

UPDATE ON NAF RESEARCH

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University of Florida



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GOALS AND IMPACT OF NAF FUNDING

- Birds-Eye View of Funding Portfolio
 - Making scientific connections between diseases
 - Encouraging young investigators to pursue ataxia research
- Some Examples
 - Genetics 101-2015
 - Four Pioneer Awards
 - Therapeutic strategies to fight ataxia
 - Connections between scientists, diseases, NIH and Industry

NAF'S RESEARCH PROGRAMS

- **Research Seed-Money Grants** – Seed monies in early or pilot phases of studies and ongoing investigations that demonstrate need to attract future funding from other sources.
- **Post-Doc Fellowship Awards** – Intended for researcher to spend a 3rd year in a post-doc position to increase their chance of establishing an independent ataxia research program



NAF'S RESEARCH PROGRAMS

- **Young Investigator and Young Investigator SCA Awards** - Awarded to encourage young investigators to pursue a career in the field of ataxia
- **Pioneer SCA Translational Awards** - This grant is intended for research investigations that will facilitate the development of treatments for the spinocerebellar ataxias (SCAs).

RESEARCH APPLICATIONS RECEIVED

- 33 Seed-money grant applications were received
- 22 Post-Doc Fellowship Applications received
- 15 Young Investigator (non-SCA) applications received
- 14 Pioneer SCA Translational Research Applications received
- 18 Young Investigator SCA Applications received

TOTAL of 102 Applications were submitted

13 COUNTRIES REPRESENTED

- United States
- United Kingdom
- Brazil
- The Netherlands
- Germany
- Portugal
- Australia
- Canada
- Spain
- France
- Italy
- Cyprus
- Mexico

TYPES OF ATAXIA

- SCAs
- Friedreich Ataxia
- Gordon Holmes Syndrome
- Joubert Syndrome
- Mitochondrial
- FXTAS
- A-T
- Episodic Ataxia
- ARSACS
- Spastic ataxia
- AOA
- SCAN
- Sporadic

REVIEW PROCESS

Reviewers are assigned by Dr. Harry Orr and Dr. Laura Ranum

All applications are peer-reviewed by 2 reviewers

Pioneer applications peer-reviewed by 3 reviewers

More than 60 reviewers worldwide

Conference calls are attended by a sub-group of NAF Medical Research Advisory Board members to make funding recommendations

NAF Board of Directors make the final funding decisions

NAF FUNDED RESEARCH

27 Research Projects were Funded totaling more than
\$1.1 million of research

9 Seed Money Research Grants
4 Young Investigator Awards
7 Post-doc Fellowship Awards
4 Young Investigator-SCA Awards
3 Pioneer Translational SCA Awards

26% of submitted grants were funded
Success rate is substantially higher than the NIH

RESEARCH SEED GRANTS 2016

Speech Rehabilitation
SCA3
Gene discovery
SCA36
Brain Imaging
Inflammation in Setx
Ataxia Database

Vogel, Adam, PhD
Centre for Neuroscience of Speech Parkville, Victoria, Australia
[Intensive home based speech rehabilitation for adults with degenerative ataxia](#)

Simões, Ana, PharmD, PhD
University of Coimbra, Portugal
[Calpain-mediated proteolysis in Machado-Joseph disease](#)

Karnebeek, Clara, MD, PhD
University of British Columbia, Vancouver, BC, Canada
[Whole Exome Sequencing in the Diagnosis and Management of Atypical Childhood Hereditary Ataxia Conditions](#)

He, Fang, PhD
Texas A&M University, Kingsville, TX
[Development of a Drosophila model for Spinocerebellar Ataxia type 36 \(SCA36\)](#)

Oz, Gulin, PhD
University of Minnesota, Minneapolis, MN
[Launching the US-Europe Neuroimaging Partnership in SCA](#)

Burmeister, Margit, PhD
University of Michigan, Ann Arbor, MI
[Role of VPS13D in Ataxia](#)

Lavin, Martin, PhD
University of Queensland Centre for Clinical Research (UQCCR) Australia
[Assessing the role of senataxin in cellular inflammation, gene regulation, and innate immunity in Setx-/- mice and a human neuronal model](#)

Ribeiro, Sandra, PhD
Instituto de Biologia Molecular e Celular Porto, Portugal
[New therapeutic approaches for Machado-Joseph Disease: Chaperoning protein self-assembly](#)

Perlman, Susan, M.D.
University of California Los Angeles, Los Angeles, CA
[Web-based National Ataxia Database](#)

NAF GRANT RECIPIENTS YOUNG INVESTIGATORS IN 2016

INVESTING IN THE NEXT GENERATION OF ATAXIA RESEARCHERS

Gene Regulation
Removal of toxic proteins in SCA3
Friedreich's – mitochondrial stress
Cell-Cell communication SCA27
Astrocyte signaling in SCA1
Turning up Frataxin with Crispr/Cas
ASO SCA3
Chemical pathways in Friedreich's
Pathways for polyQ regulation SCA3
Protein networks in ataxia
PUMILIO1 – a new ataxia gene

Yao, Bing, PhD

Emory University, Atlanta, GA

[Epigenetic Modulation Mediated by RNA-Binding Proteins in Neurodegeneration](#)

Schmidt, Jana, PhD

University of Tuebingen, Germany

[Alleviation of proteasomal inhibition as a therapeutic approach for SCA3](#)

Butler, Jill Sergesketter, PhD

University of Alabama at Birmingham, AL

[Reduced expression of mitochondrial aldehyde dehydrogenases contributes to metabolic stress in Friedreich's ataxia](#)

Ben-Johny, Manu, PhD

Johns Hopkins University, Baltimore, MD

[Aberrant Regulation of Voltage-gated Na channels in the Pathophysiology of Spinocerebellar Ataxia 27](#)

Cvetanovic, Marija, PhD

University of Minnesota, Minneapolis, MN

[Role of astrocyte calcium signaling in the pathogenesis of SCA1](#)

Nageshwaran, Sathiji, MD

Harvard University, Boston, MA

[Transcriptional activation using CRISPR/Cas mutant proteins as a novel therapy for Frataxin gene silencing](#)

Khurana, Vikram, MD, PhD

Brigham and Women's Hospital and Harvard Stem Cell Institute, Boston, MA

[Systematic edgotyping of ataxin proteins in cellular systems from yeast to patient neurons.](#)

Gennarino, Vincenzo, PhD

Baylor College of Medicine, Houston, TX

[PUMILIO1 deficiency: understanding a new ataxia gene and its role in cerebellar dysfunction in mice and humans](#)

NAF GRANT RECIPIENTS POSTDOCTORAL FELLOWSHIPS 2016

INVESTING IN THE NEXT GENERATION OF ATAXIA RESEARCHERS

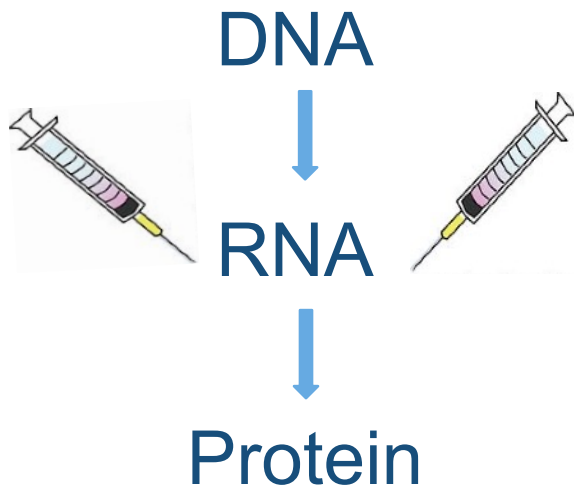
Brain Structure
Small Molecule Drugs
Motor Neuron Degeneration
RNA Degradation of CAG repeats
Cell-cell communication and protection
Protein dysregulation

