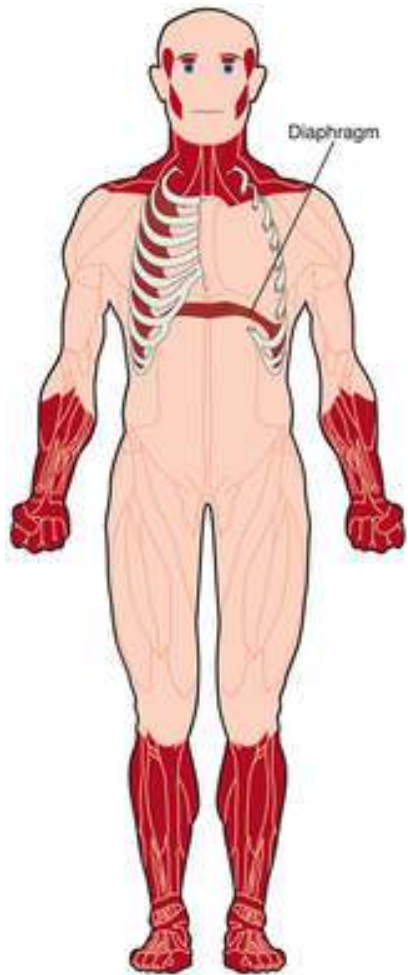
Type 2 (DM2)
CNBP gene mutation

DM2 is caused by a tetranucleotide CCTG repeat in the first *CNBP* intron



RNA-gain-of-function is a prominent mechanism in both type 1 and 2

DM2 has a wide range of symptoms



**Muscle
wasting**

**Muscle
weakness**

Myotonia

**Early onset
cataracts**

**Heart
problems**

**Muscle
pain**

More mild than DM1

Prevalence ~1/8000

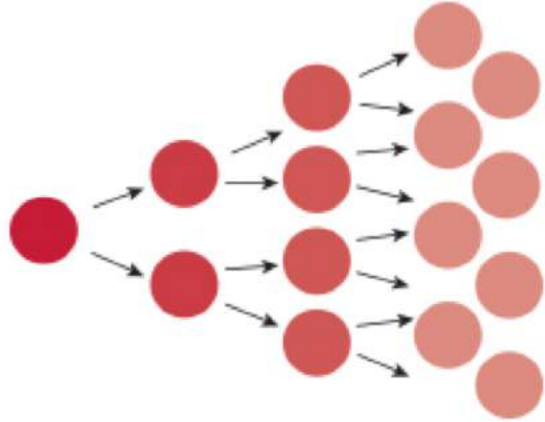
Onset at 20-30 years

What is the **CNBP** protein?

Homo sapiens
Humans

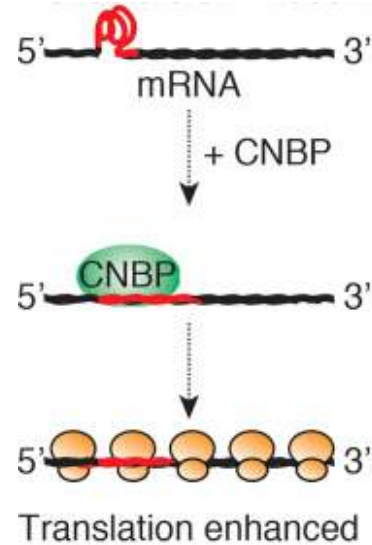


Biological Processes



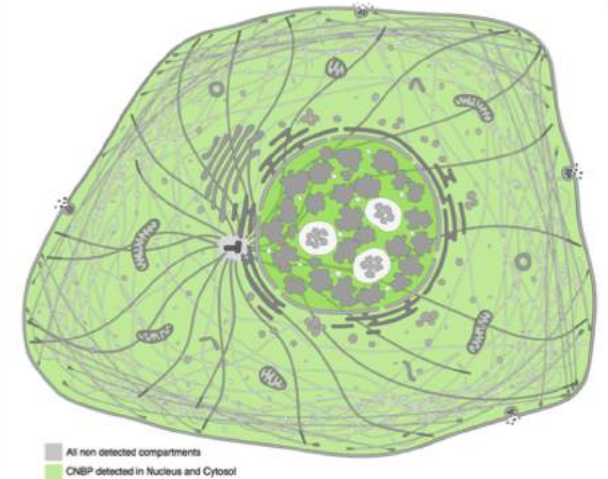
Reg. of transcription/translation
Reg. of cell proliferation
Cholesterol biosynthetic process

