

Genetic Approaches to Tackle Dominantly Inherited Ataxias

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Spinocerebellar Ataxia Type 1 (SCA1)



Dominantly inherited

Affects 1/100,000 people

Loss of balance and coordination

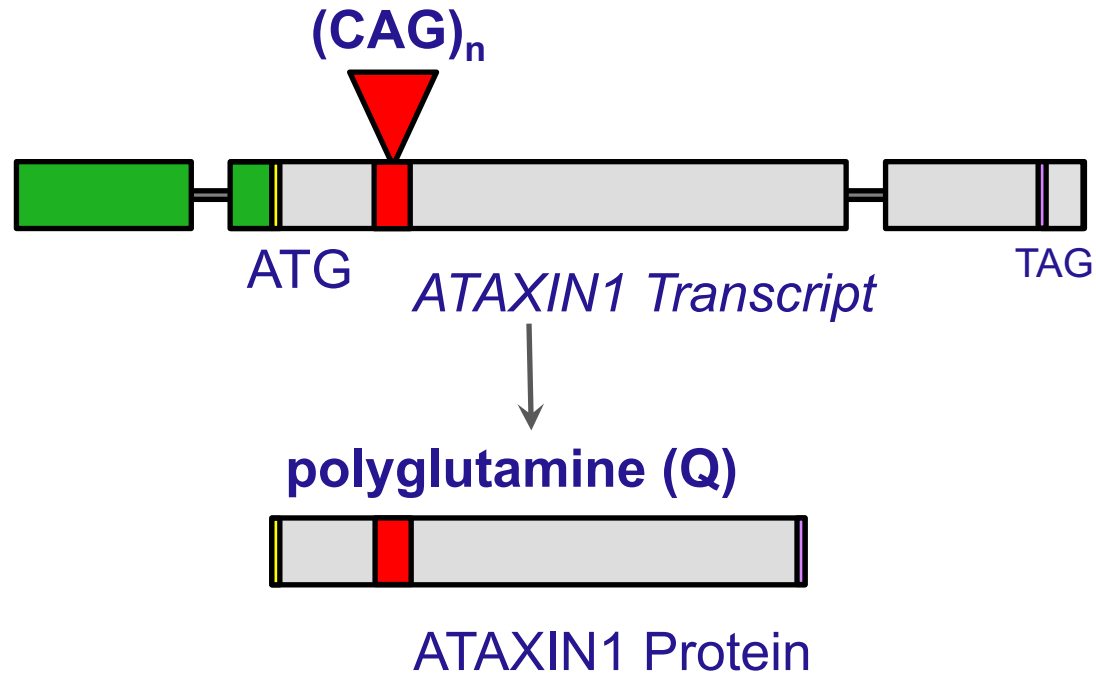
Difficulty with swallowing & breathing

Loss of cerebellar Purkinje cell and brainstem neurons

Premature death

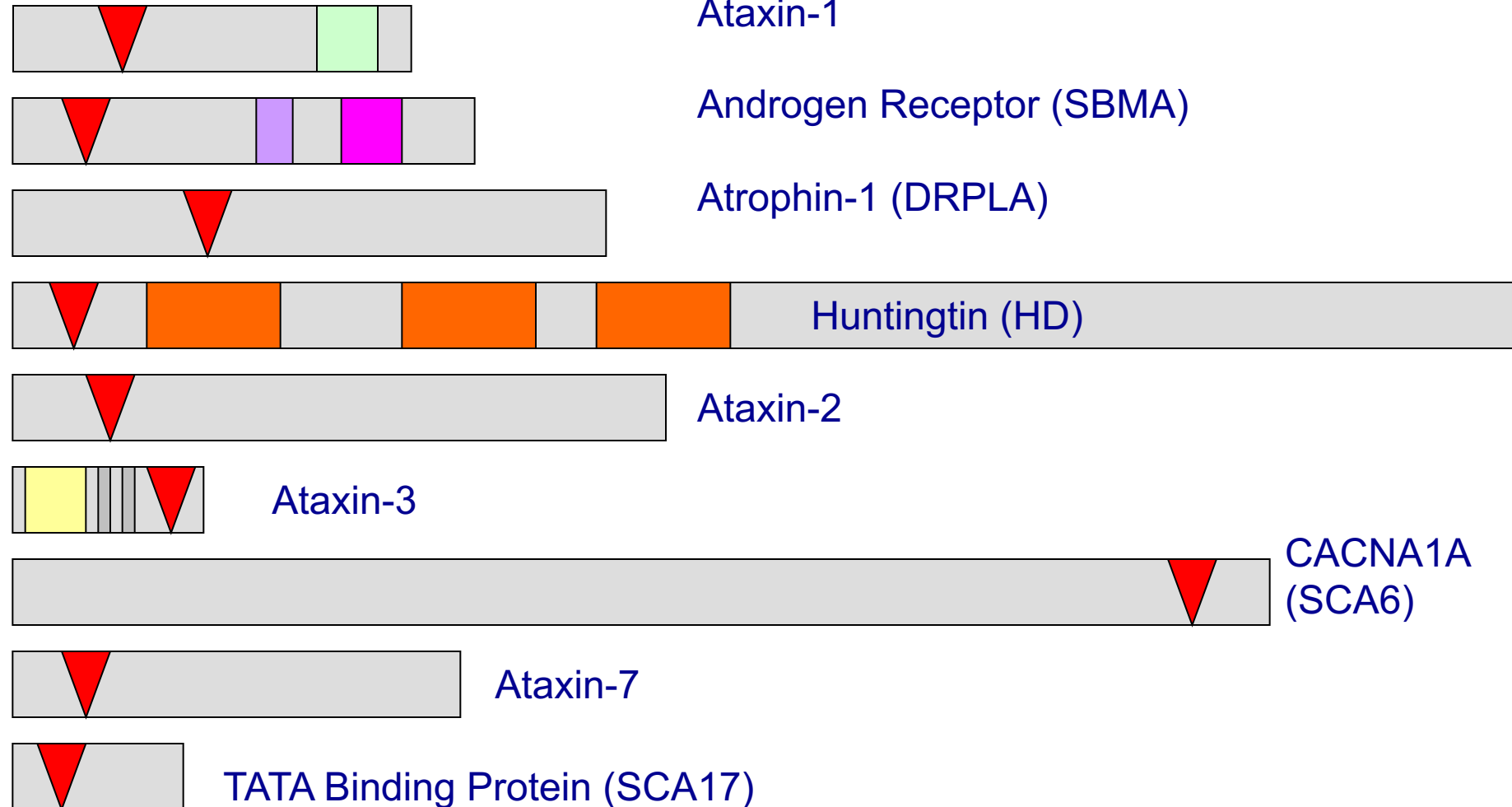
SCA1 is caused by expansion of unstable CAG repeats

39-82 repeats
(affected)



Orr *et al*, *Nat. Genet.*, 1993

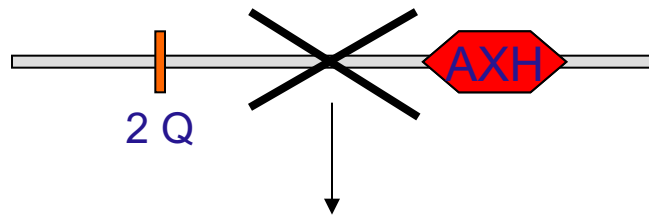
SCA1 is one of nine polyglutamine diseases



SCA1 pathogenesis is caused mainly by a gain-of-function mechanism



Atxn1^{-/-}

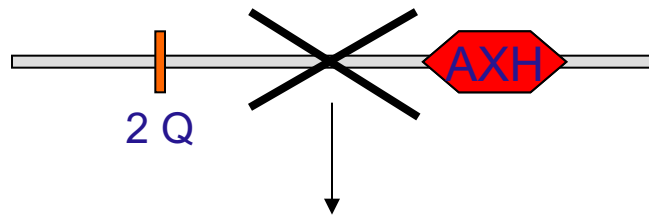


*No SCA1 phenotypes,
but some neurological
deficits*

SCA1 pathogenesis is caused mainly by a gain-of-function mechanism



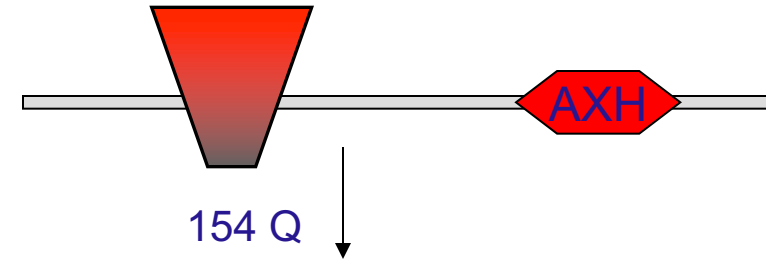
Atxn1^{-/-}



*No SCA1 phenotypes,
but some neurological
deficits*



Atxn1^{154Q/+}



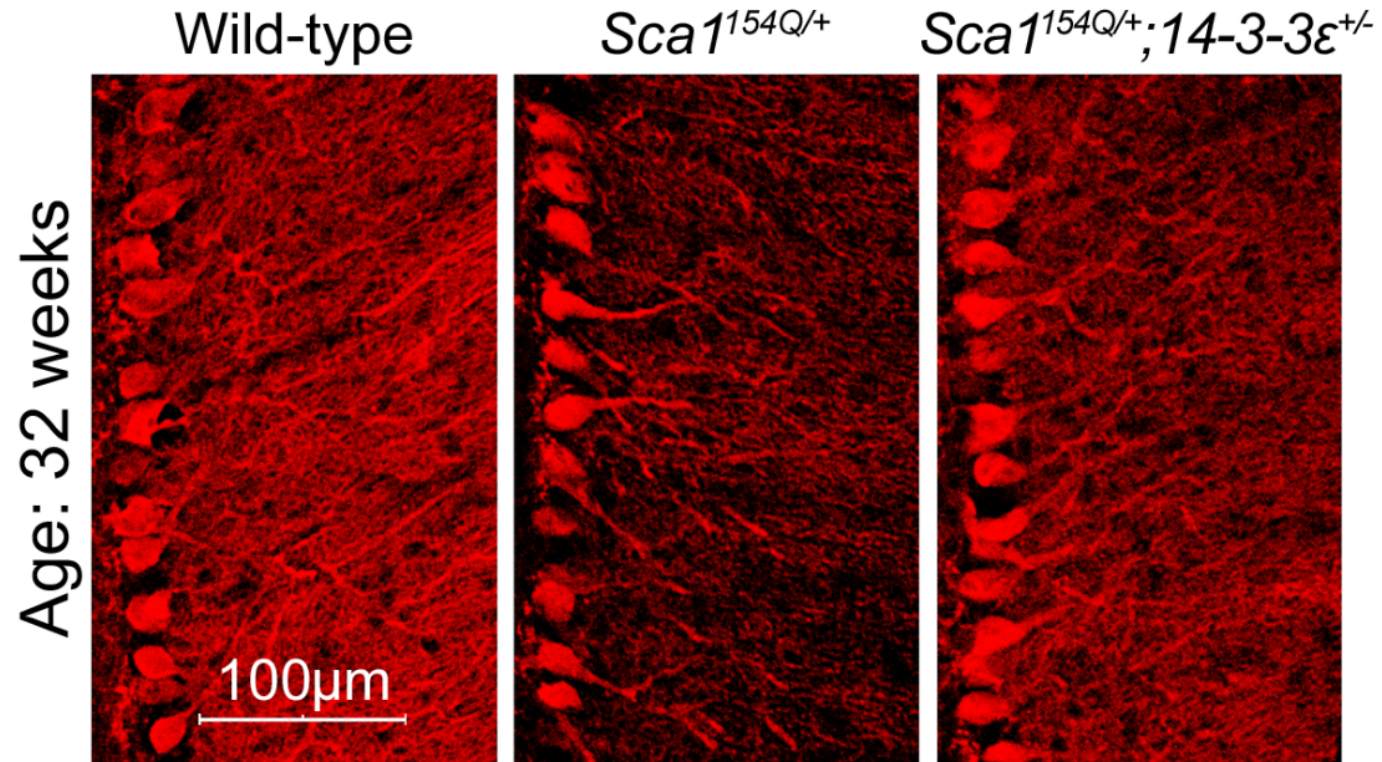
Recapitulate SCA1 phenotypes
Ataxia
Brain stem dysfunction
Purkinje cell loss
Reduced lifespan