- Anderson, Collin J., PhD
University of Utah, Salt Lake City, UT
[Development and mechanistic study of deep brain stimulation of dentate nucleus for the treatment of degenerative ataxia.](#)
- Orengo, James, MD, PhD
Baylor College of Medicine, Houston, TX
[Unraveling the mechanisms of motor neuron degeneration in Spinocerebellar Ataxia, type1](#)
- Chen, Jonathan, PhD
Scripps Florida, Jupiter, FL
[Rapid structure-based lead optimization of a small molecule drug that target r\(CAG\)exp](#)
- Bott, Laura C., PhD
Northwestern University, Evanston, IL
[Transcellular regulation of the proteostasis network in Spinocerebellar ataxia type 3](#)
- Chopra, Ravi, PhD
University of Michigan, Ann Arbor, MI
[Identifying Dendro-Protective Ion Channels in Cerebellar Ataxia](#)
- Seminara, Stephanie, MD
Massachusetts General Hospital, Boston, MA
[Ataxia with hypogonadotropic hypogonadism due to ubiquitin ligase dysregulation](#)

NAF GRANT RECIPIENTS IN 2016

PIONEER SCA TRANSLATIONAL RESEARCH AWARDS

Moving towards therapies

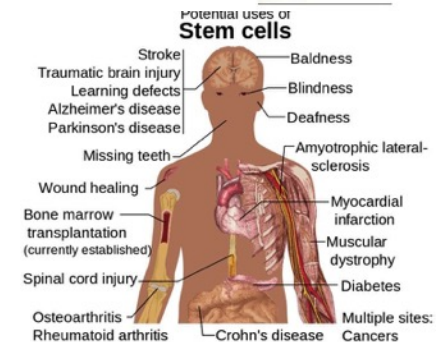


•Orr, Harry, PhD
University of Minnesota, Minneapolis, MN
[Towards an ASO Therapy for Spinocerebellar Ataxia Type 1](#)

•Roon-Mom, Willeke M.C. van, PhD
Leiden University Medical Center, The Netherlands
[Advancing the therapeutic potential of exon skipping](#)
[for Spinocerebellar ataxia type 3](#)

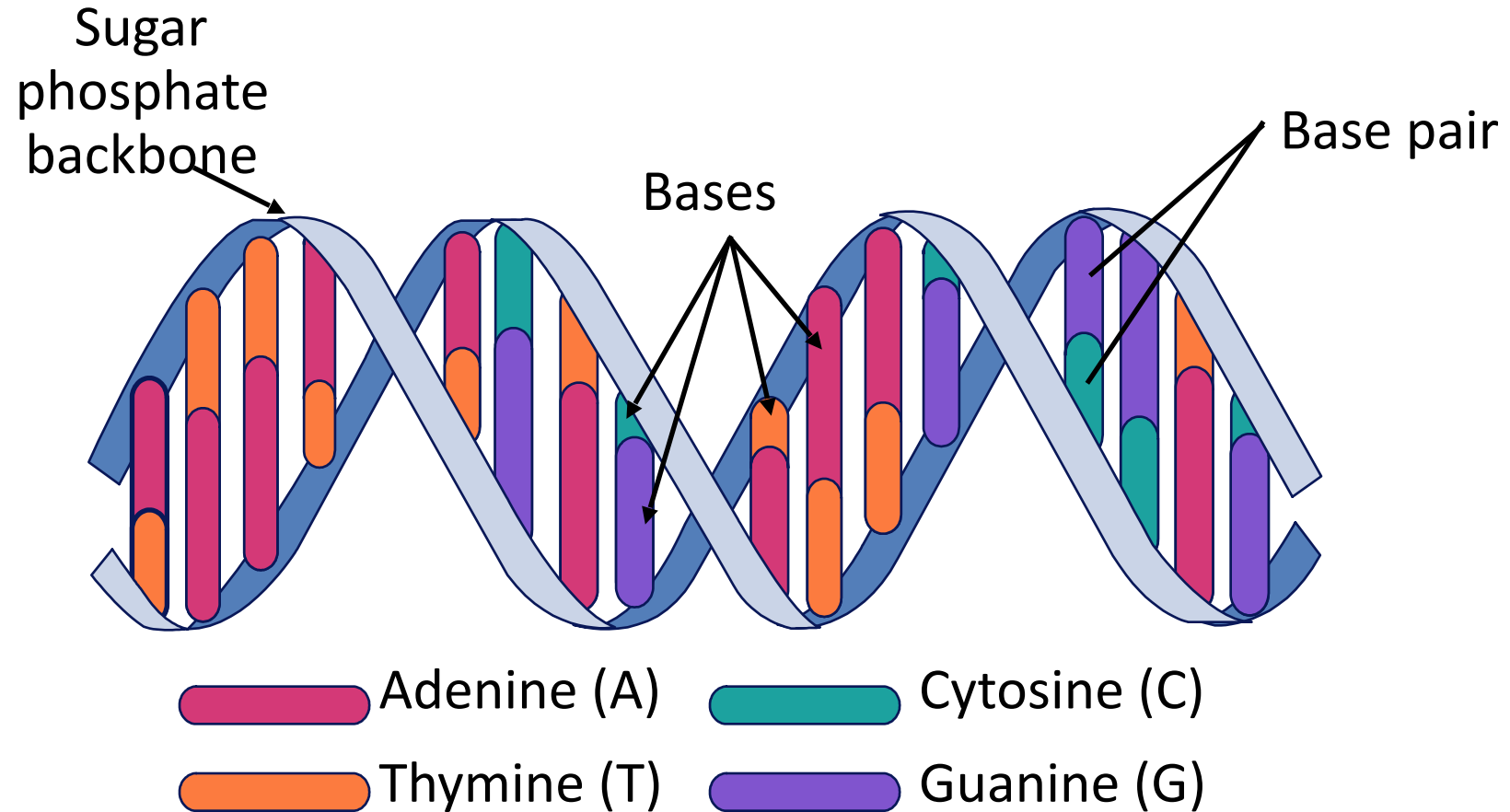
BASIC RESEARCH STEM CELLS

•Maciel, Patrícia, PhD
University of Minho, Braga, Portugal
[Testing the therapeutic potential of Mesenchymal Stem Cells and their secretome in an animal model of spinocerebellar ataxia type 3](#)