Molecular Function



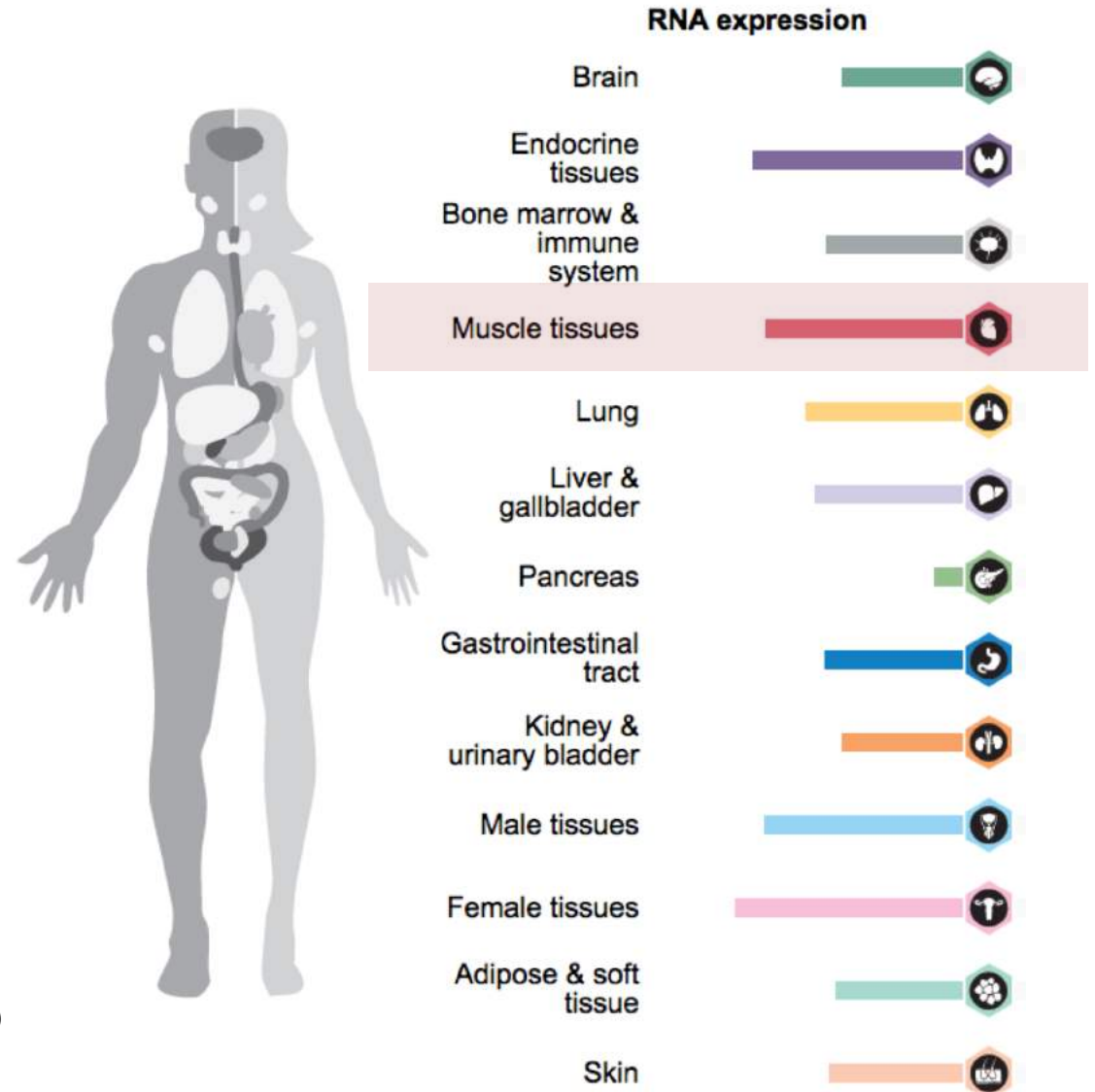
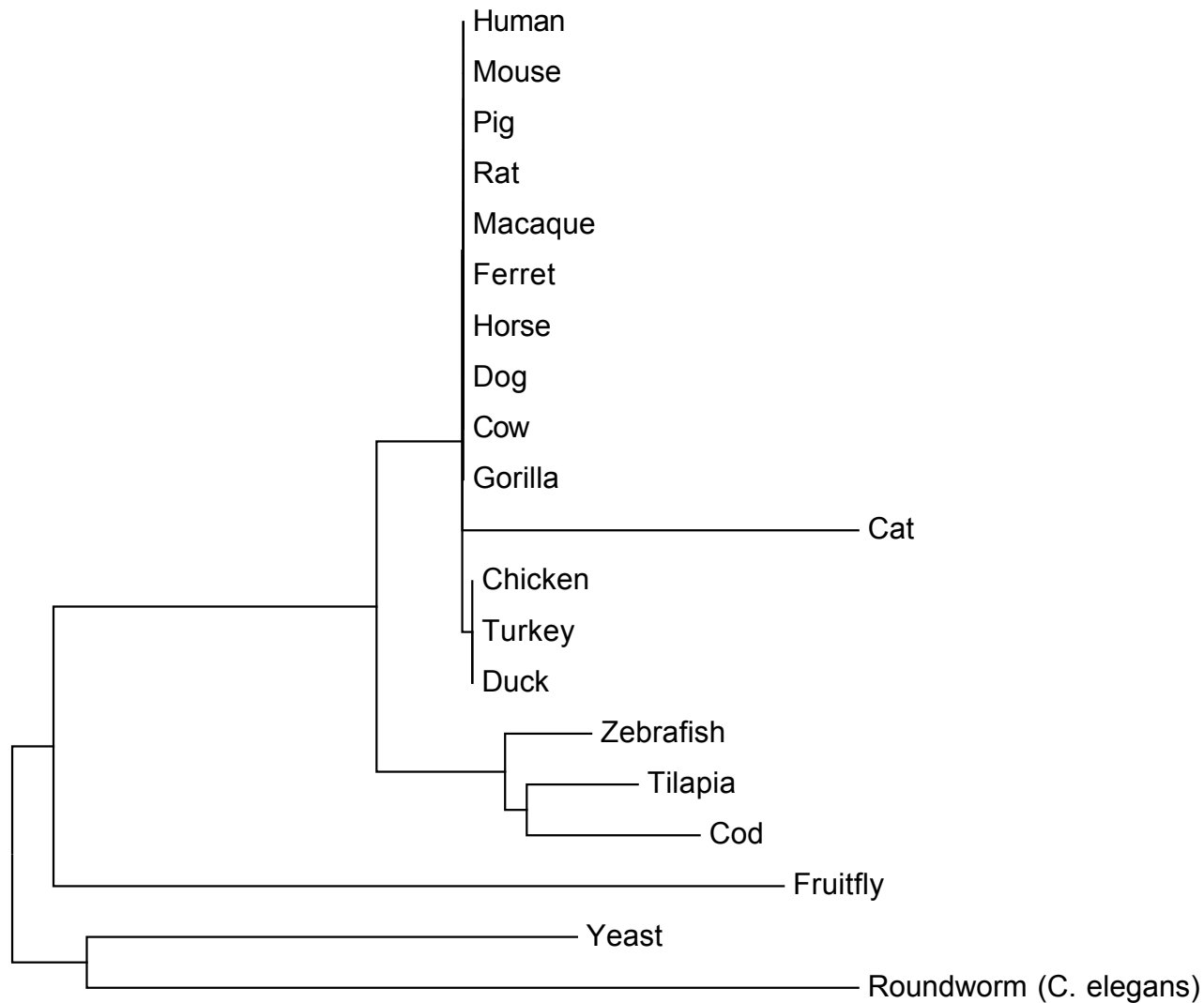
Metal ion binding
DNA & RNA binding