Insights from genetic & biochemical studies



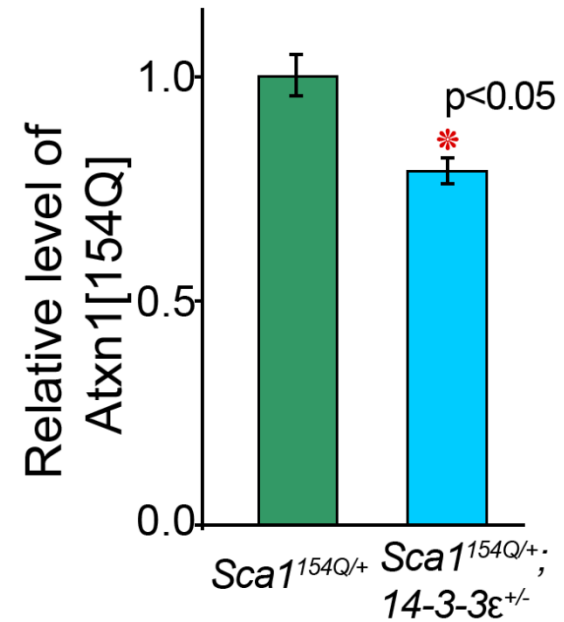
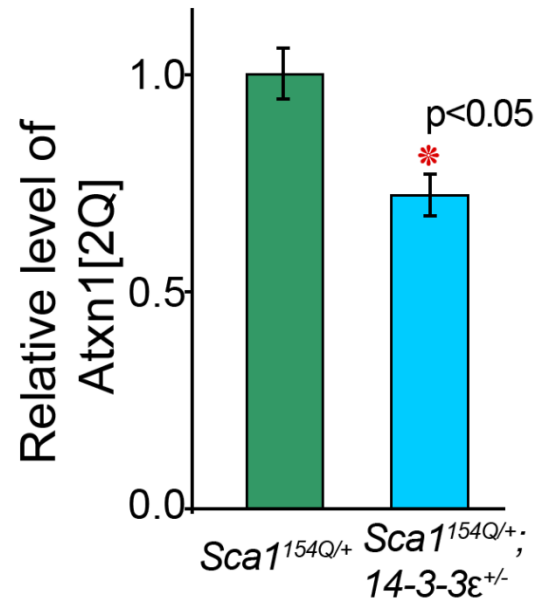
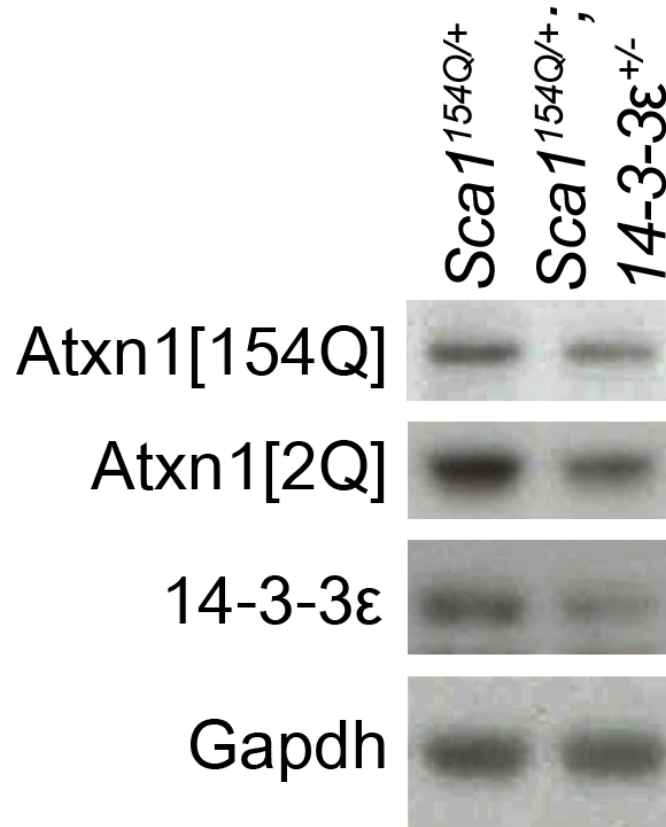
- The glutamine (Q) expansion in Ataxin-1 (ATXN1) makes ATXN1 more stable (resistant to degradation), so its interactions with its protein partners are prolonged and ATXN1 gradually accumulates in cells
- Increased levels of even wild-type ATXN1 cause neurodegeneration
- Phosphorylation at serine 776 in ATXN1 is necessary for SCA1 pathogenesis

Reducing 14-3-3 ϵ rescues Purkinje cell degeneration in SCA1 mice



Calbindin immunofluorescence

Reducing 14-3-3 ϵ decreases the levels of ATXN1 in SCA1 mice

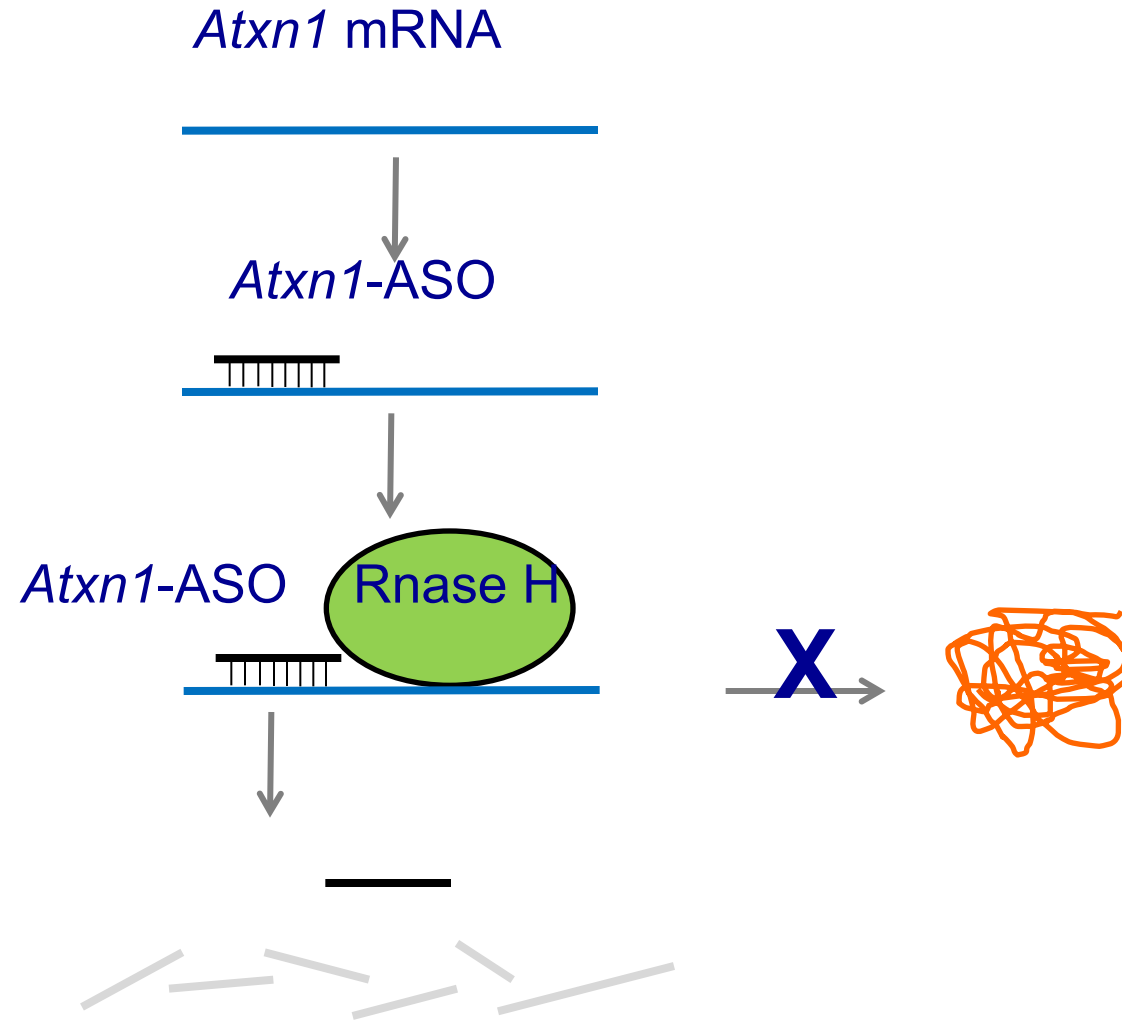


Would lowering ATXN1 levels using a strategy that directly target ATXN1 mRNA (Anti Sense Oligonucleotide—ASO) alleviate symptoms?

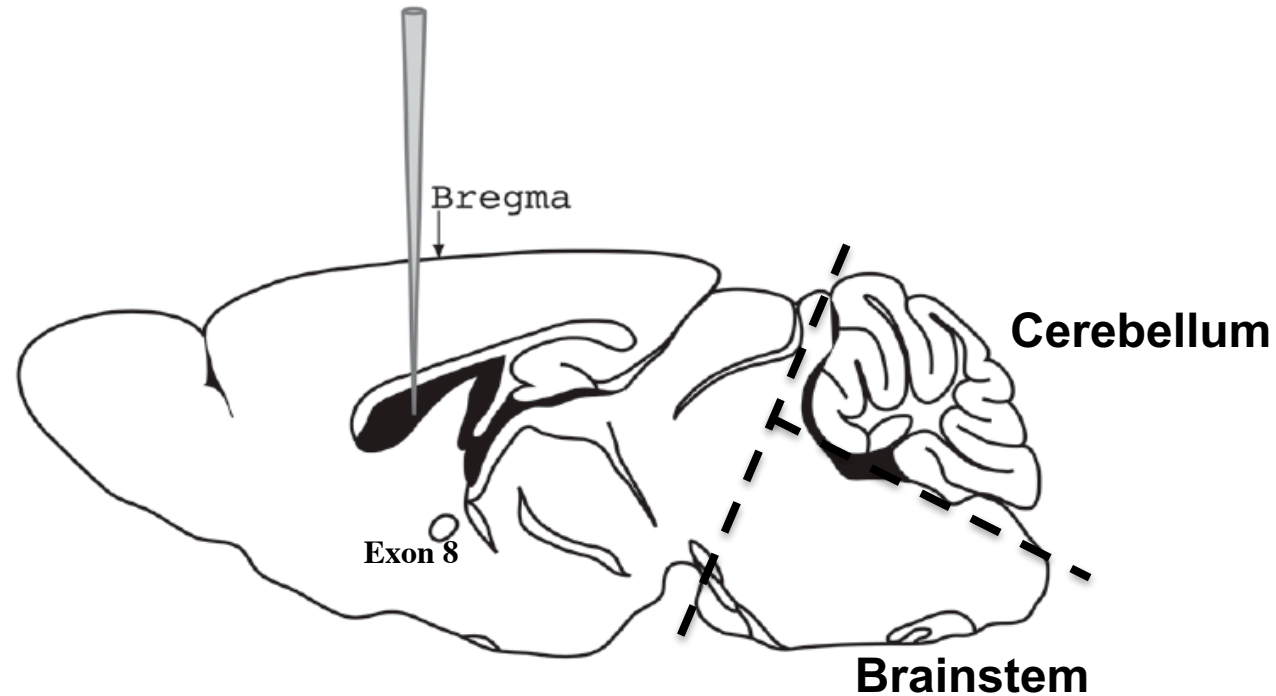
Orr lab in collaboration with Holly Kordasiewicz