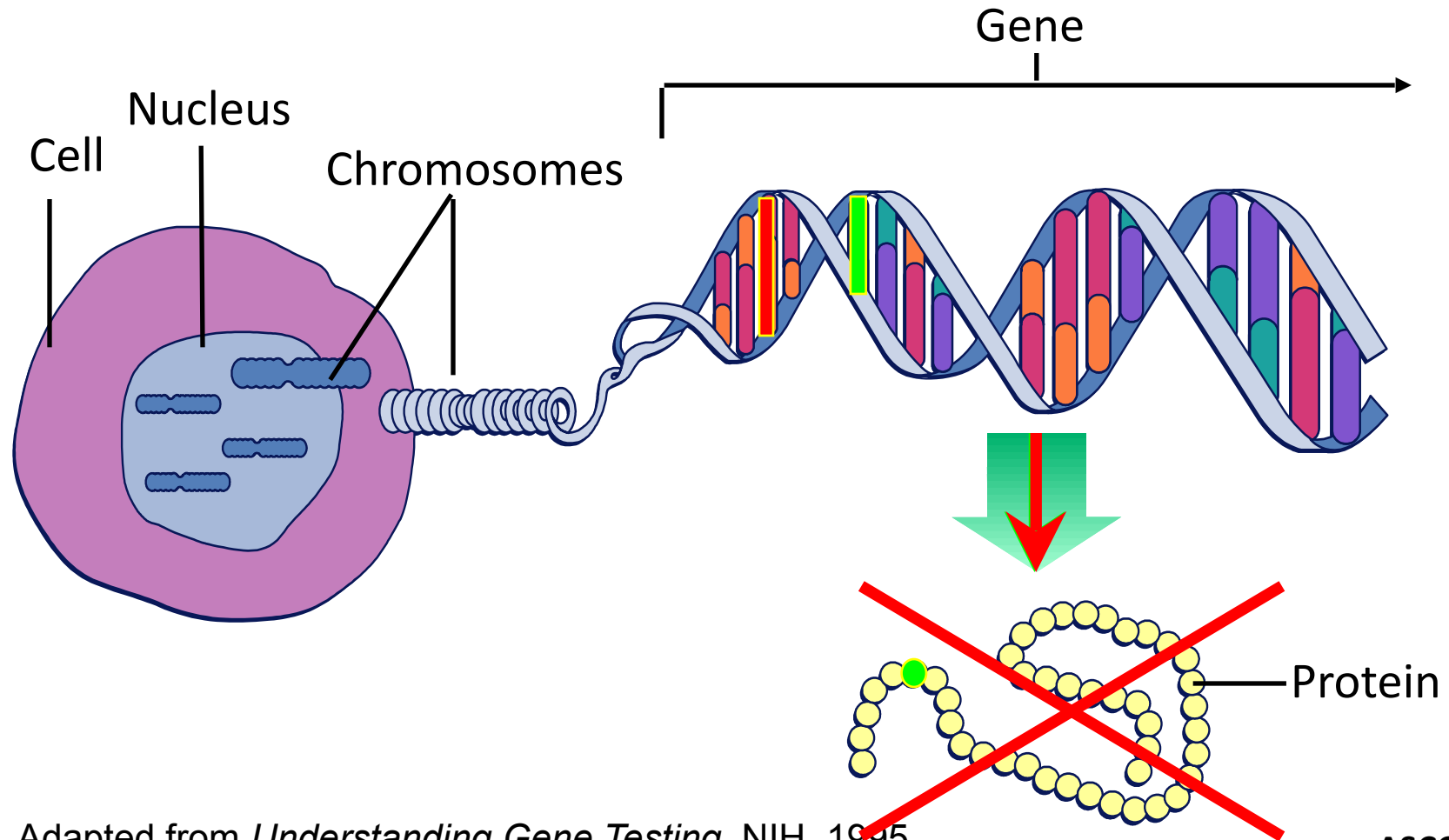
GENE DISCOVERY TO GENE BASED THERAPIES

The DNA Double Helix



Overview

Chromosomes, DNA, Genes, Proteins



Adapted from *Understanding Gene Testing*, NIH, 1995

ASCO

Central Dogma

DNA



RNA



Protein

GENOMIC SOFTWARE
“The Code”

DNA → RNA CODONS → Protein

Genomic DNA

THYBIGRYDDOGRANOUT

Messenger RNA

THEBIGREDDOGRANOUT

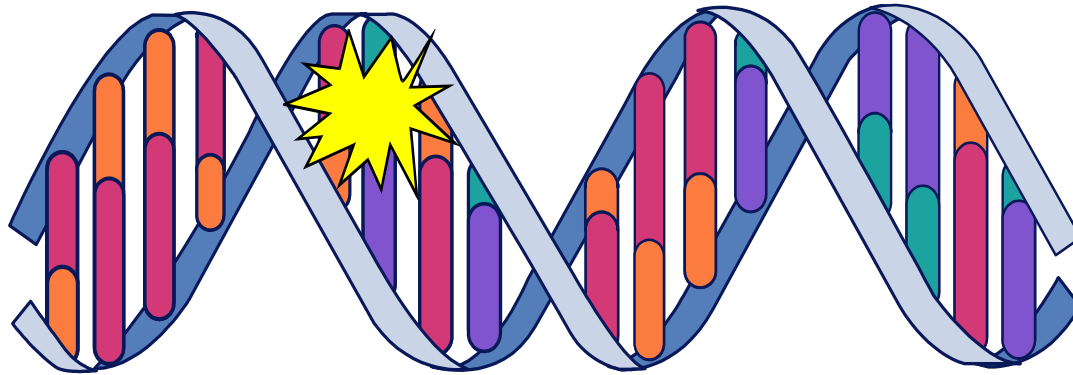
Translated Message

THE BIG RED DOG RAN OUT

Genetics of Disease

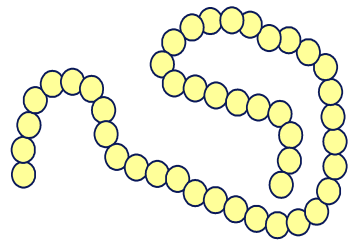
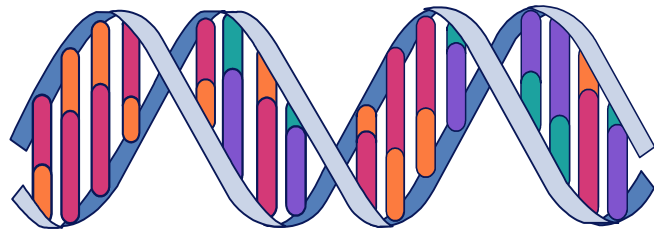
Disease-Associated Mutations

A **mutation** is a change in the normal base pair sequence

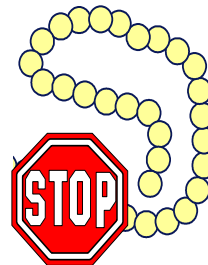
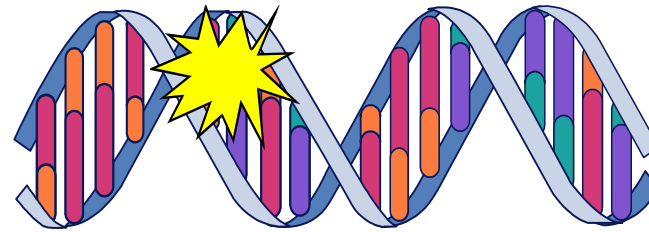


Commonly used to define DNA sequence changes that alter protein function

Disease-Associated Mutations Can Alter Protein Function



Functional protein



Nonfunctional or
missing protein

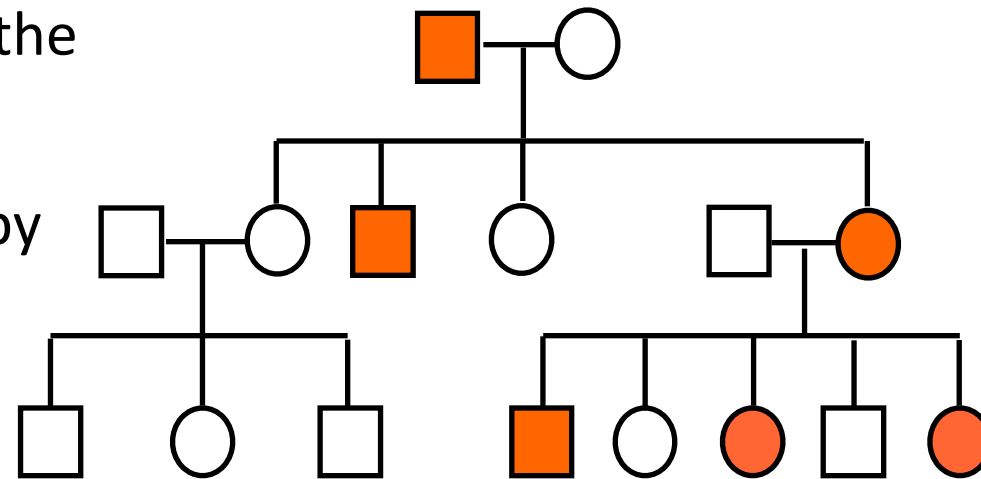
Point Mutations

Normal	THE BIG RED DOG RAN OUT.
Missense	THE BIG RAD DOG RAN OUT.
Nonsense	THE BIG RED.
Frameshift (deletion)	THE BRE DDO GRA.
Frameshift (insertion)	THE BIG RED ZDO GRA.