Cellular Components

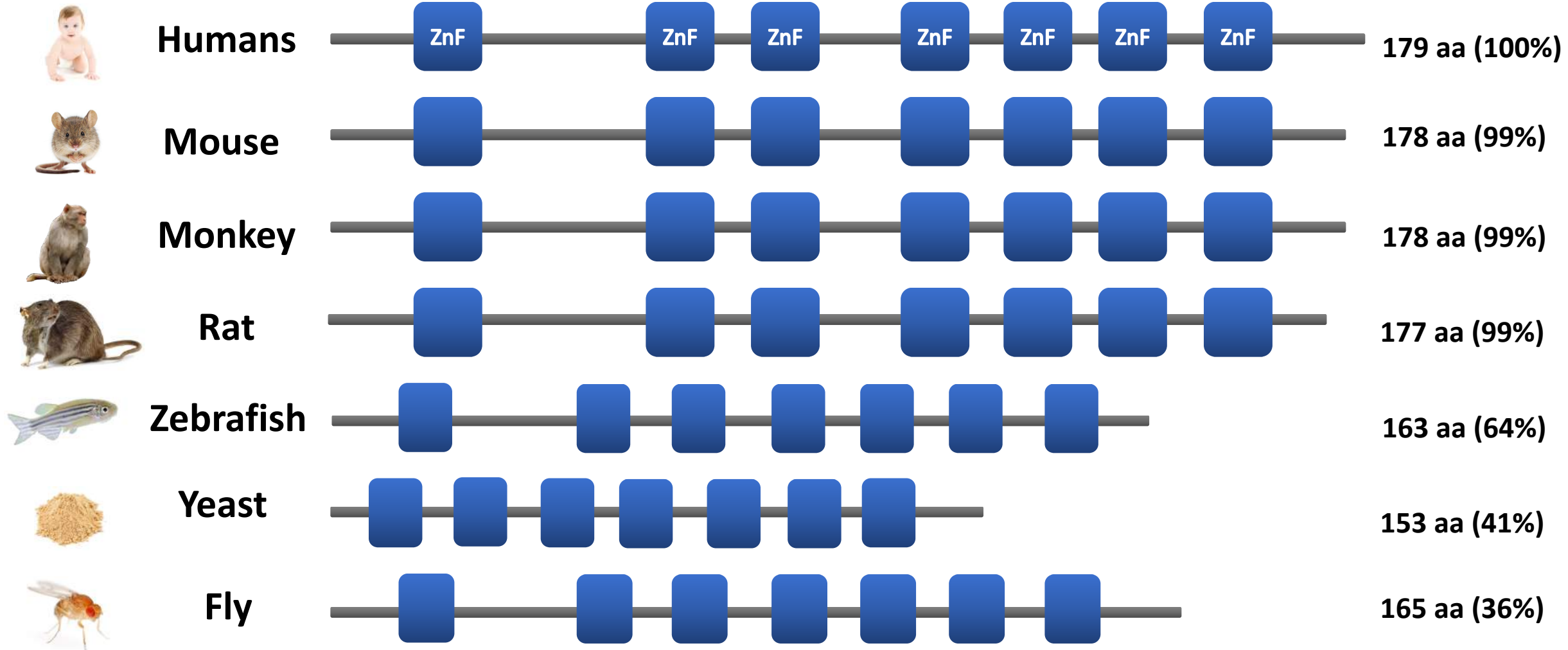


Cytosol
Nucleus
ER

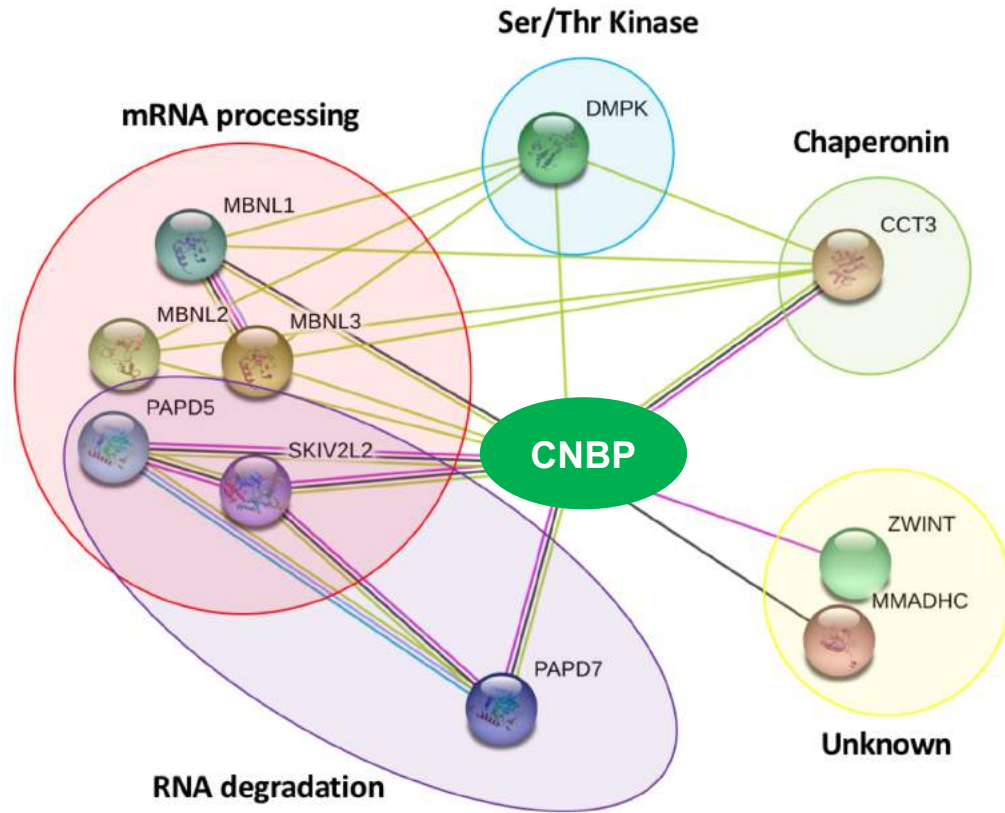
Phylogeny & expression of CNBP



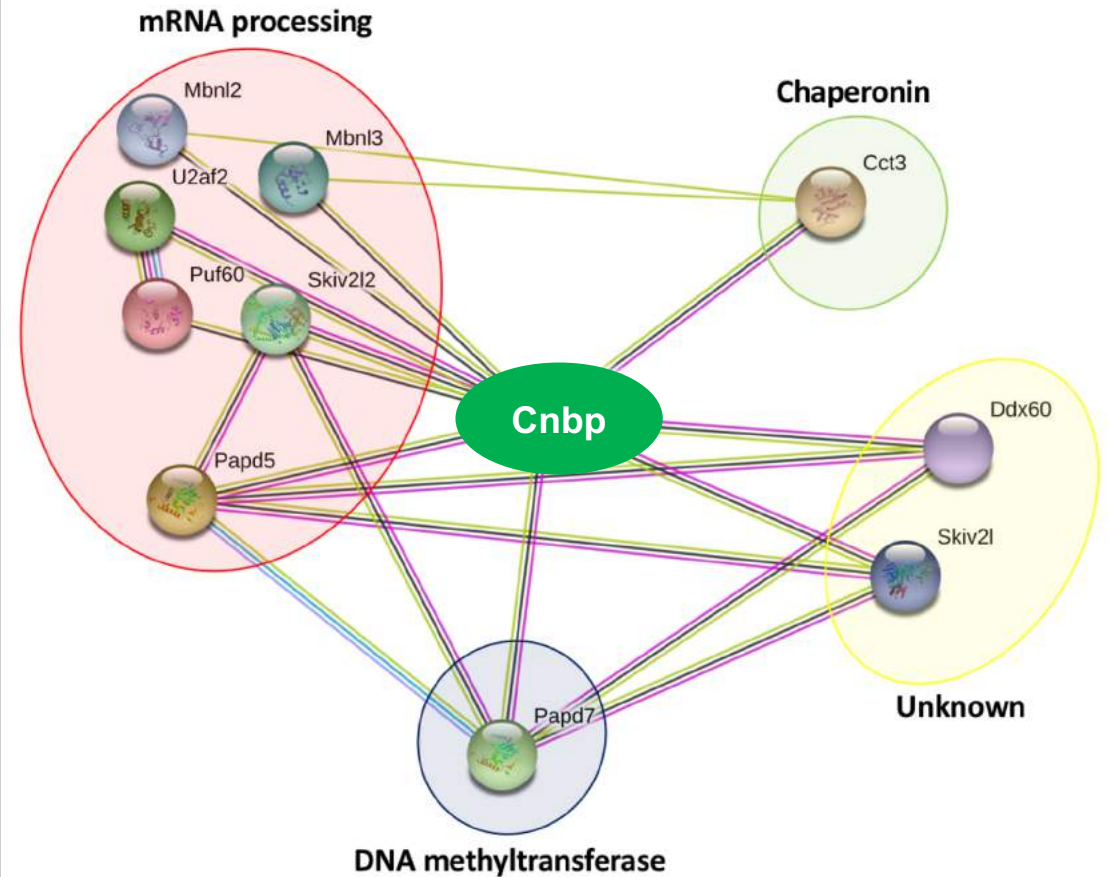
Is **CNBP** conserved across species?