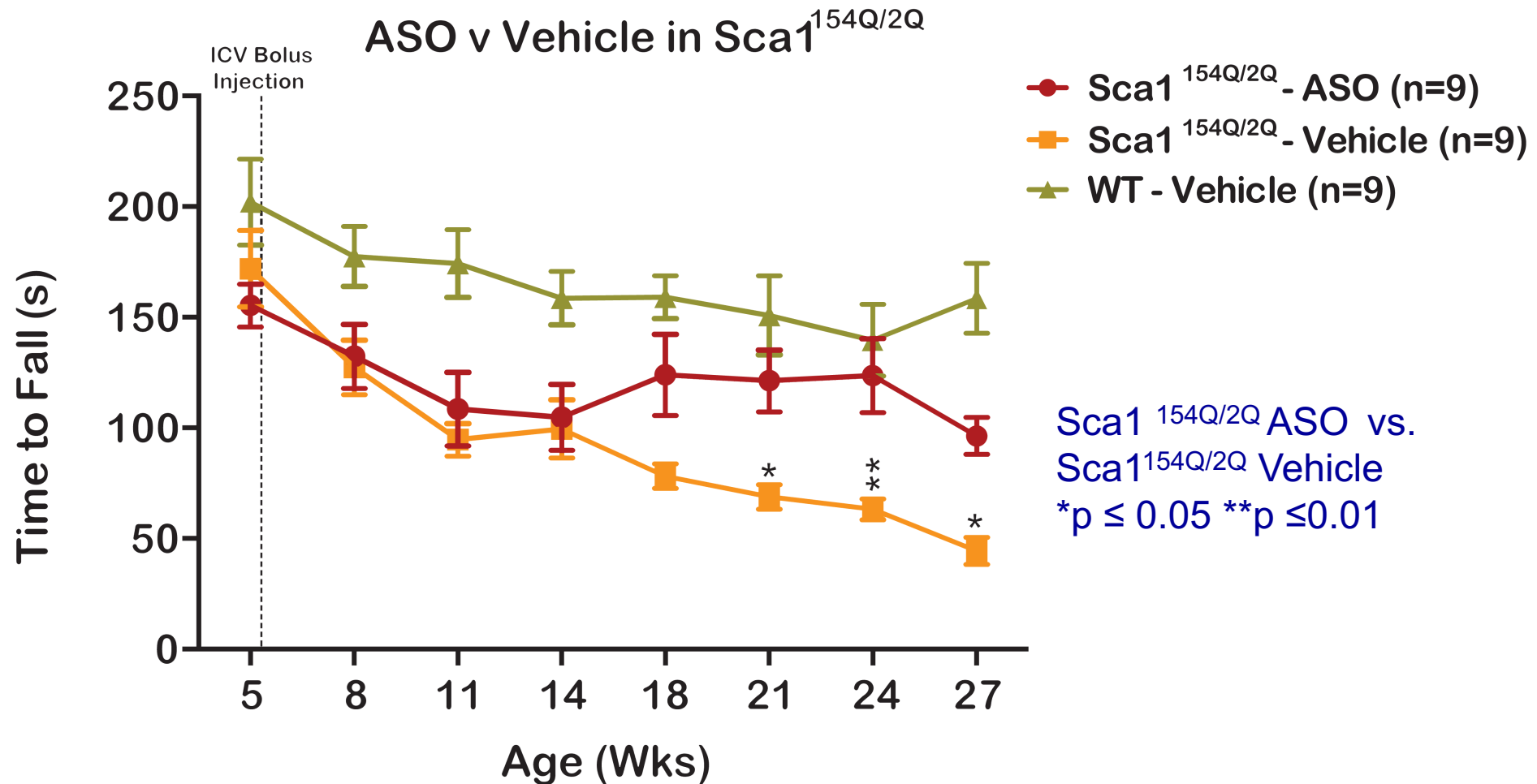
Molecular mechanism of ASO-mediated gene silencing



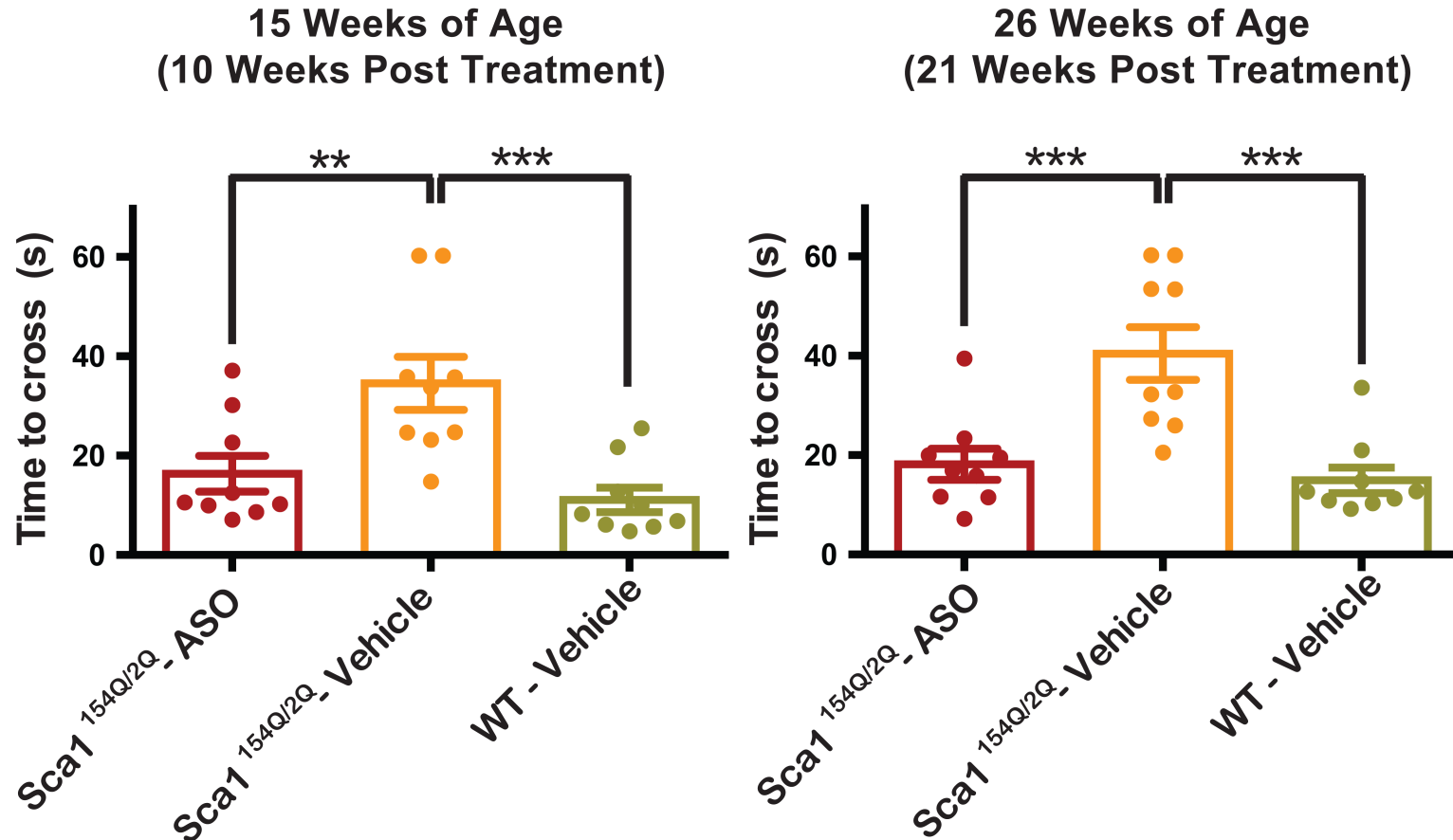
Intraventricular delivery of 500 μ g Atxn1-ASO to five-week old *Sca1*^{154Q/+} mice