Point mutation: a change in a single base pair

Autosomal Dominant Inheritance

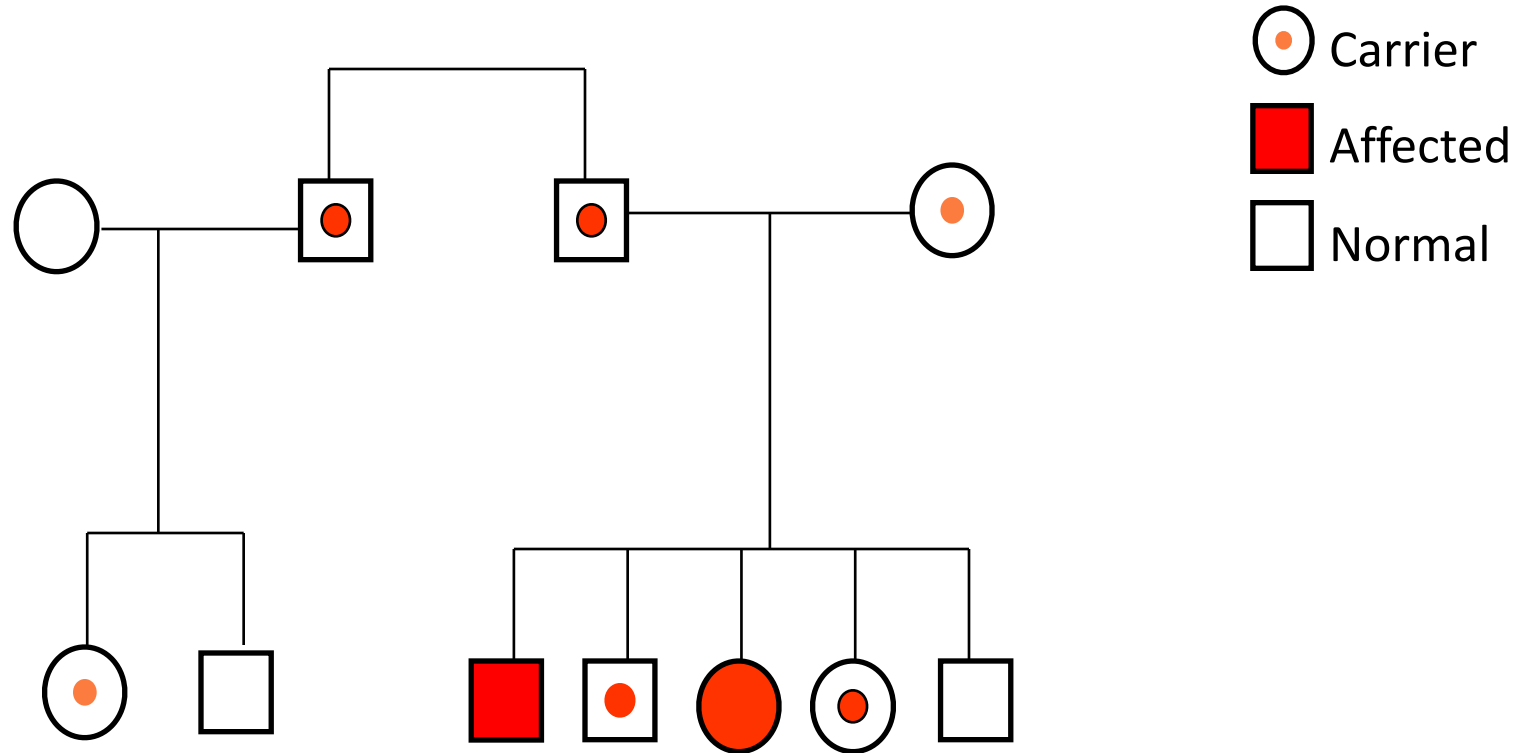
- Each child has 50% chance of inheriting the mutation
- Equally transmitted by men and women