CNBP protein-protein interactions



Human

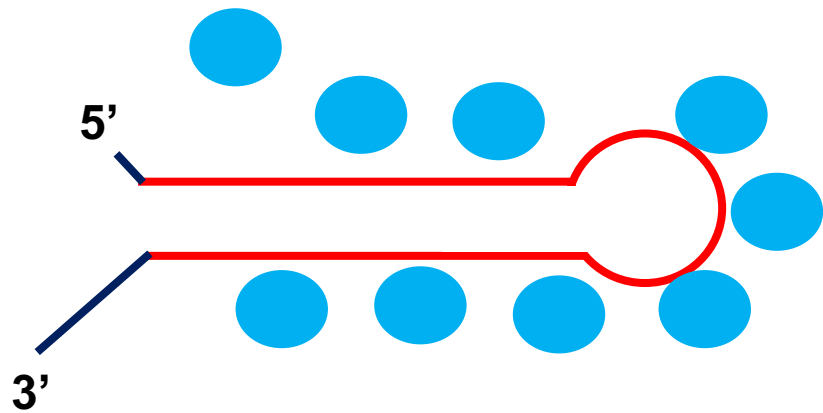


Mouse

What is currently known about DM2 & CNBP?

CNBP pre-mRNA is toxic, sequesters proteins, and not translated

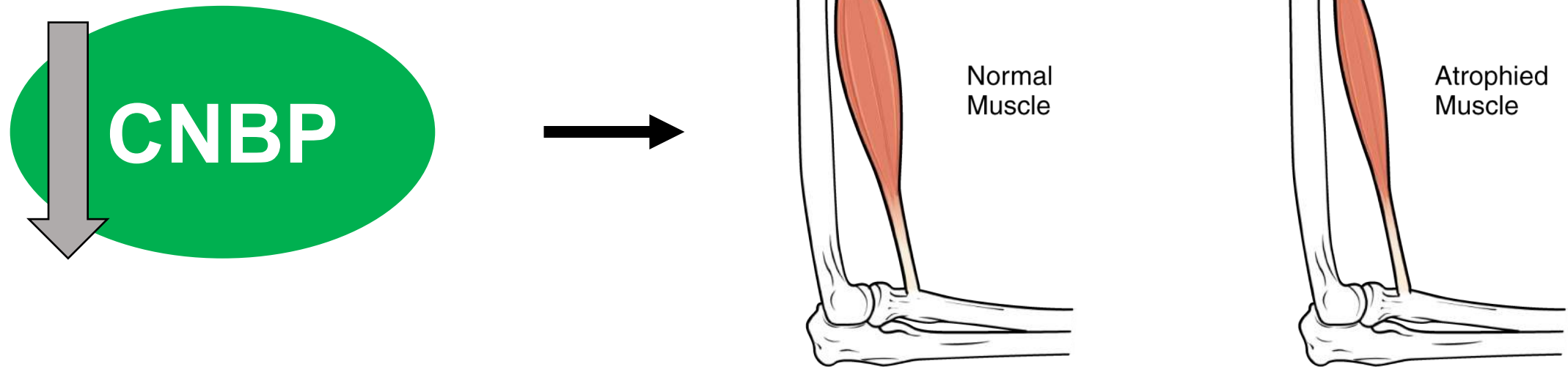
CNBP downregulated in DM2



***Present in both DM1
& DM2**



It is unknown how **decreased CNBP** contributes to **muscle wasting** and **weakness**

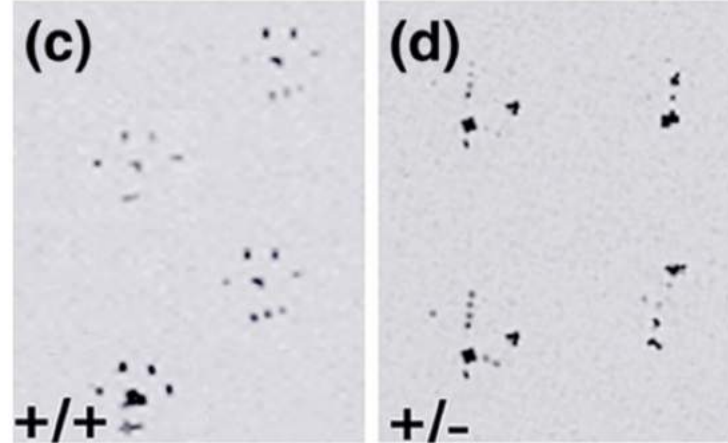


Also unclear why DM2 less severe than DM1

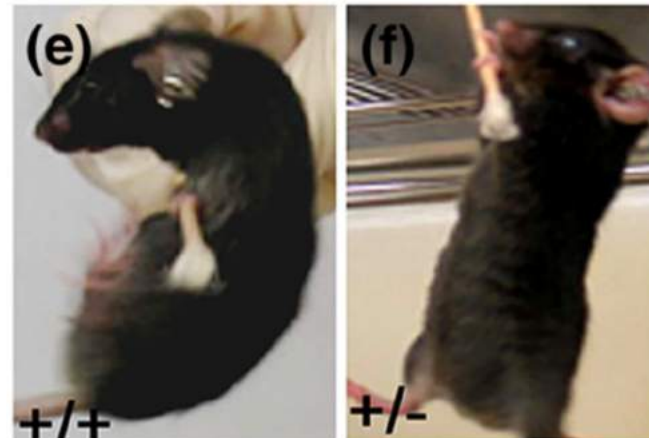
Model organism phenotypes



Mus musculus



Defective walking & unclear footprints

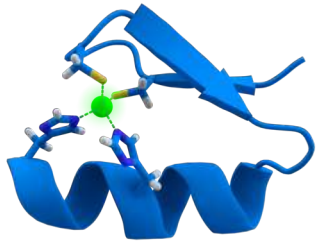


Extremely weak & proximal/distal muscle wasting

Haploinsufficient mice exhibit DM2 phenotypes

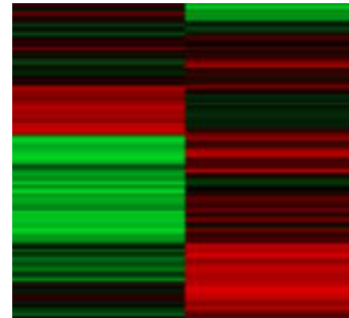
Goal: To study how low levels of **CNBP** contributes to **muscle wasting** and **weakness**