Atxn1-ASO improves rotating rod performance in *Sca1*^{154Q/+} mice



Atxn1-ASO improves beam crossing in *Sca1*^{154Q/+} mice

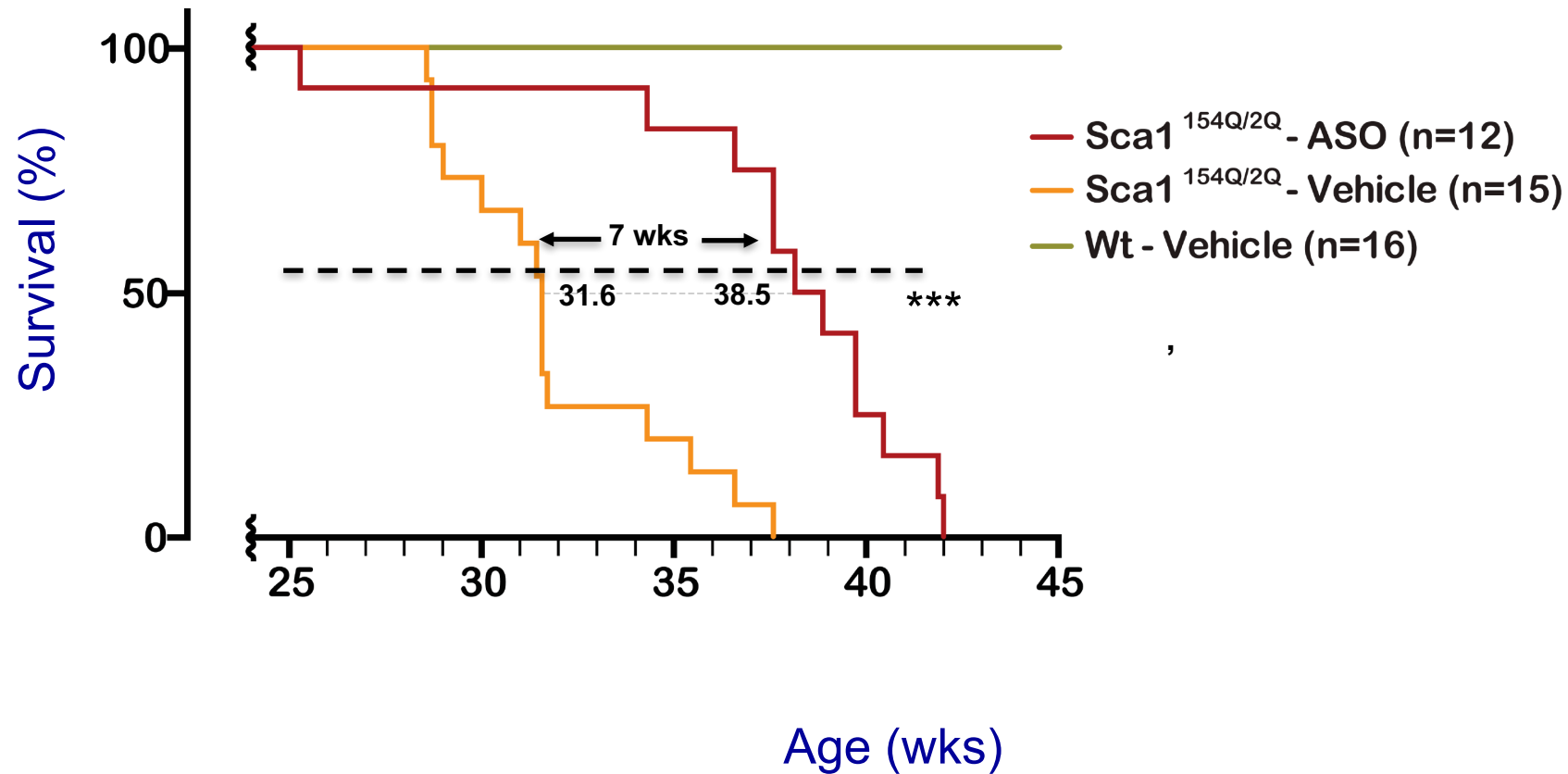


* $p \leq 0.05$ ** $p \leq 0.01$

Jill Friedrich

- *Sca1*^{154Q/2Q} - ASO (n=9)
- *Sca1*^{154Q/2Q} - Vehicle (n=9)
- ▲ WT - Vehicle (n=9)

Atxn1-ASO improves survival of *Sca1*^{154Q/+} mice



***p=0.0001

Jill Friedrich

ASO therapy advantages and limitations

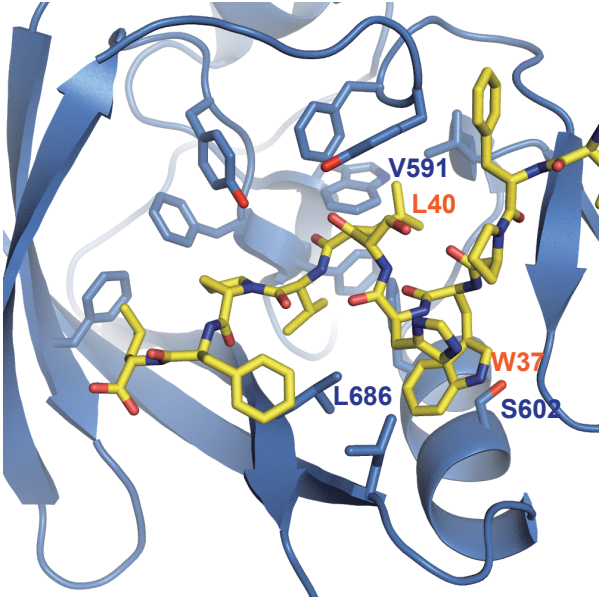
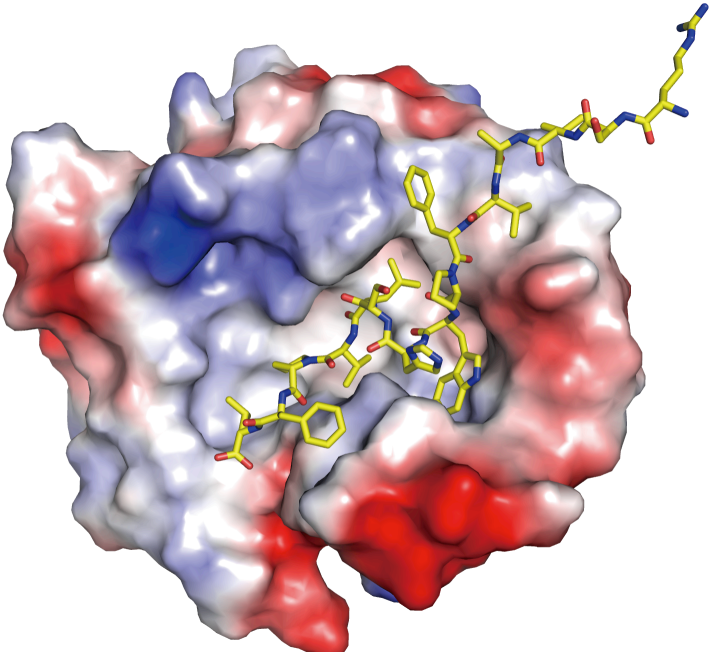
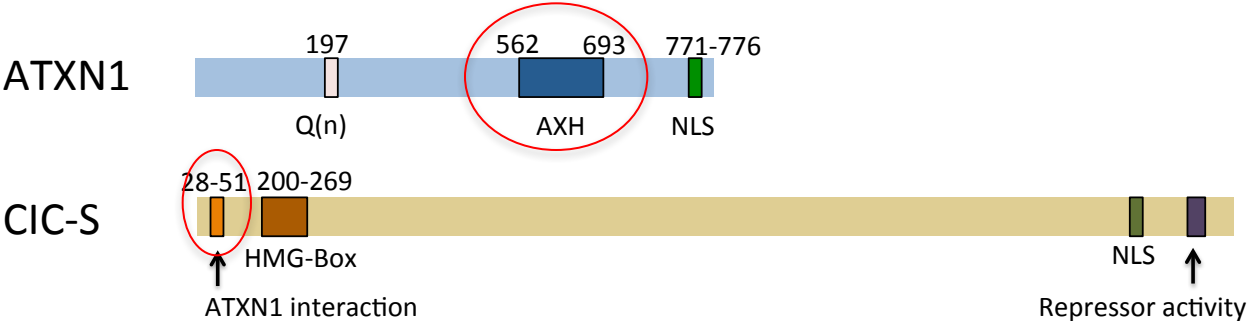
Advantages:

- It is specific and targets the disease-driving protein
- It can be rapidly advanced into clinical trials given precedent for other neurological disorders (SMA)

Limitations:

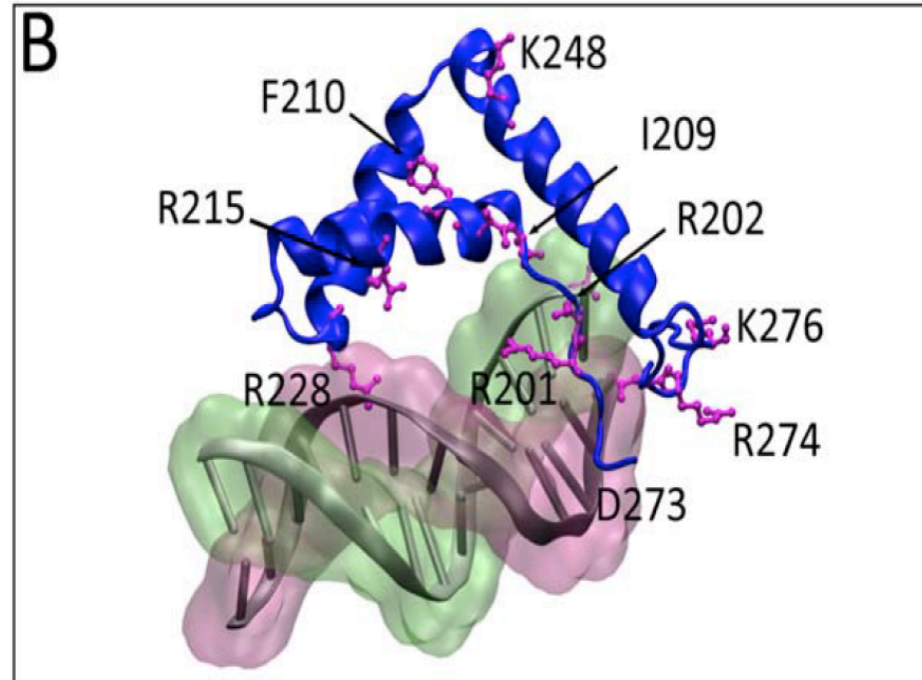
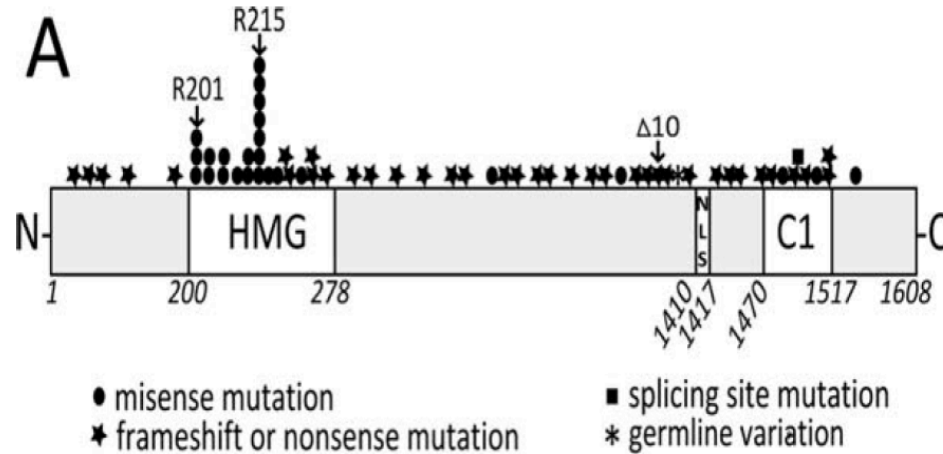
- It requires monitoring of the target protein (ATXN1) levels to insure that the reduction is mild to moderate (20-40%) given the important role of ATXN1 with Capicua
- It is invasive requiring repeated intrathecal injections

ATXN1-CIC interact and function together for normal brain function and to suppress tumors

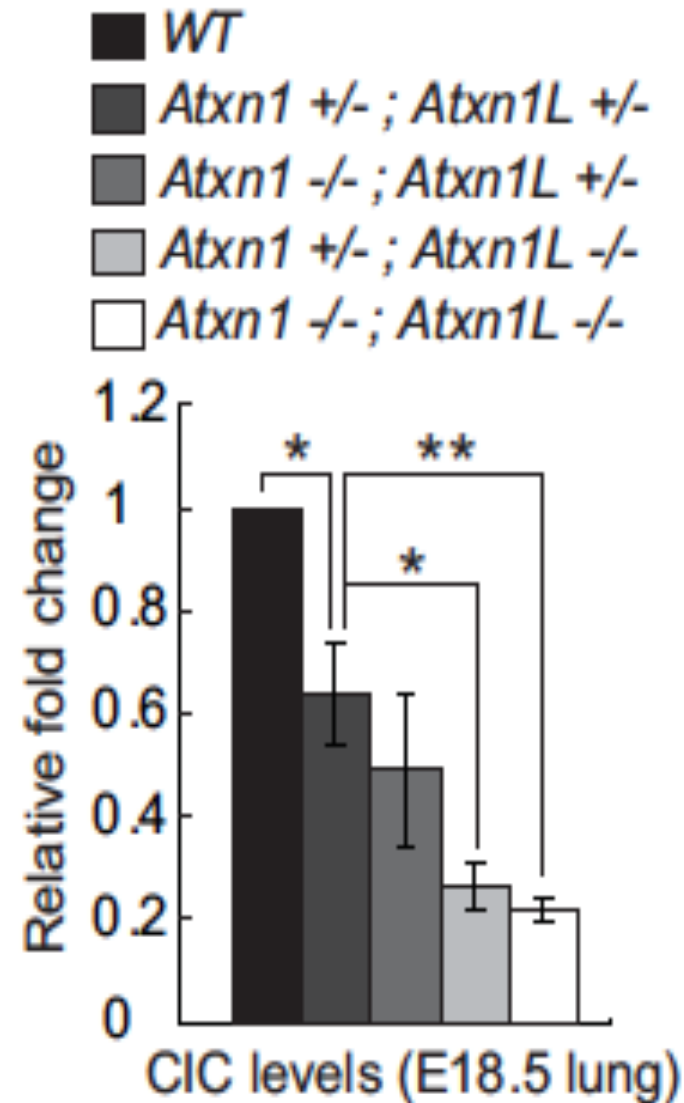


(Kim E, et al., *Genes and Development* 2013)