□ ○ Normal

■ ● Affected

Autosomal Recessive Inheritance



Genetics 101 → 2017

DNA → RNA CODONS → Protein

Genomic DNA

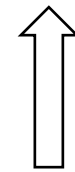
THYBIGRYDDOGRANOUT

Messenger RNA

THEBIGREDDOGRANOUT

Translated Message

THE BIG RED DOG RAN OUT



The code is read three letters at a time and the ATG sets the reading frame

DNA → RNA → Protein

Genomic DNA

THYBIGRYDRYDRYDRYDDOGRANOUT

Messenger RNA

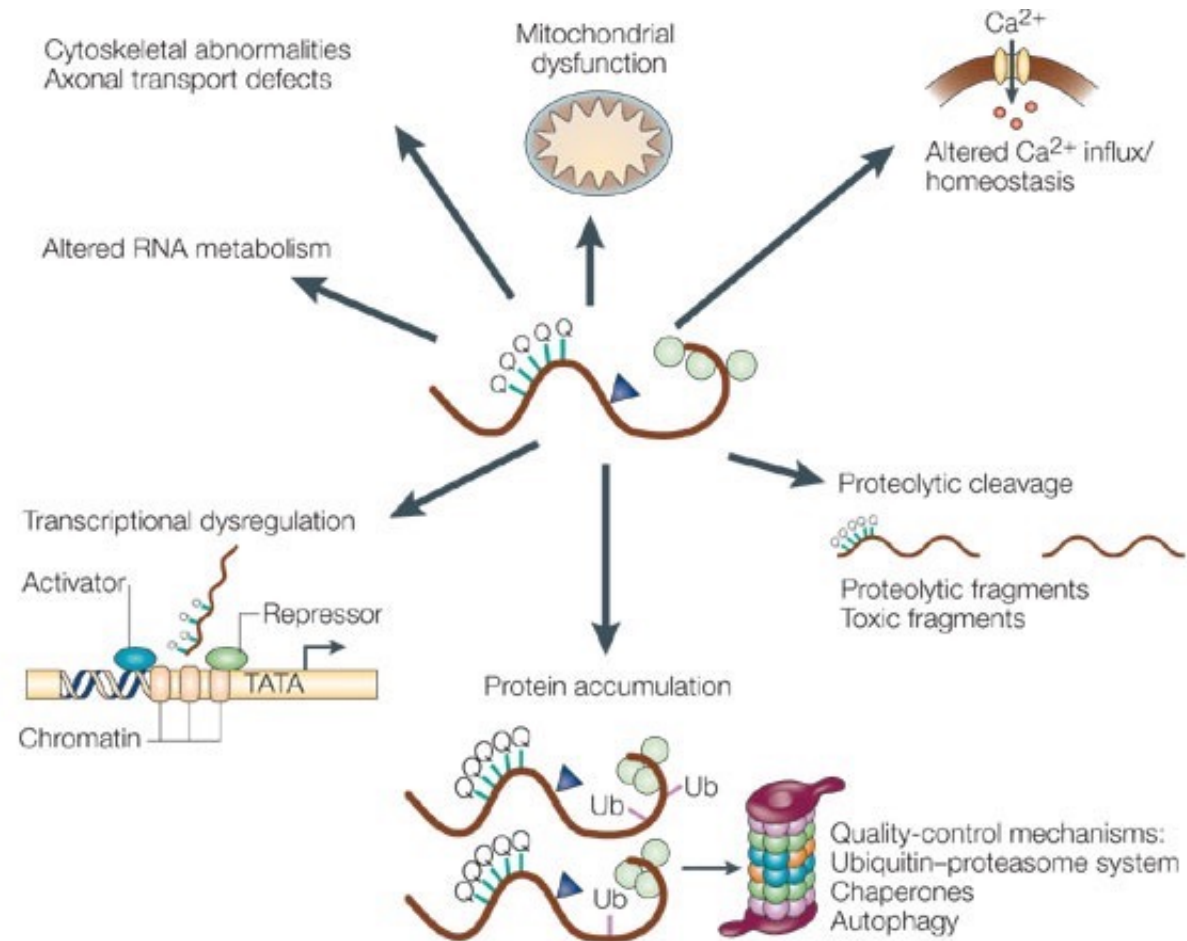
THEBIGREDREDREDREDDOGRANOUT

Translated Message



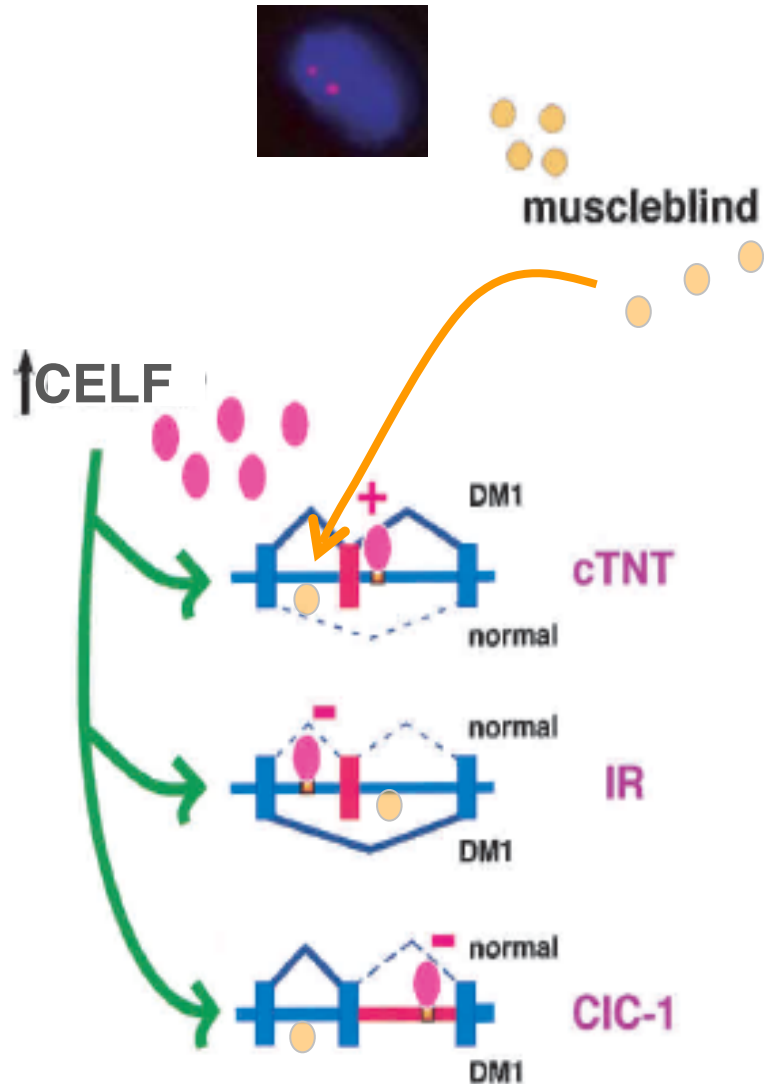
THE BIG RED RED RED RED DOG RAN OUT

PolyQ Protein Problems

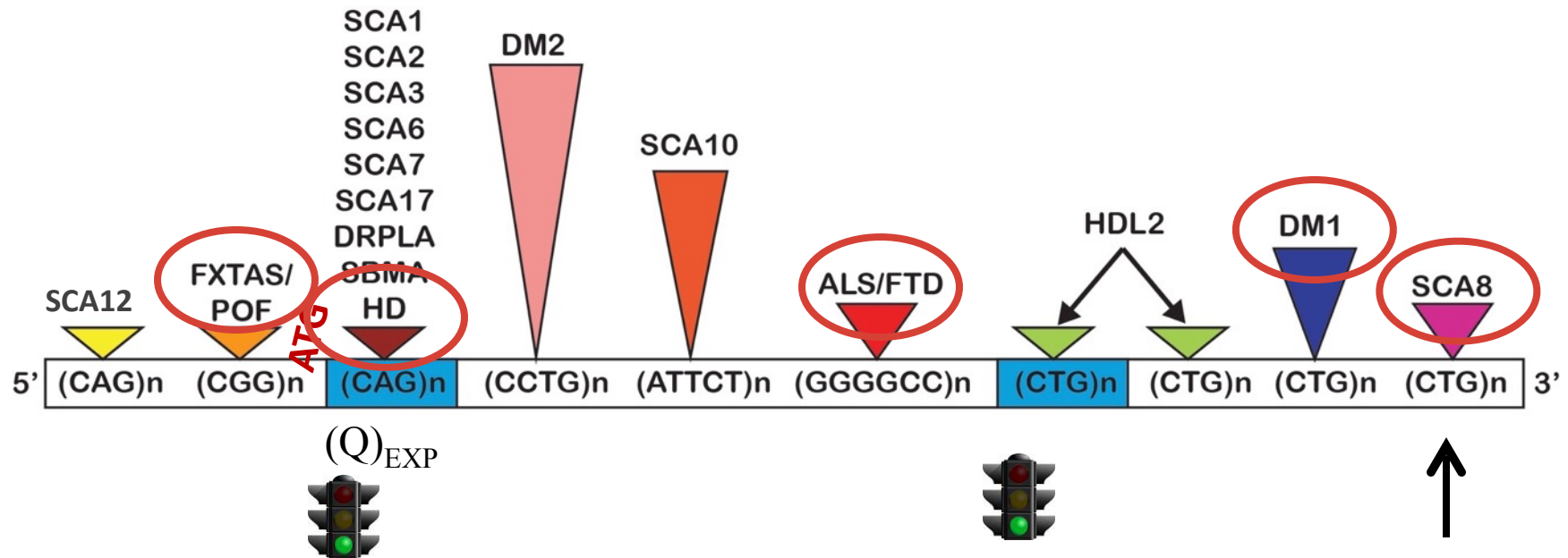


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Mutant RNA problems – myotonic dystrophy