Aim 1



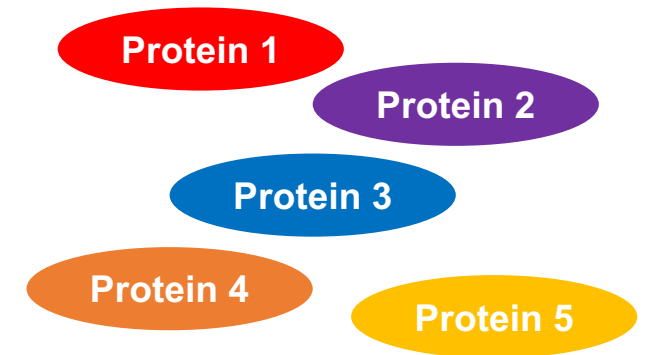
Identify conserved sites of CNBP important to proper muscle function

Aim 2



Identify genes important to proper muscle function

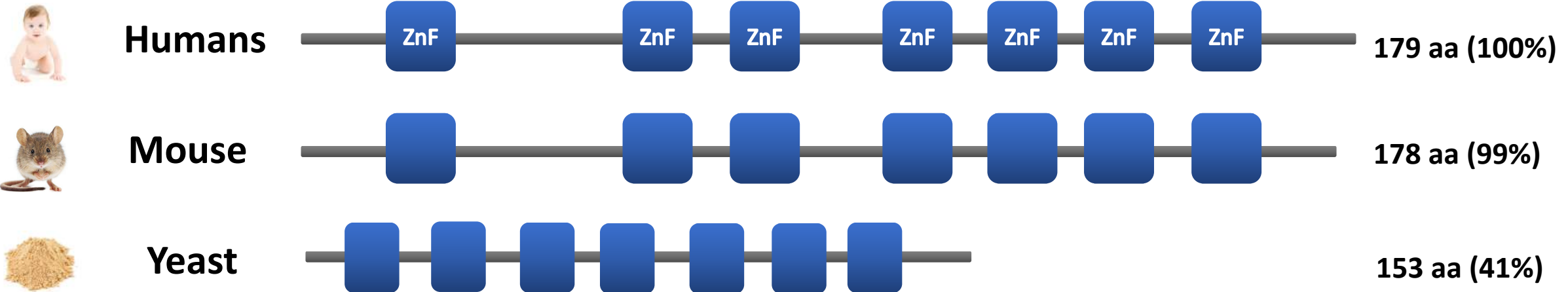
Aim 3



Quantify dysregulation of proteins involved in skeletal muscle function

Hypothesis: **CNBP loss** contributes to weakness and wasting by affecting pathways not associated with the RNA-gain-of-function mechanism

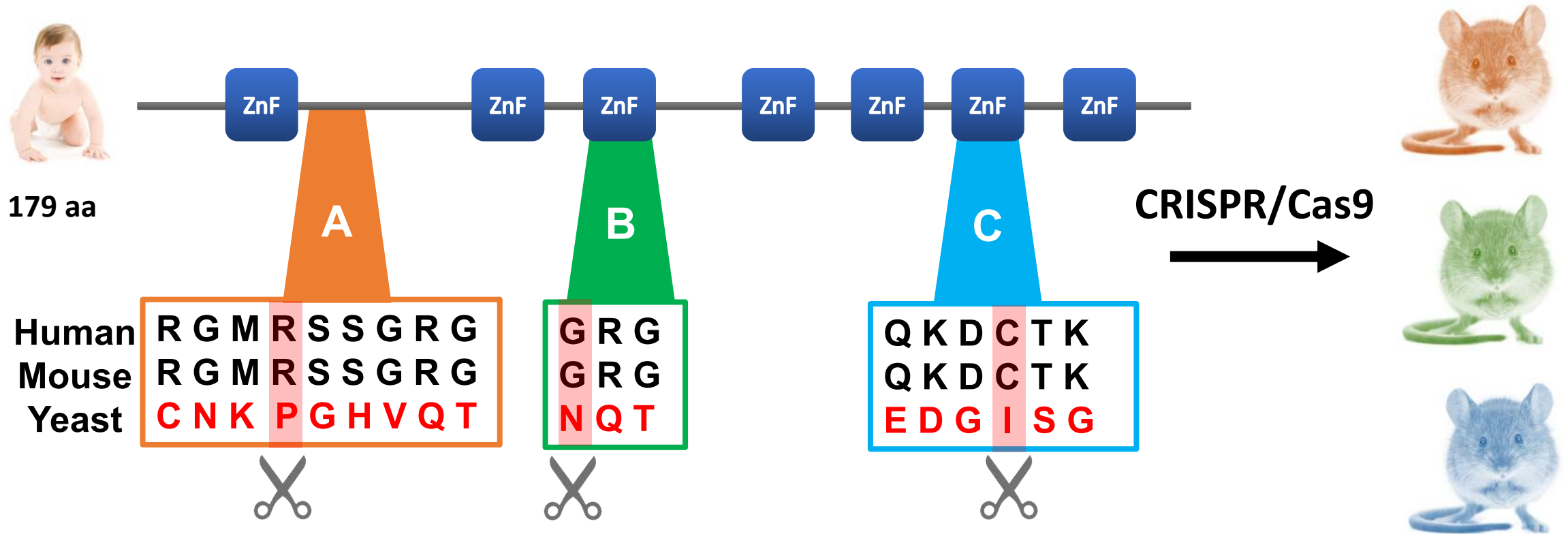
Aim 1: Identify conserved sites of CNBP critical to muscle function