C/C mutations were observed in 60 patients with oligodendroglioma



Capicua stability depends on ATXN1 and ATXN1Like levels

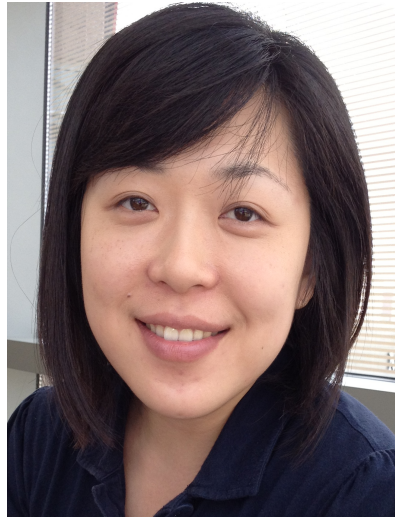


Ongoing studies to optimize feasibility of translational ASO therapy

- Establish the extent of the effects of ASO therapy on ATXN1 levels and CIC levels
- Determine the ideal dose that partially lowers ATXN1 without impacting CIC levels and that is effective
- Determine the distribution of ASO after intrathecal injection in non-human primates and its effects on ATXN1 and CIC levels

Can we develop a complementary strategy
that barely lowers ATXN1 and has the
potential of being used chronically

There should be targets that, when inhibited,
reduce ATAXIN1 levels and could provide
therapeutic entry points for SCA1

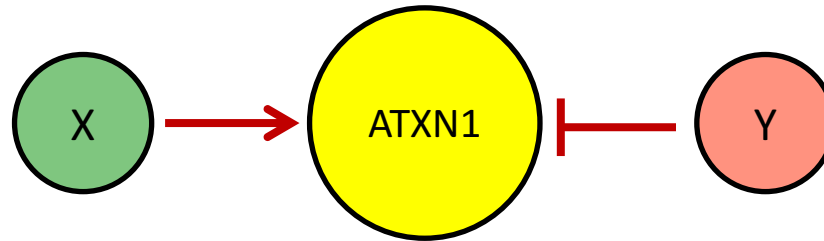


Jeehye Park, PhD

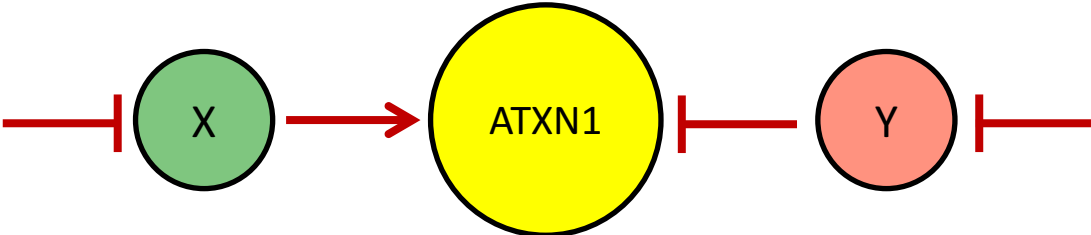


Ismael Al-Ramahi, PhD

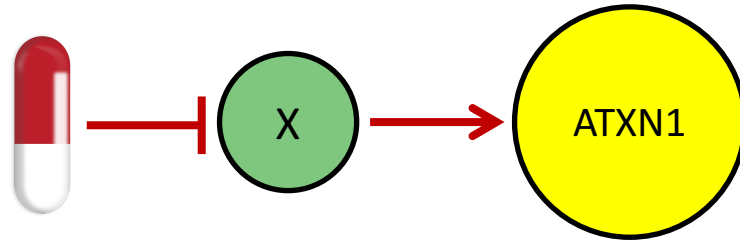
Find genes that regulate ATXN1 levels



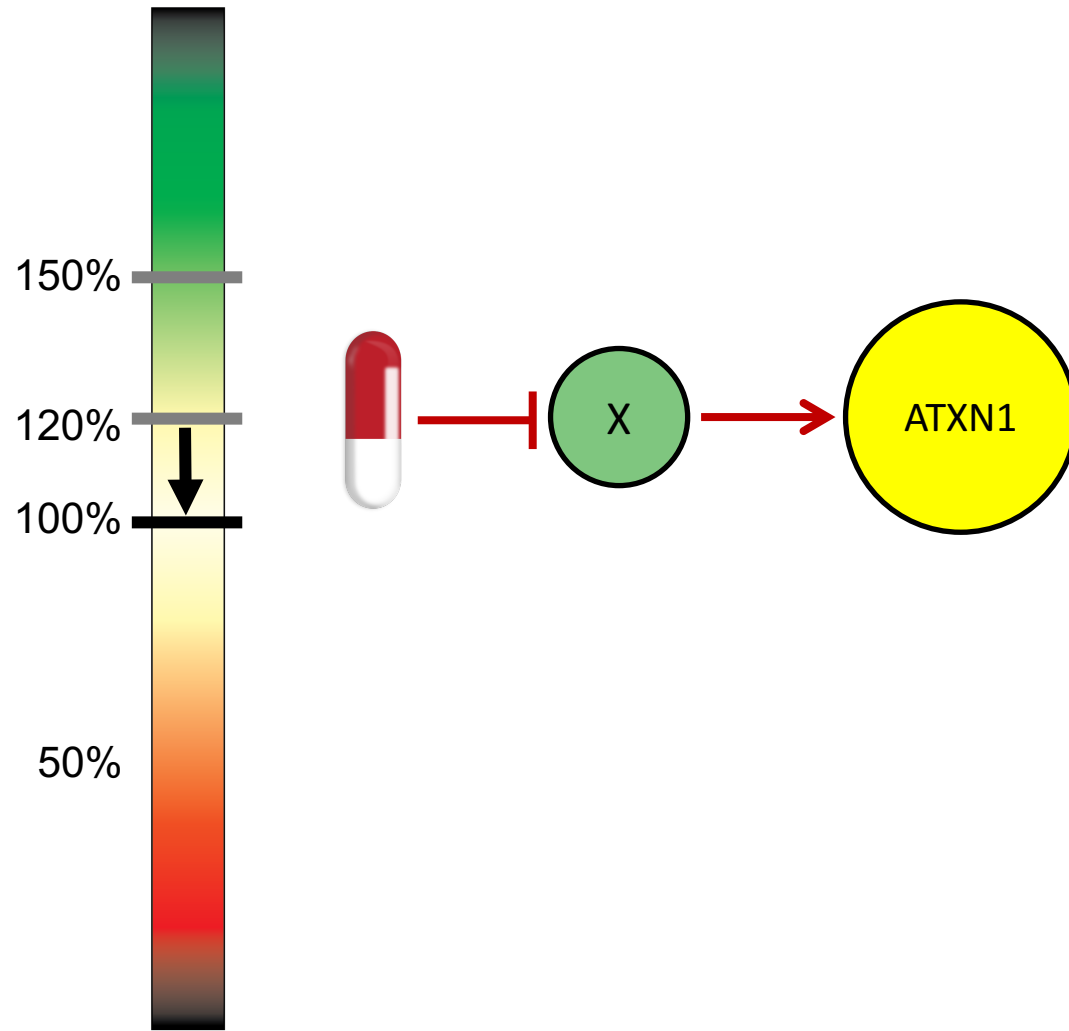
Find genes that regulate ATXN1 levels by systematically inhibiting them using genetics



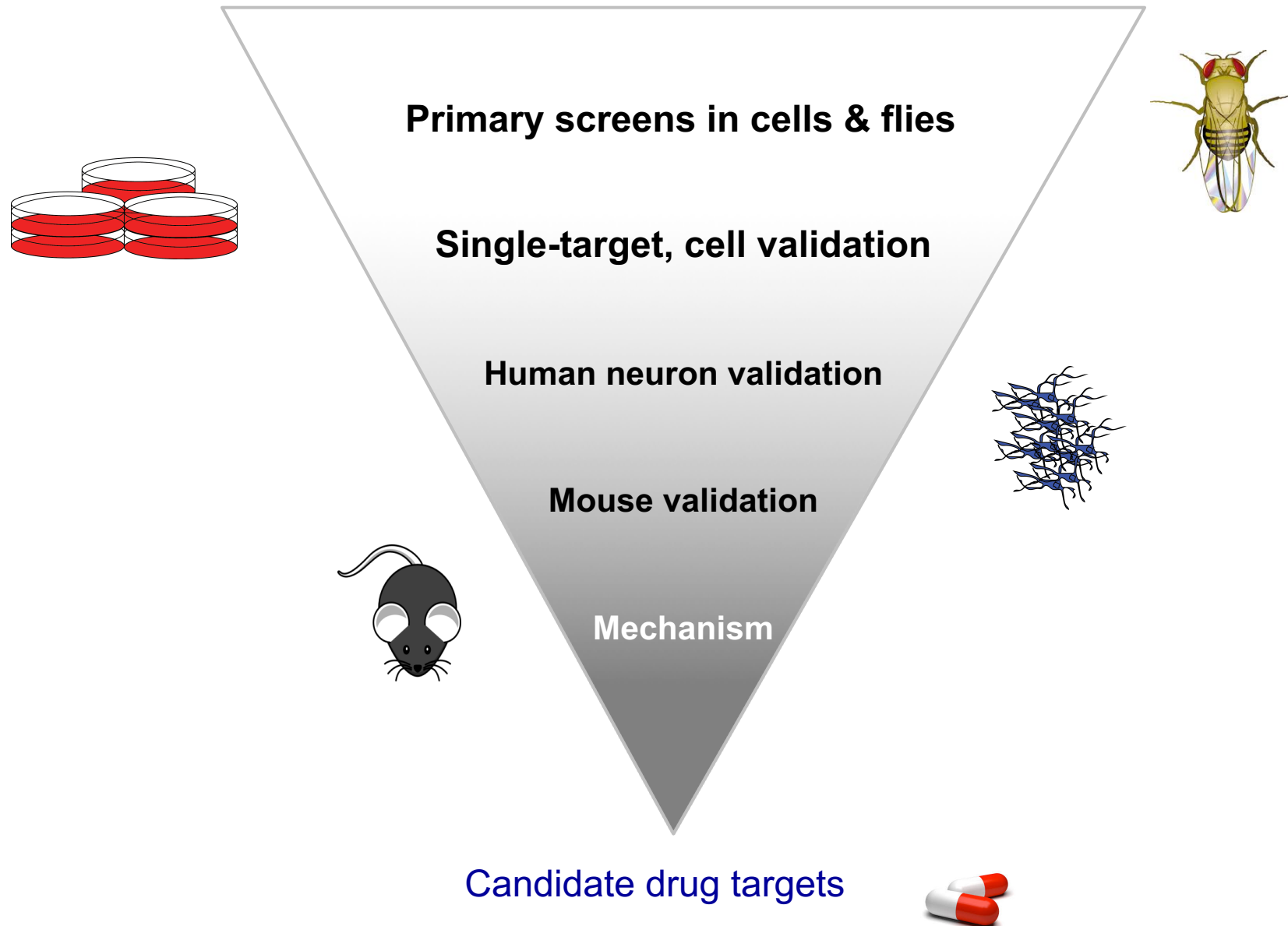
Identify regulators of ATXN1 levels & develop a pharmaceutical to inhibit them