Repeat Expansion Mutations in Neurologic Disease



Protein Loss / Protein Gain / RNA Gain of Function

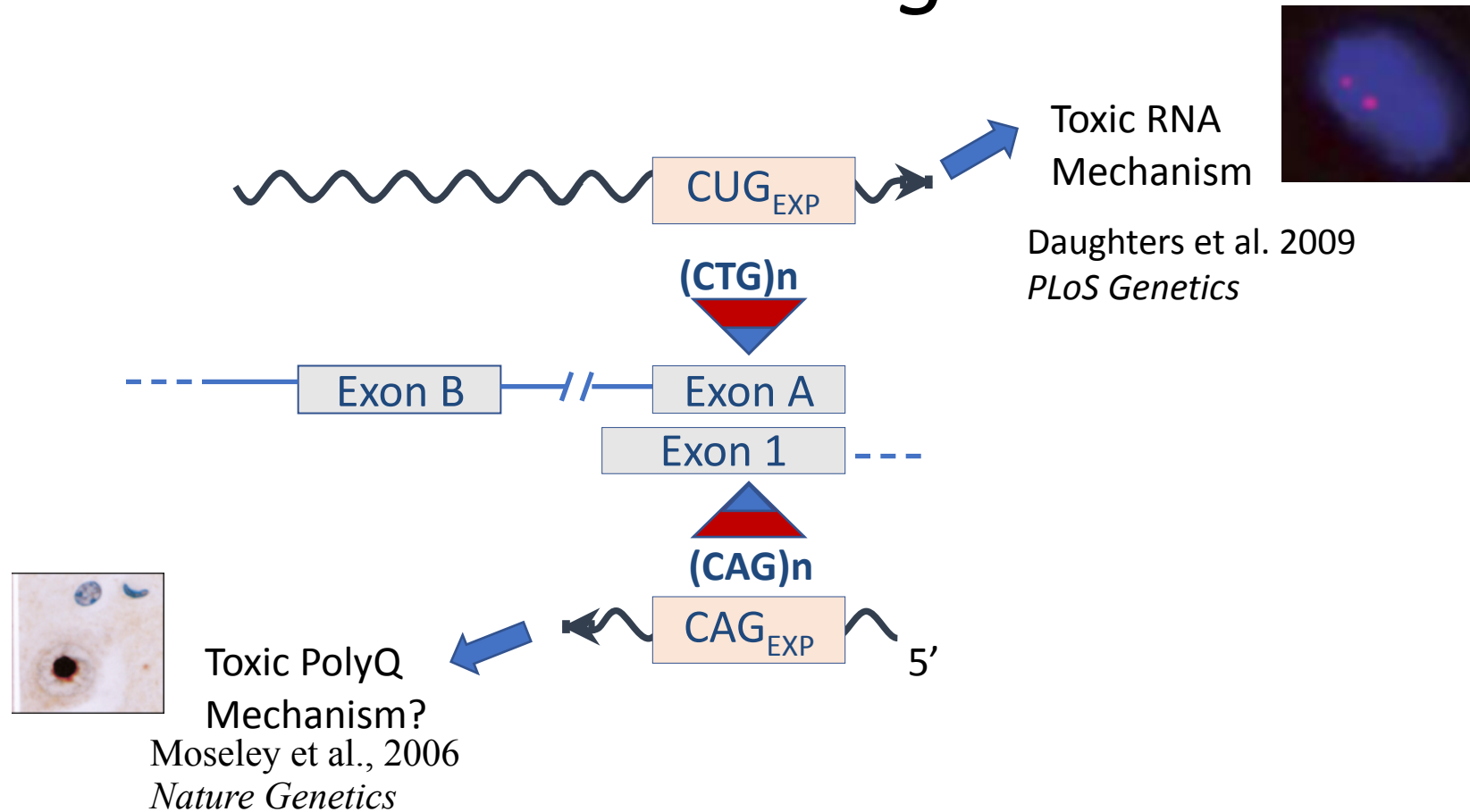
Bidirectional Expression

Repeat Associated Non-ATG (RAN) translation

Expansion mutations produce unexpected proteins in multiple frames

Spinocerebellar Ataxia Type 8
SCA8

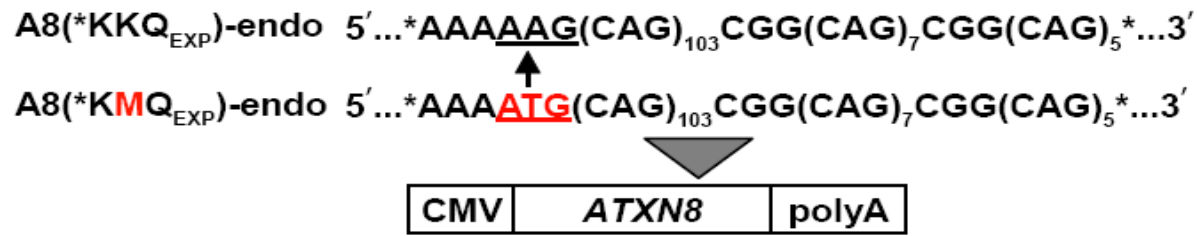
SCA8: one mutation two genes



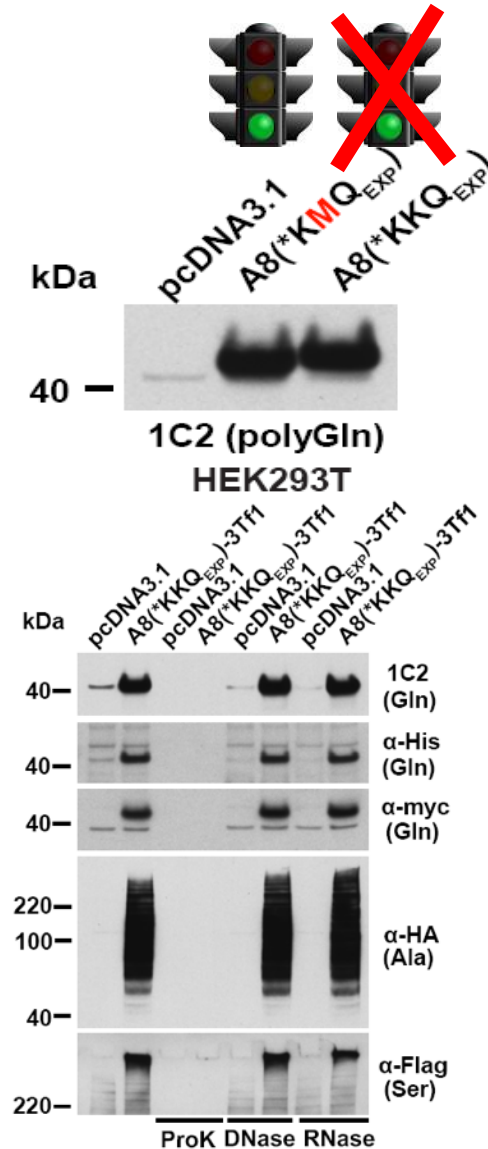
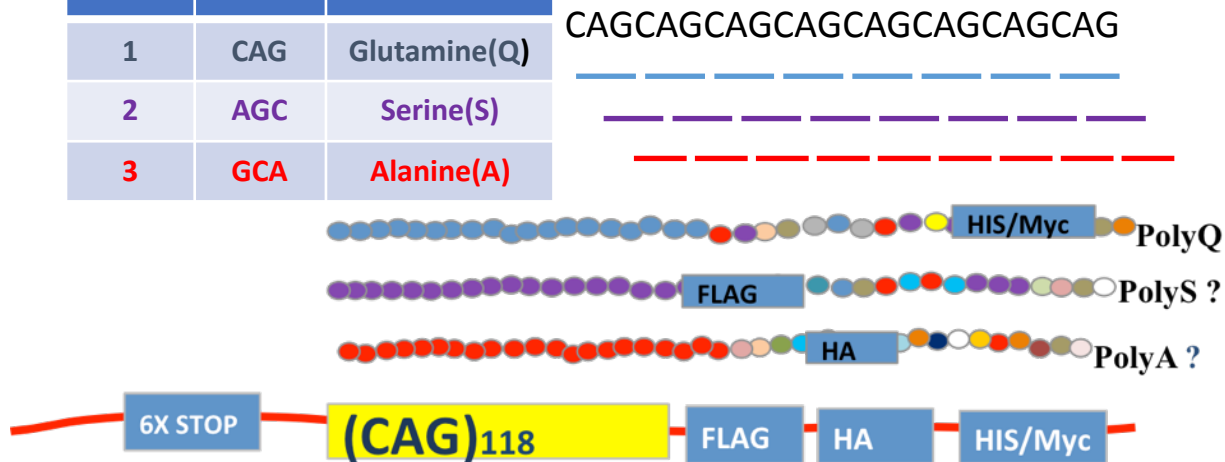
*AAA**ATG**(CAG)_{EXP}***



Discovery of Repeat-Associated Non-ATG Translation (RAN translation)



Frame	Codon	Amino Acid
1	CAG	Glutamine(Q)
2	AGC	Serine(S)
3	GCA	Alanine(A)



DNA → RNA → Protein

Genomic DNA

THYBIGRYDRYDRYDRYDDOGRANOUT

Messenger RNA

THEBIGREDREDREDREDDOGRANOUT

Translated Message

 THE BIG RED RED RED RED DOG RAN OUT

 RED RED RED RED DOG RAN OUT

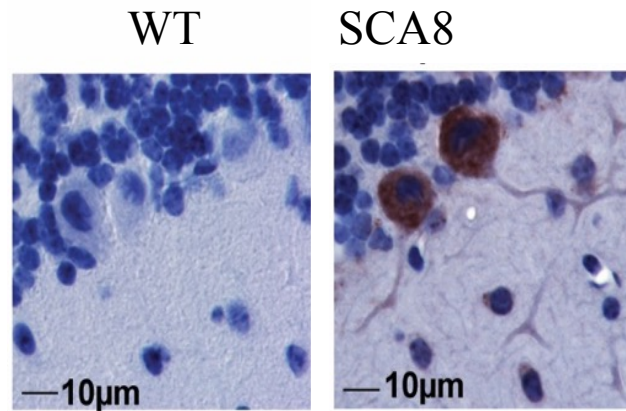
 EDR EDR EDR EDD OGR ANO UT

 DRE DRE DRE DDO GRA NOU T

Repeat Associated Non-ATG (RAN) translation

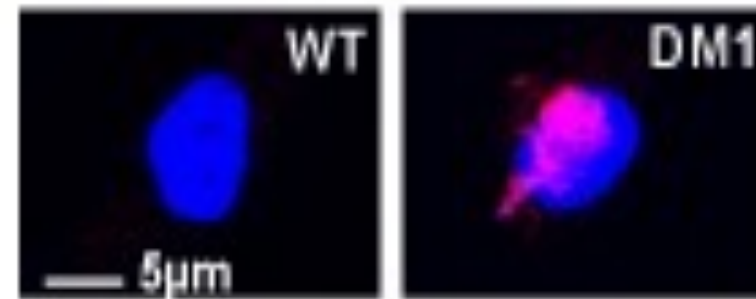
RAN Translation in vivo: SCA8 and myotonic dystrophy