Species/Abbrv	Group Name								*																											*																				
1. Human		R	G	R	G	M	R	S	R	G	R	G	G	F	T	S	D	R	G	F	Q	F	V	S	S	S	L	P	D	I	C	Y	R	C	G	E	S	G	H	L	A	K	D	C	D	L	Q	E	D	V	E	A	C	Y		
2. Mouse		R	G	R	G	M	R	S	R	G	R	G	G	F	T	S	D	R	G	F	Q	F	V	S	S	S	L	P	D	I	C	Y	R	C	G	E	S	G	H	L	A	K	D	C	D	L	Q	E	D	E	-	A	C	Y		
3. Fruitfly		G	G	P	G	G	V	G	G	G	G	G	G	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	G	G	G	G	M	R	G	N	D	G	G	G	M	R	R	N	R	E	-	-	-	-	-	-	-	K	C	Y
4. Tilapia		R	G	R	G	K	-	G	R	G	R	G	K	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	D	L	F	C	Y	R	C	G	E	L	G	H	V	A	R	D	C	E	R	T	E	D	-	-	A	C	Y
5. Cow		R	G	R	G	M	R	S	R	G	R	G	-	-	-	-	-	-	-	F	Q	F	V	S	S	S	L	P	D	I	C	Y	R	C	G	E	S	G	H	L	A	K	D	C	D	L	Q	E	D	-	-	A	C	Y		
6. Yeast		L	C	Y	N	C	N	K	P	G	H	V	Q	-	-	-	-	T	D	C	T	M	P	R	T	V	E	F	K	Q	C	Y	N	C	G	E	T	G	-	-	H	V	R	S	E	C	T	V	Q	-	-	R	C	F		
7. Dog		R	G	R	G	M	R	S	R	G	R	G	G	F	T	S	D	R	G	F	Q	F	V	S	S	S	L	P	D	I	C	Y	R	C	G	E	S	G	H	L	A	K	D	C	D	L	Q	E	D	V	E	A	C	Y		
8. Horse		R	G	R	G	M	R	S	R	G	R	G	-	-	-	-	-	-	F	Q	F	V	S	S	S	L	P	D	I	C	Y	R	C	G	E	S	G	H	L	A	K	D	C	D	L	Q	E	D	-	-	A	C	Y			

1. Identify protein domains using SMART and align FASTA sequences with ClustalW

Aim 1: Identify conserved sites of CNBP critical to muscle function

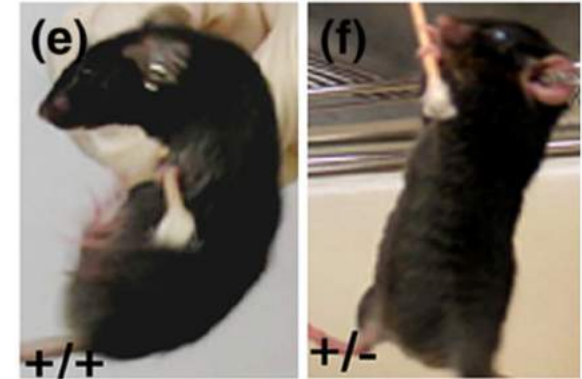
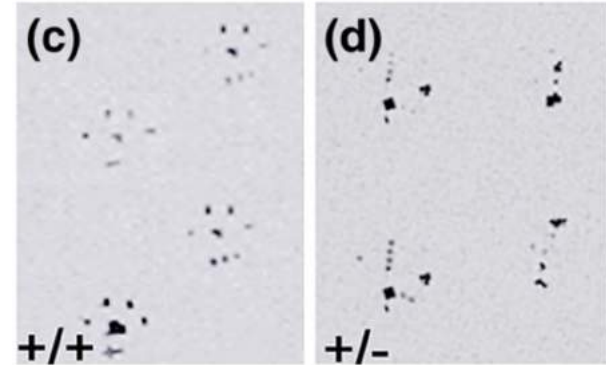


2. Create mutants in conserved regions with CRISPR/Cas9

Aim 1: Identify conserved sites of CNBP critical to muscle function



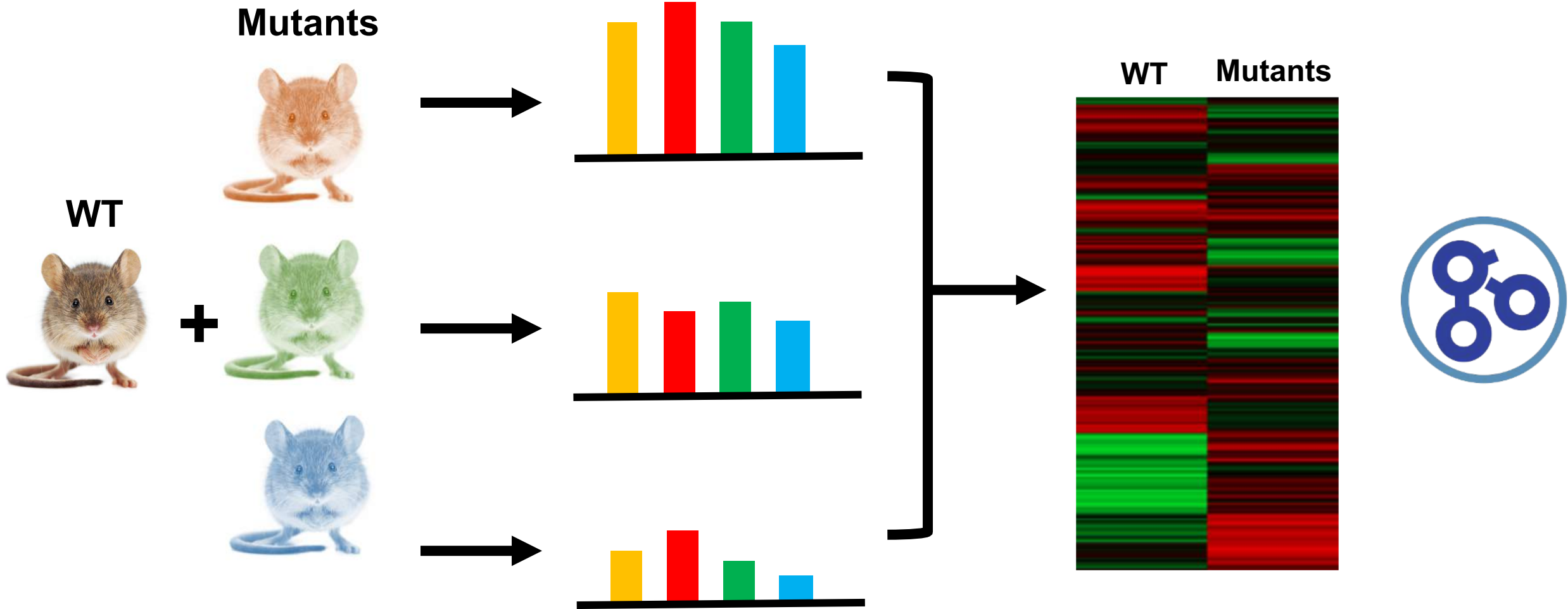
Phenotype
screen



Hypothesis: Specific sites conserved only within muscular organisms will be critical in regulating muscle function