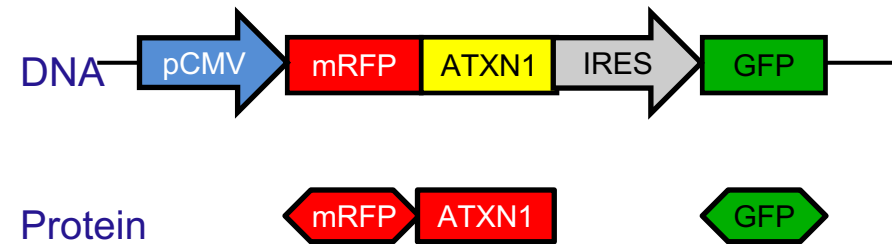
Aim to lower ATXN1 protein levels by ~20%



The strategy to identify ATXN1 regulators

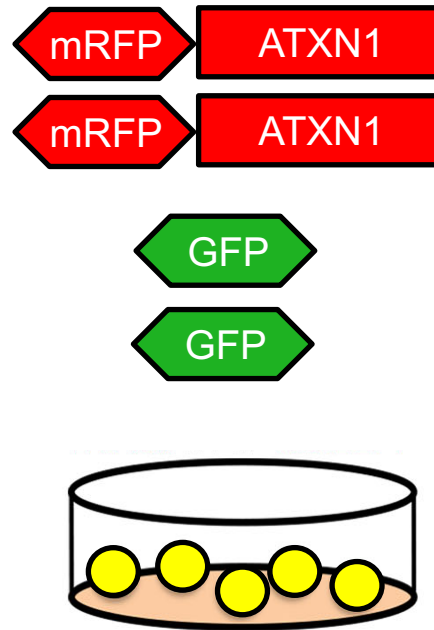


Transgenic cell line that allows monitoring ATXN1 protein levels



Red:green ratio helps identify regulators

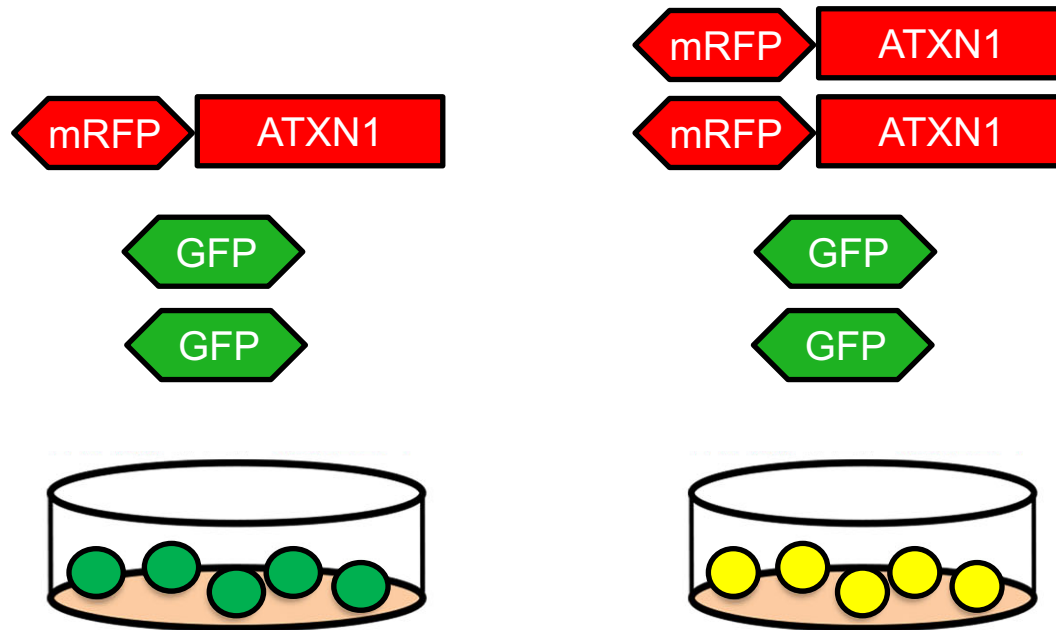
No treatment



Red:green ratio helps identify regulators

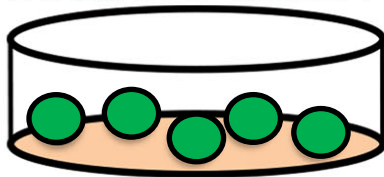
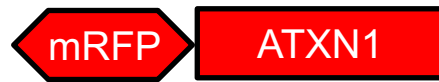
ATXN1 decrease

No treatment

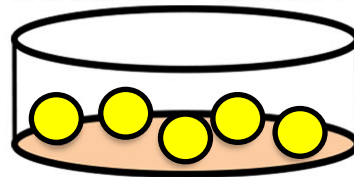
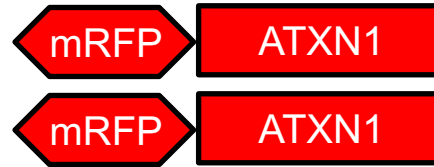


Red:green ratio helps identify regulators

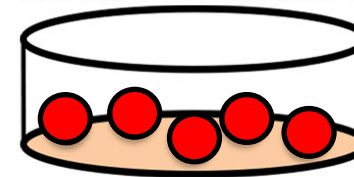
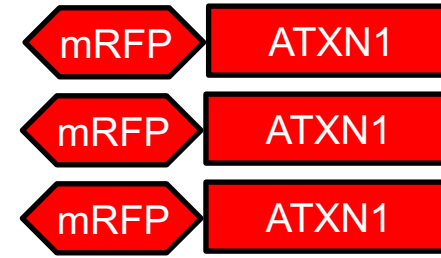
ATXN1 decrease