α -SCA8 poly-GCA-Ala



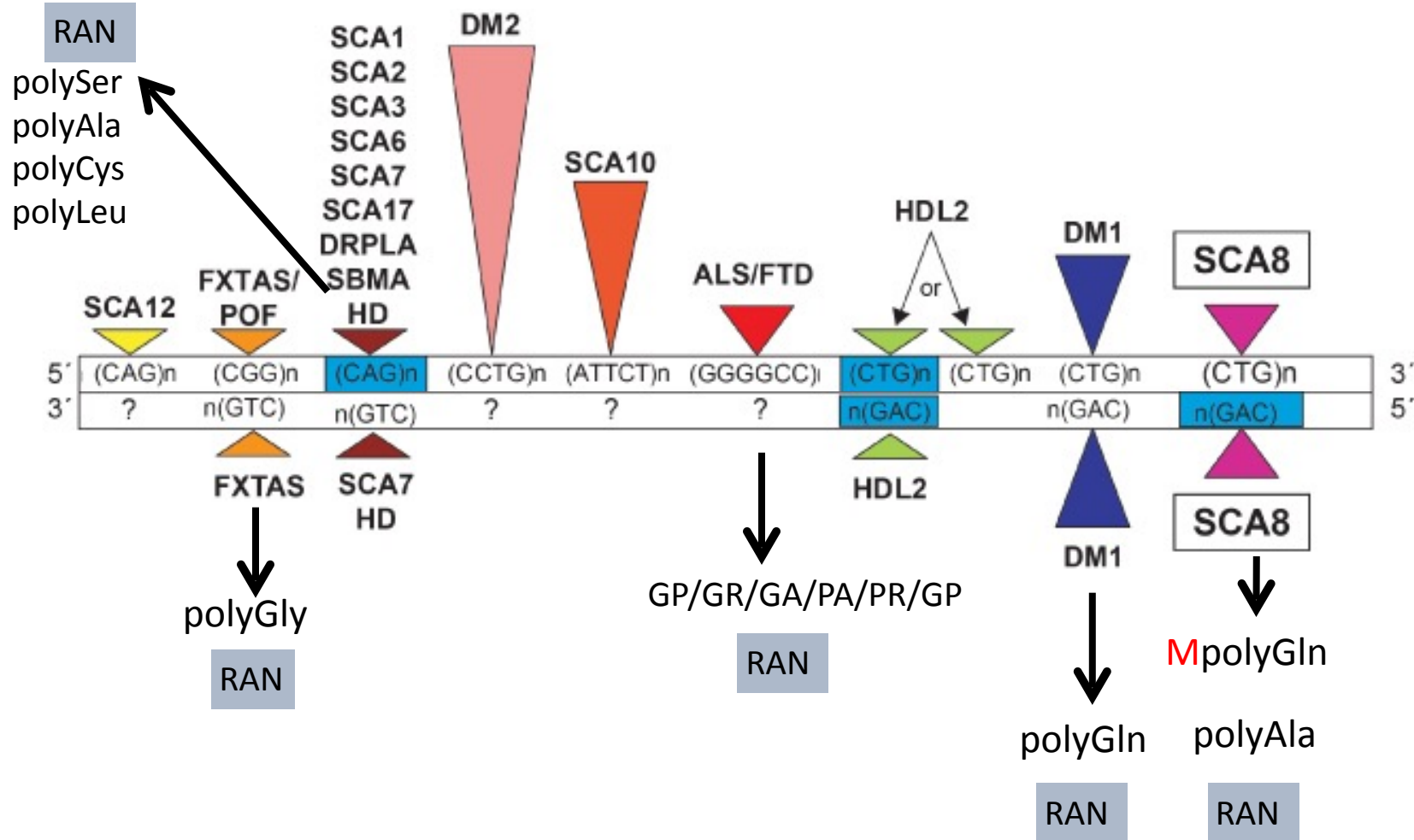
Mouse Cerebellum

α -DM1 poly-CAG-Gln



Human Myoblasts

Growing trends in many expansion diseases



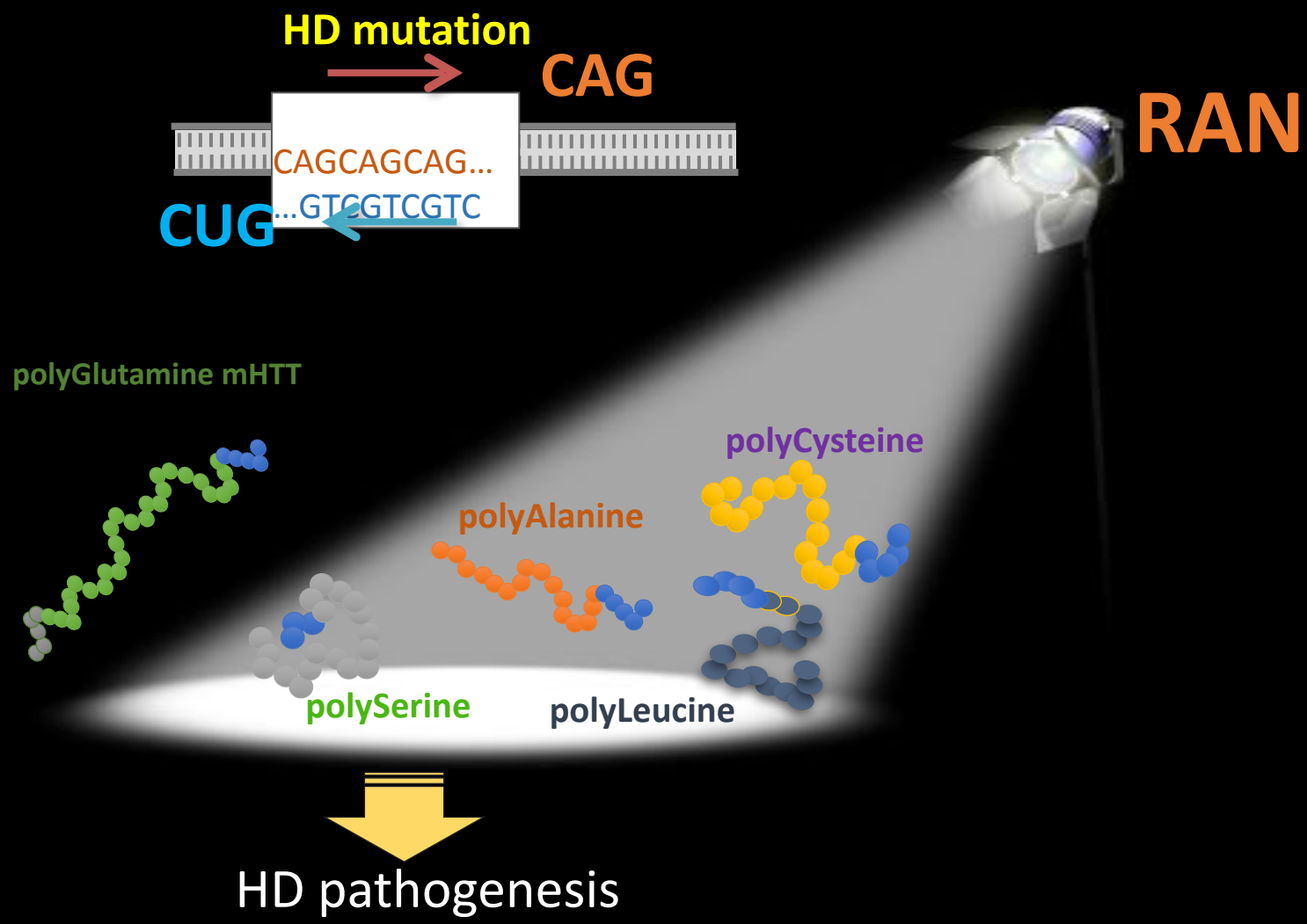
Framework for understanding disease mutations needs revision.

Bidirectional expression is common

Repeats express unexpected proteins

RAN proteins now found in RNA and "polyQ" diseases

RAN Proteins in Huntington's Disease



WHAT ABOUT THE POLY-Q ATAXIAS?