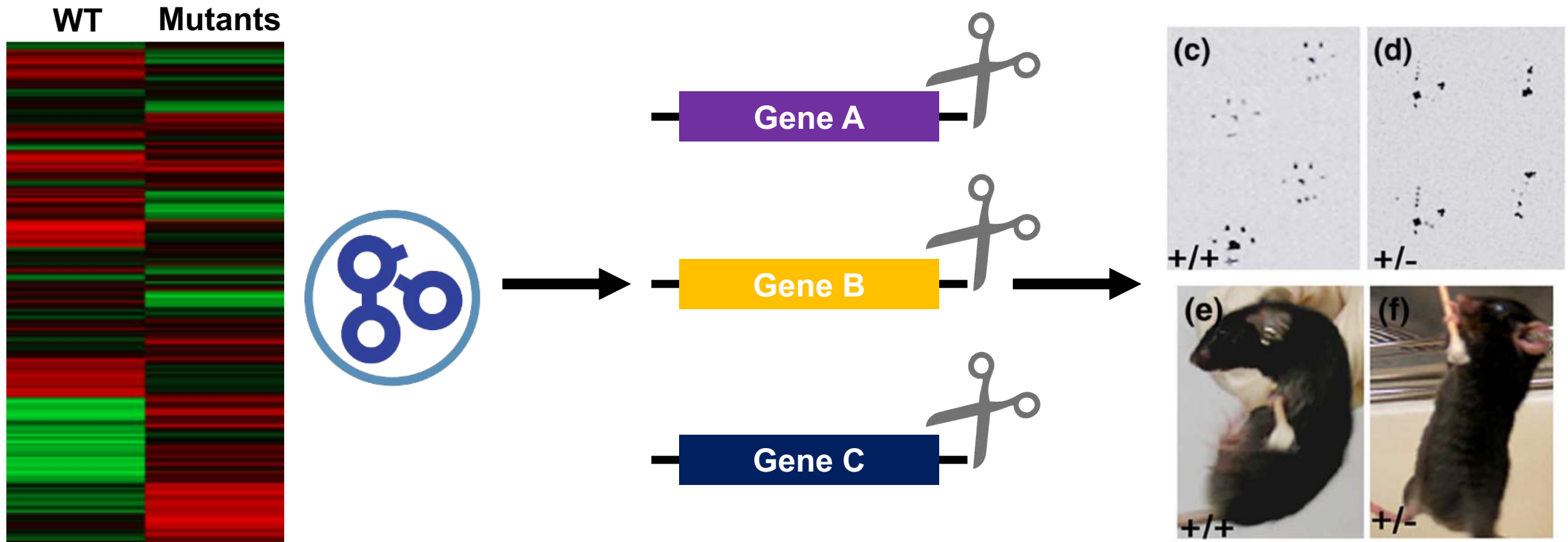
3. Screen for mutant phenotypes

Aim 2: Identify genes that are important in regulating muscle function



1. Perform RNA-seq in mouse muscle tissue and sort with GO

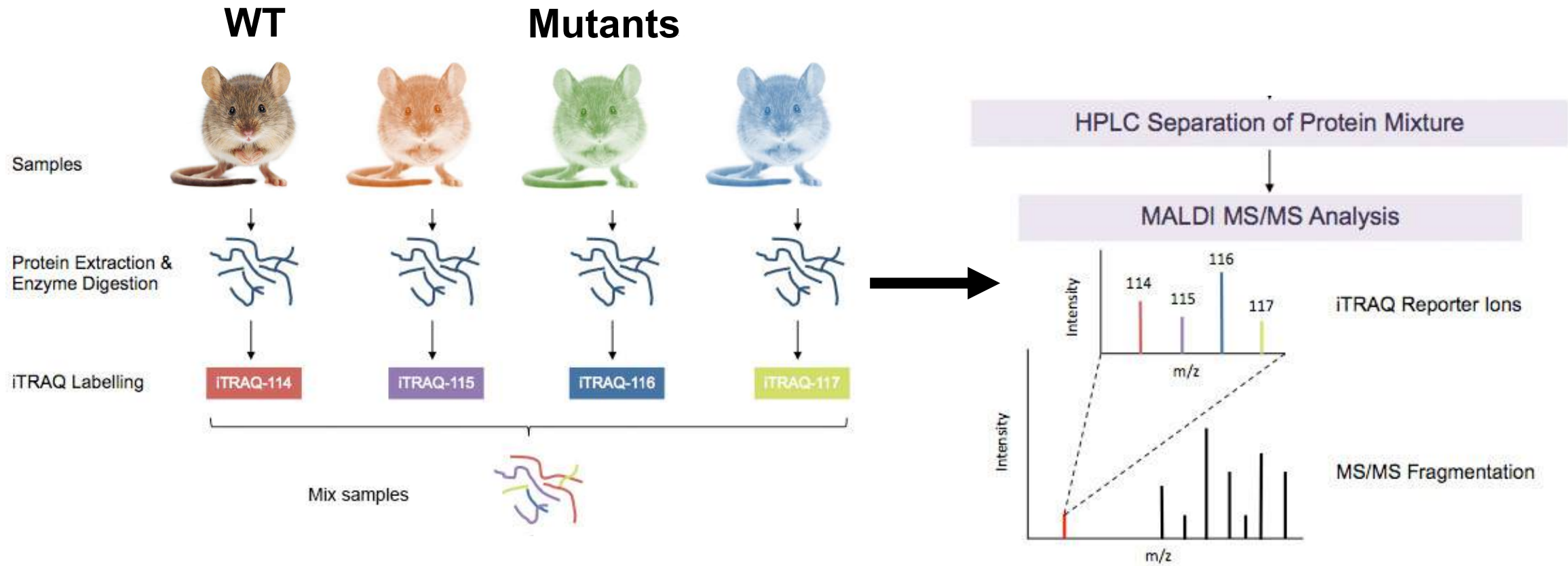
Aim 2: Identify genes that are important in regulating muscle function



Hypothesis: CNBP mutants will have differentially expressed genes due to changes in CNBP-DNA binding and these genes are critical to regulating muscle function