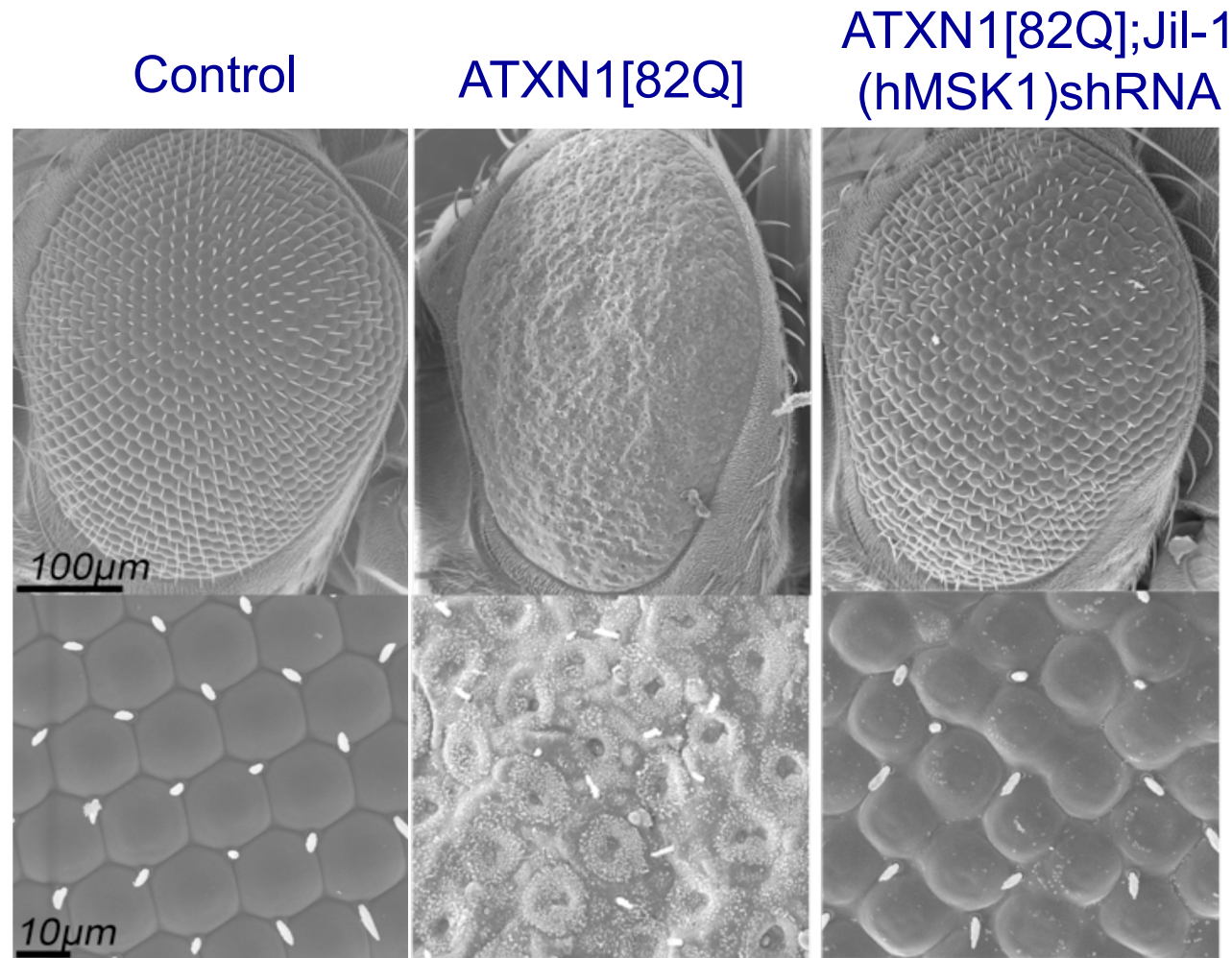
No treatment



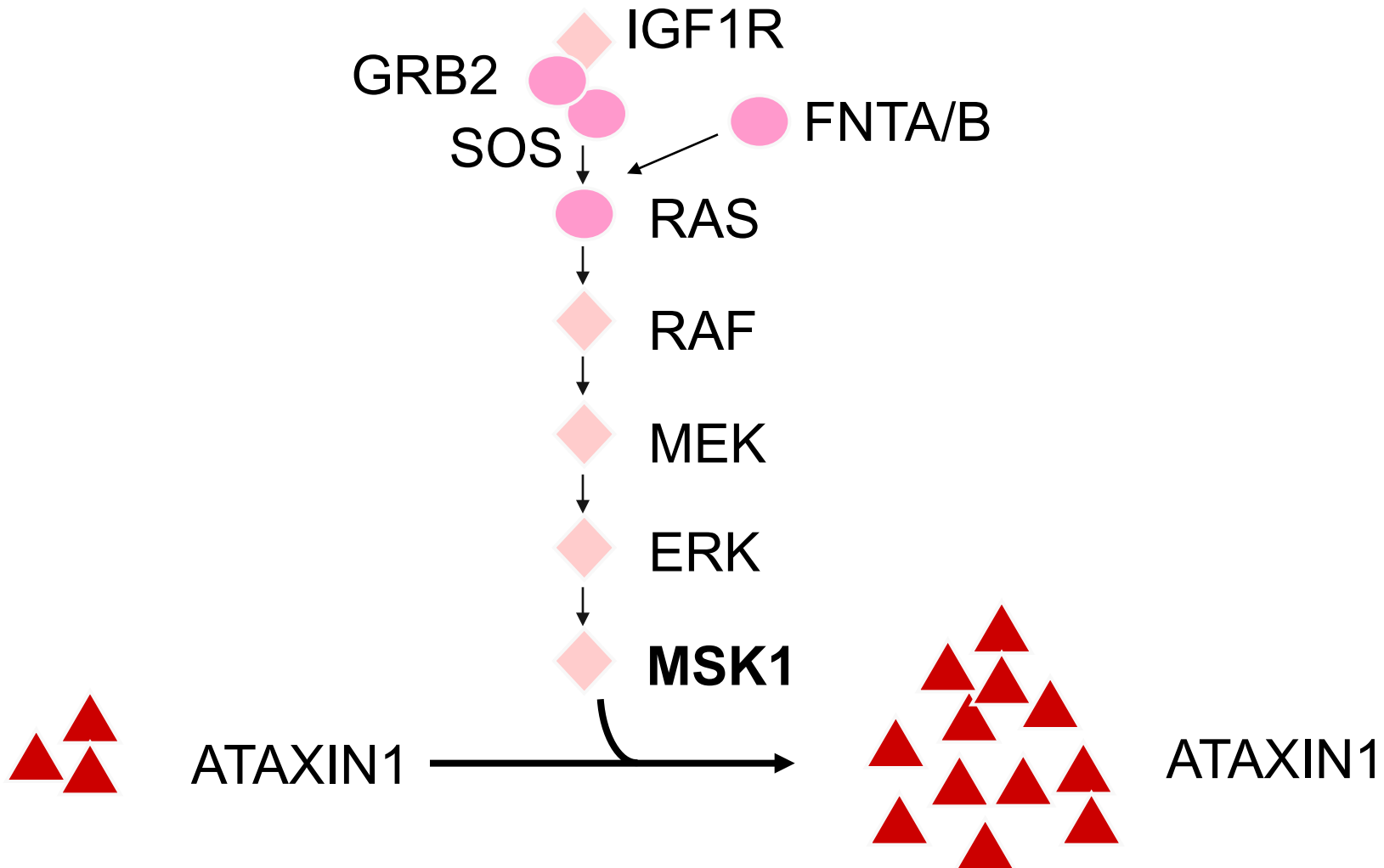
ATXN1 increase



Reduction of the *Drosophila* homolog of MSK1 rescues neurodegeneration



MAPK pathway modulates ATAXIN1 level & toxicity

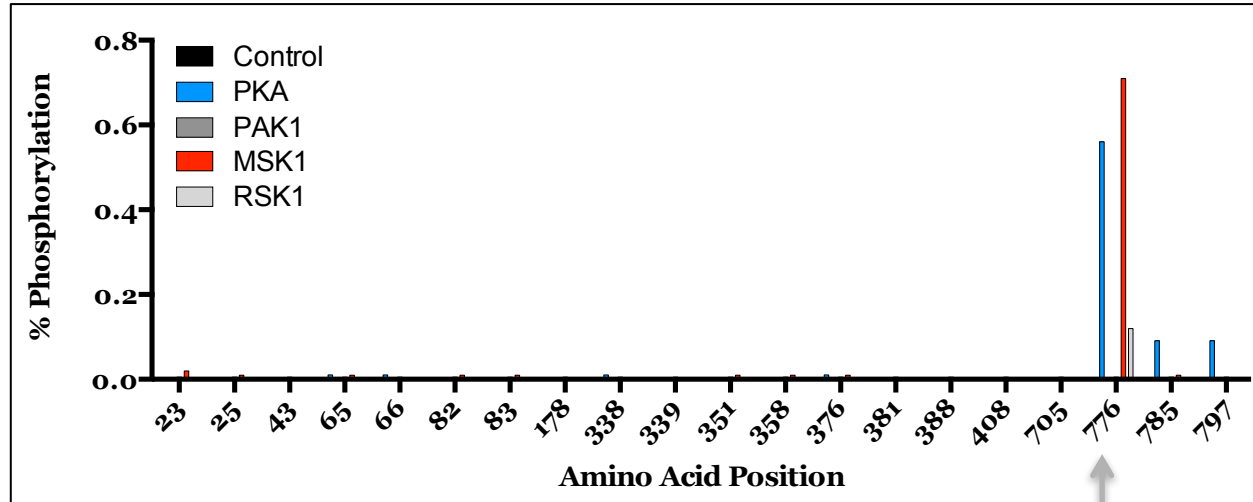


S776 falls within MSK1 consensus site

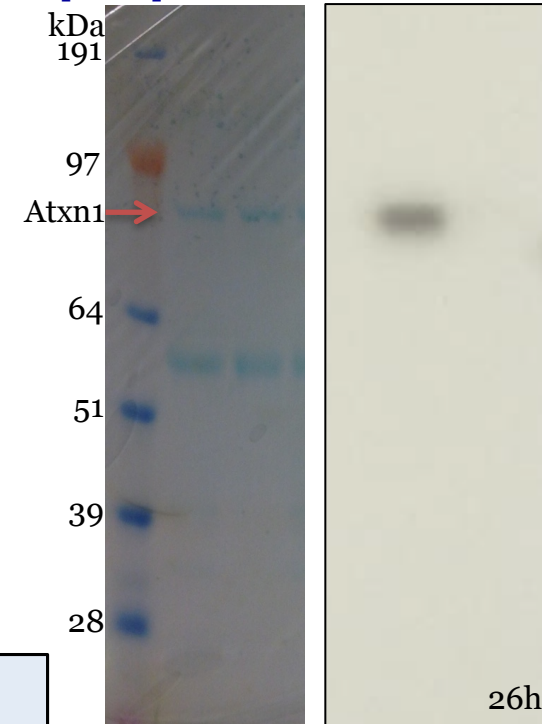
ratATXN1	PSKPTATRKRRW S APETR	754
mouseATXN1	PSKPTATRKRRW S APETR	756
cattleATXN1	PSKPAATRKRRW S APETR	782
humanATXN1	PSKPAATRKRRW S APESR	781
chickenATXN1	PSKPPATRKRRW S APESR	763
zebrafishATXN1	SSKPTG-RKRRW S APEGR	746

MSK1 consensus P site : RxxS

MSK1 phosphorylate Ataxin-1 S776



MSK1: + +
 Atxn-1 [30Q]: + +
 Atxn-1 [30Q] S776A: + +



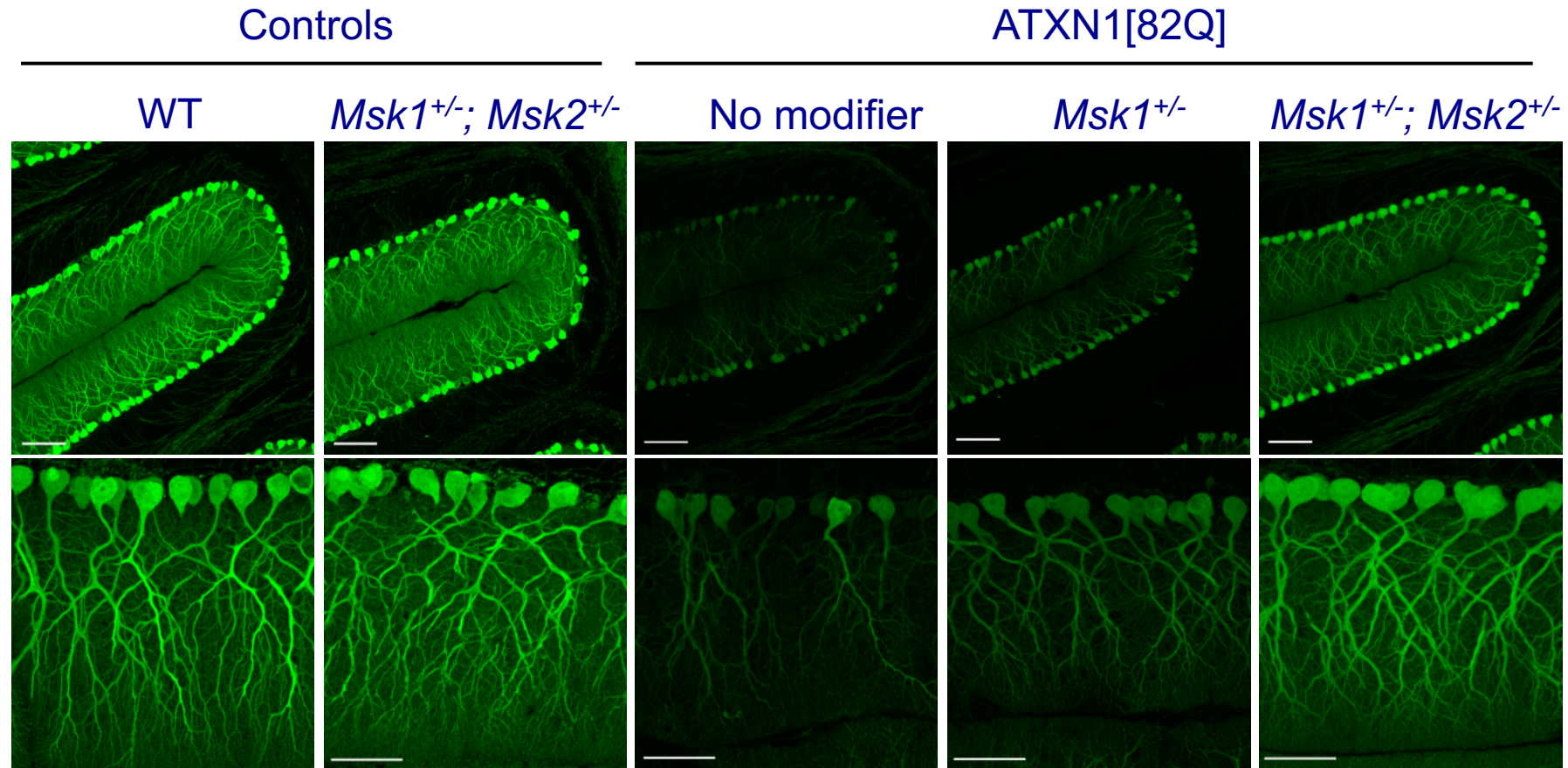
Mass Spec Results on Atxn-1[30Q] substrate