Therapy development

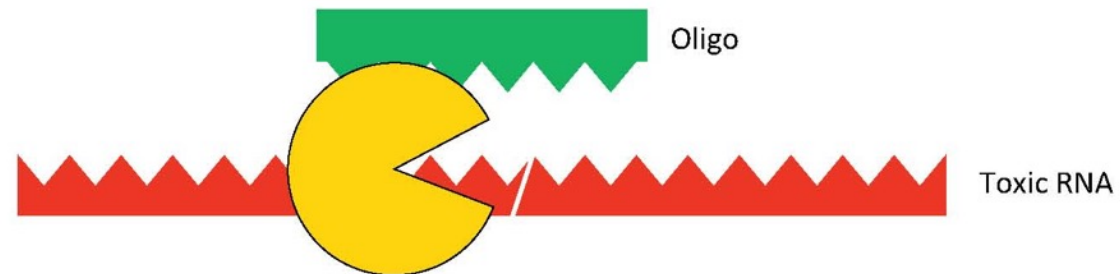
New Genetic Tools

Gene Editing

RNA Targeting

Targeting the RNA

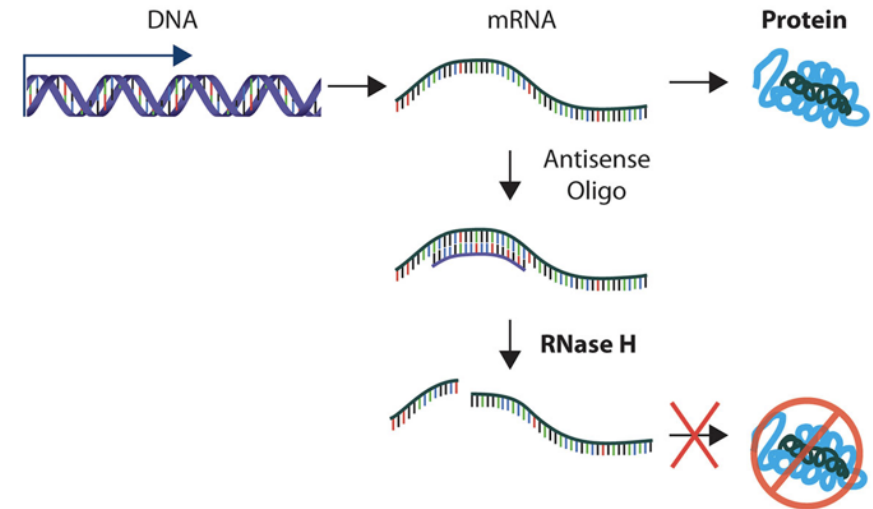
Targeted degradation of toxic RNA by
gapmer ASOs



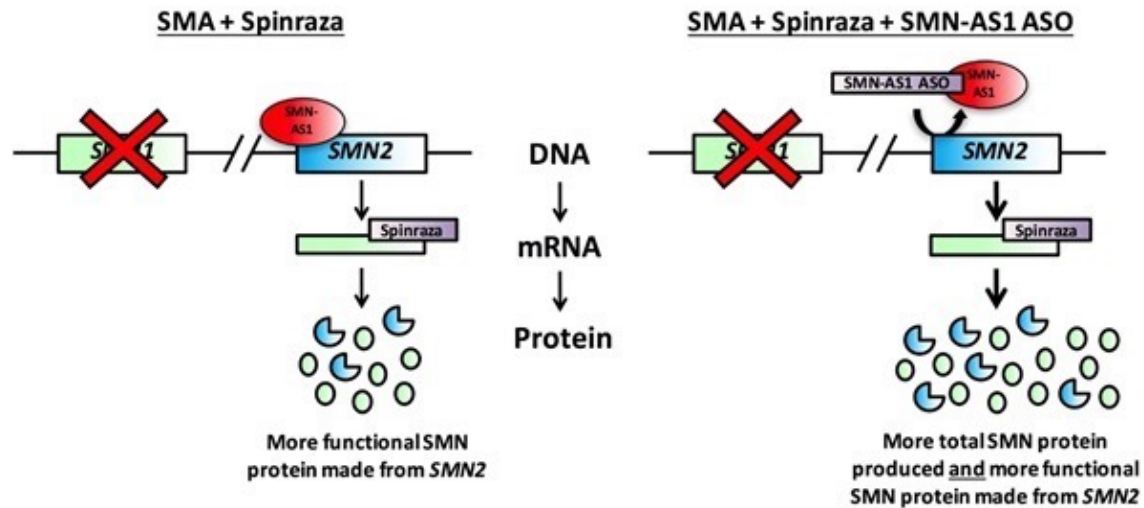
ANTISENSE OLIGONUCLEOTIDE THERAPY



FDA approval for Biogen and
Ionis Pharmaceuticals –
Spinraza
Spinal Muscular Atrophy

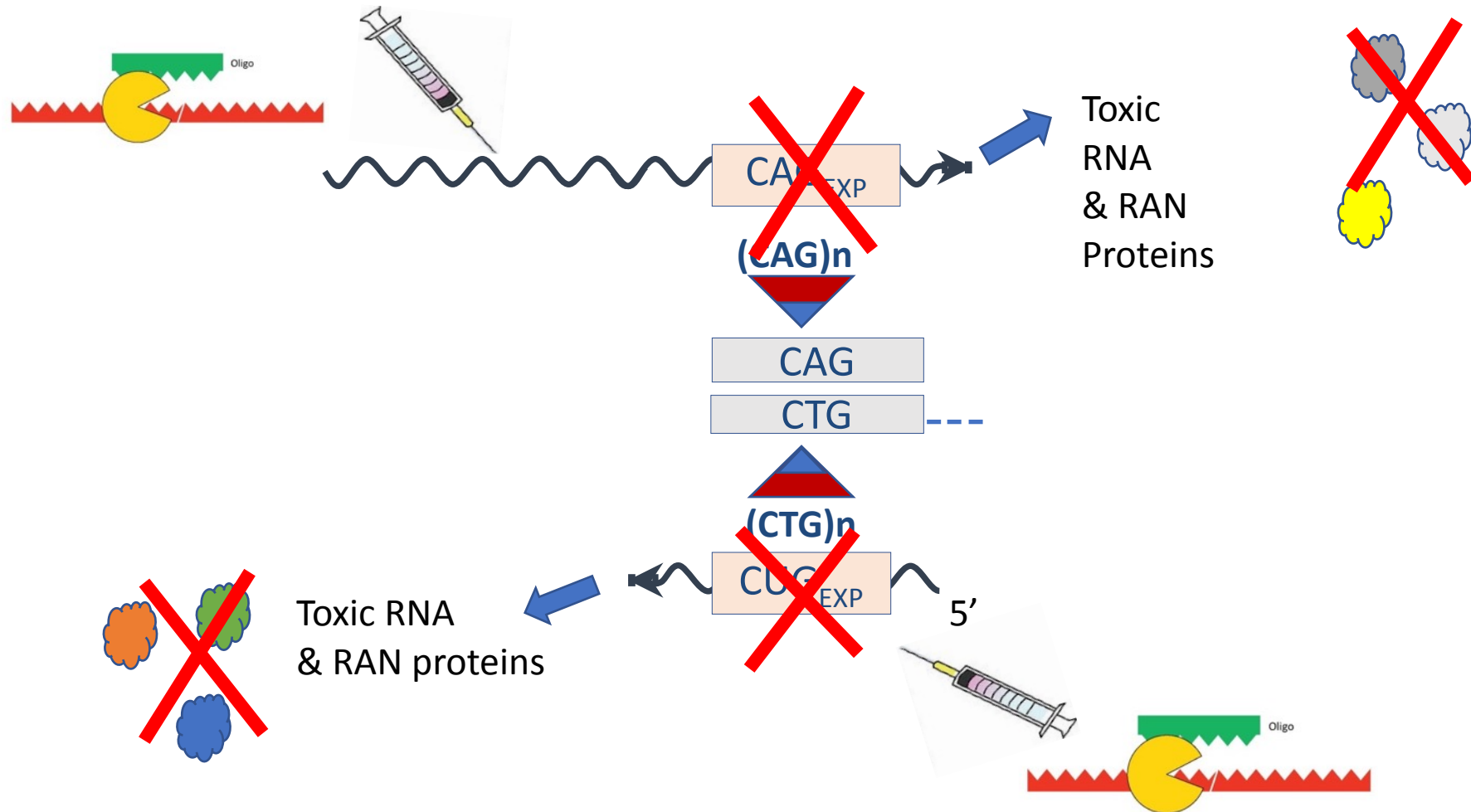


SIMILAR DRUGS CAN REMOVE RNA



DRUG WAKES UP A SLEEPING GENE

SCA8 targeting the problems: one or both RNAs?



Similar strategies being applied for many of the SCAs including SCA1 and SCA3

PARTNERING WITH YOU

GENE DISCOVERY

HOW GENETIC MUTATIONS CAUSE DISEASE

THERAPY DEVELOPMENT

Conclusions

THANK YOU!!!!!!

- NAF funding critically important for understanding the causes of ataxia and developing treatments
- Therapeutic efforts to target RNA and protein pathways and to repair genes are underway.
- Lessons from one disease connect to other diseases
 - How mutations cause disease
 - Therapy development