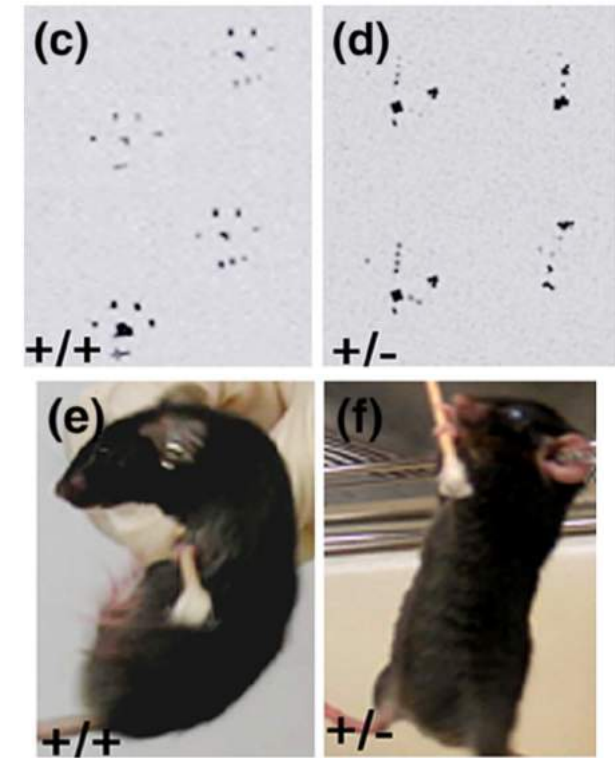
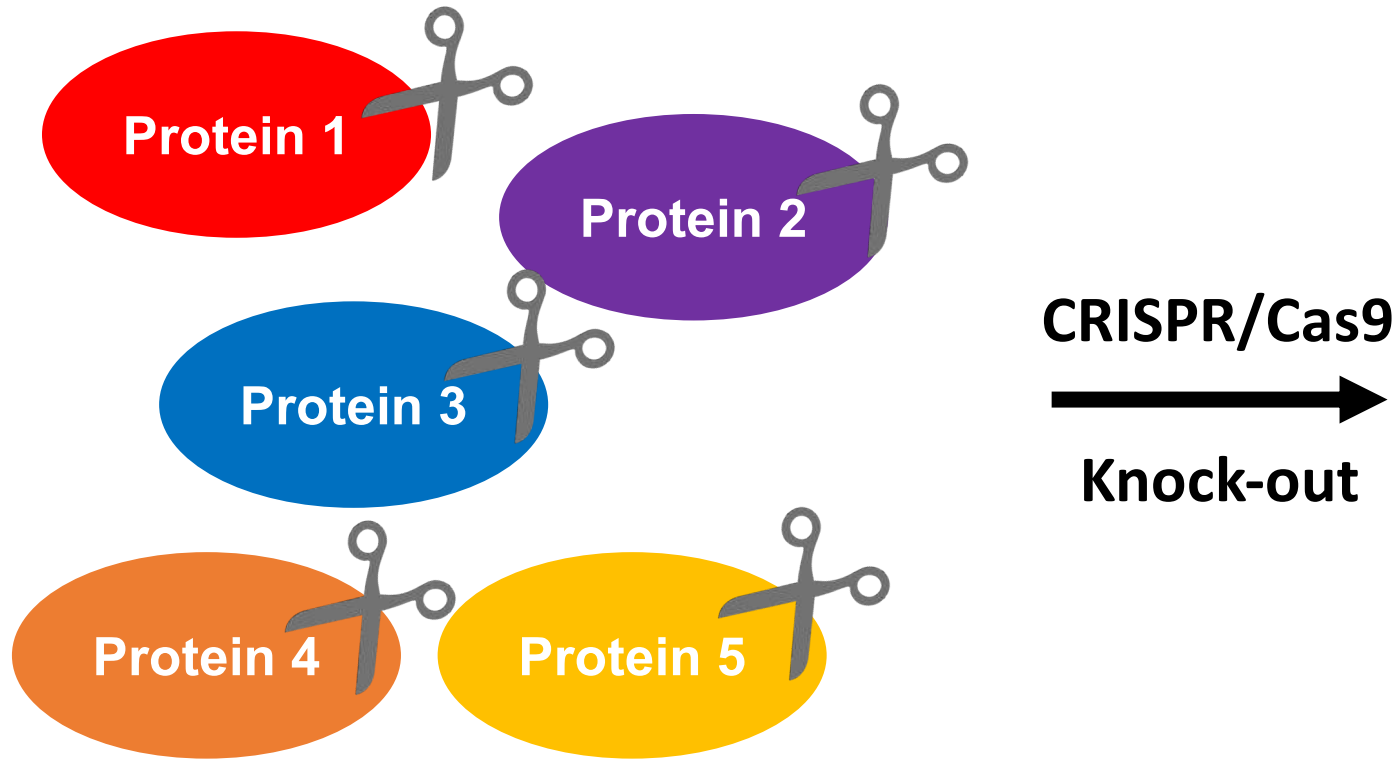
2. Knock-out differentially expressed transcripts with CRISPR & screen

Aim 3: Quantify dysregulation of proteins involved in skeletal muscle function



1. Isolate proteins from skeletal muscle of WT and CNBP mutants

Aim 3: Quantify dysregulation of proteins involved in skeletal muscle function



Hypothesis: CNBP mutants will have dysregulated proteins levels and these proteins will be critical to skeletal muscle function

2. Quantify proteins, KO dysregulated proteins and screen for mutant phenotypes

Summary of specific aims

Aim 1

Identifies regions of CNBP critical to regulating muscle function

Aim 2

Identifies mRNAs dysregulated by loss of CNBP

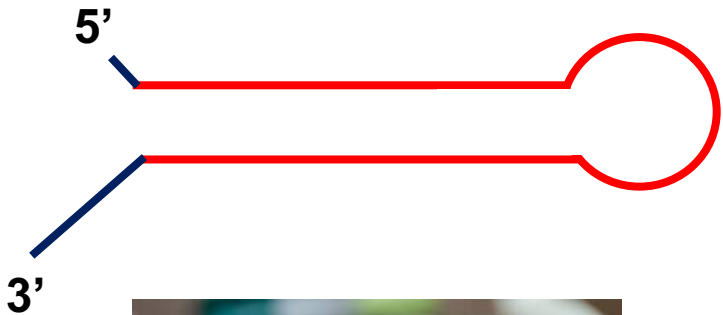
Aim 3

Identifies proteins and protein classes dysregulated by CNBP

Future directions



Characterize CNBP interactions in other tissues, such as eye and endocrine tissues

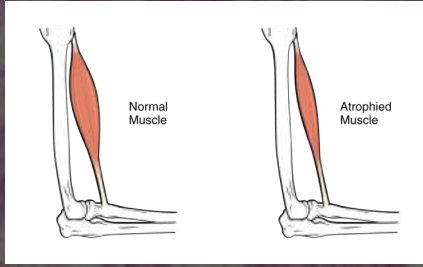


Chemical screens to remove *CNBP* pre-mRNA

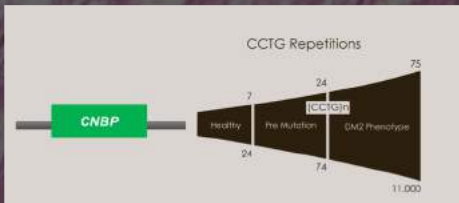


Continue developing drugs for management of symptoms, such as pain and myotonia

Conclusions



DM2 is a disease characterized by many symptoms



Nucleotide repeats in the first intron of *CNBP* causes DM2



The effects of lower CNBP levels are unclear



Learning more about each disease mechanism will aid in treatment of disease



Questions?

References

<https://www.ncbi.nlm.nih.gov/pubmed/28329689>

<https://www.ncbi.nlm.nih.gov/pubmed/28078562>

<https://www.ncbi.nlm.nih.gov/pubmed/17335846>

<https://ghr.nlm.nih.gov/gene/CNBP>

<https://ghr.nlm.nih.gov/condition/myotonic-dystrophy>