% Phosphorylation Ser776				
Control	PKA	PAK1	MSK1	RSK1
0	56	0	71	12



Carolyn Adamski, PhD

Reducing Msk1/2 rescues neurodegeneration in SCA1 mice

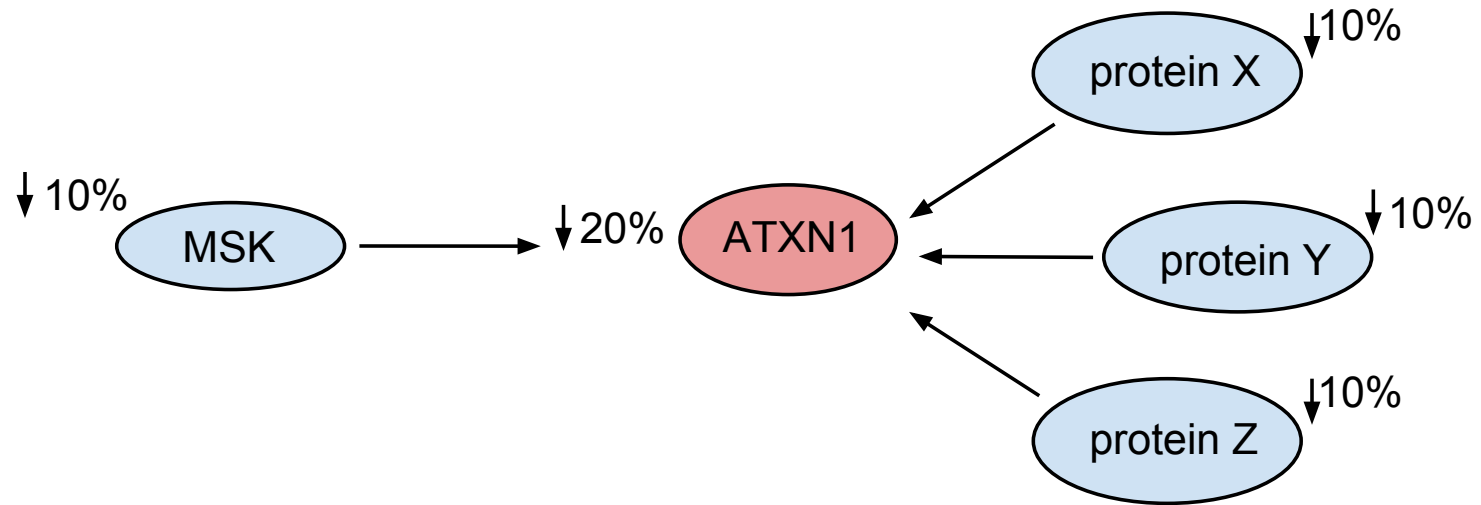


α anti-Calbindin

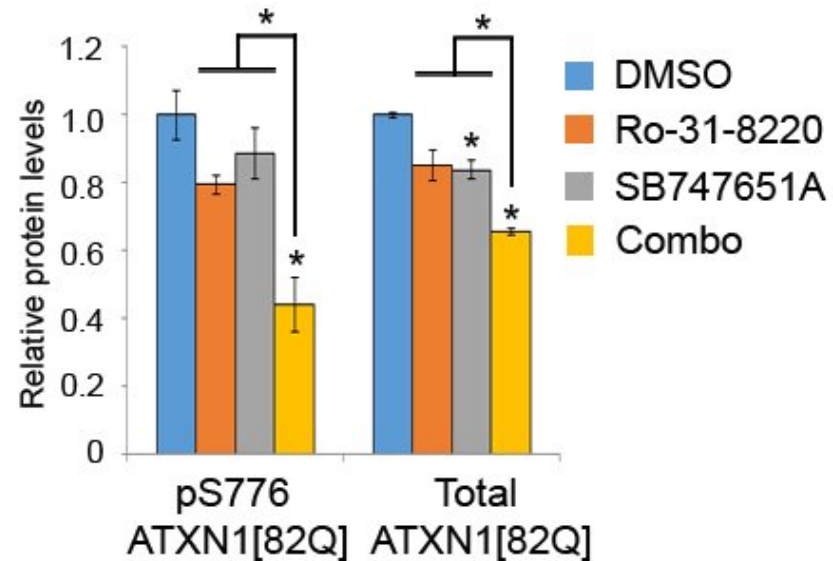
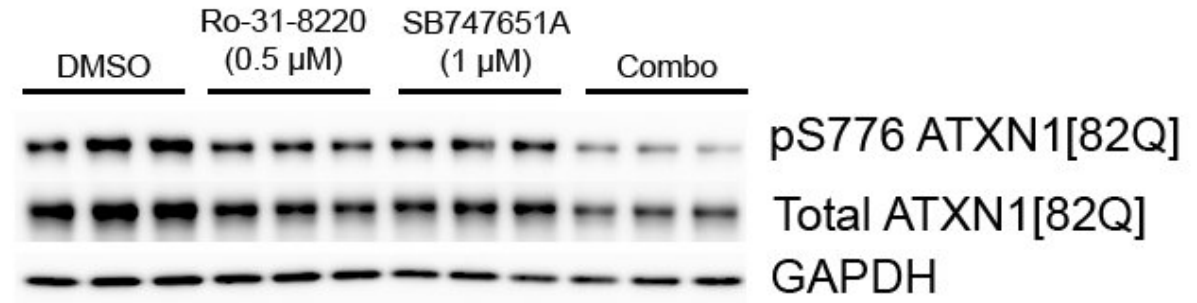
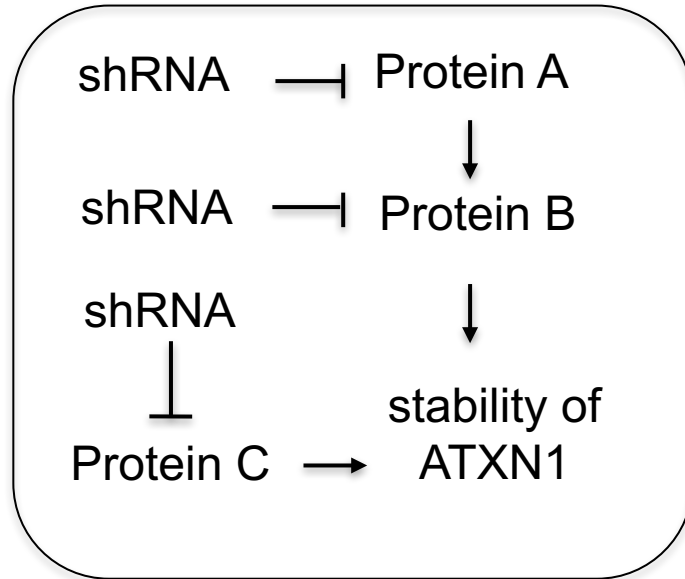
Conclusions from SCA1 studies

- Cross-species kinome screen identified modifiers that regulate ATXN1 stability and toxicity
- MSK1 increases ATXN1 stability by phosphorylating S776 and its reduction rescues degeneration in SCA1 mice
- Pharmacological inhibition of MSK1 decreases ATXN1 levels and warrants preclinical studies

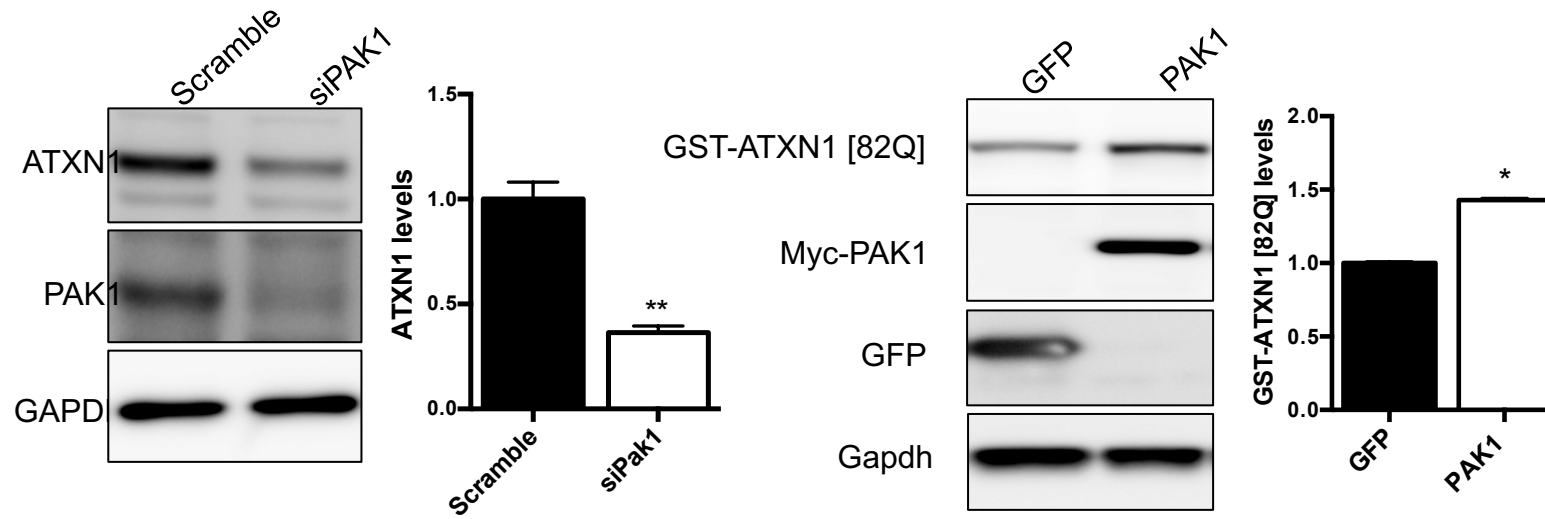
Combination therapy is more effective and likely less toxic for a chronic disease like SCA1



Combination therapy is more effective and likely less toxic for a chronic disease like SCA1



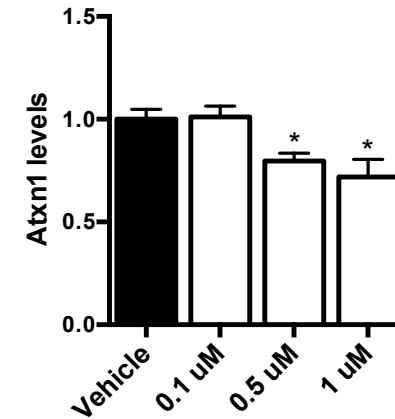
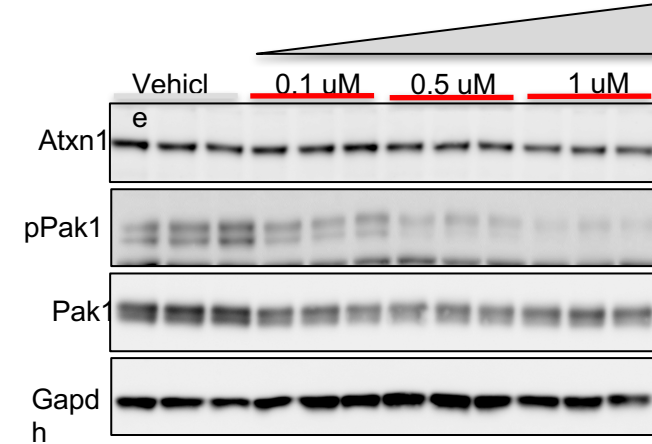
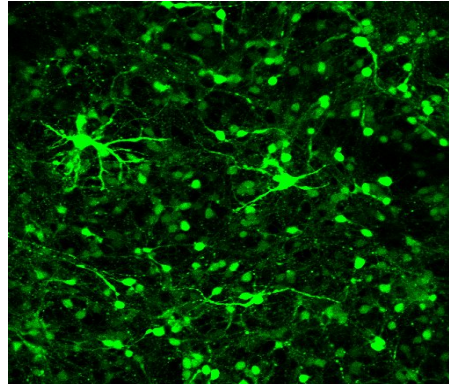
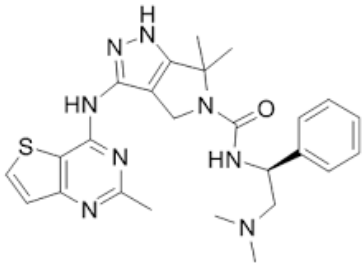
PAK1 genetic perturbation modulates ATXN1 levels



*p<0.05

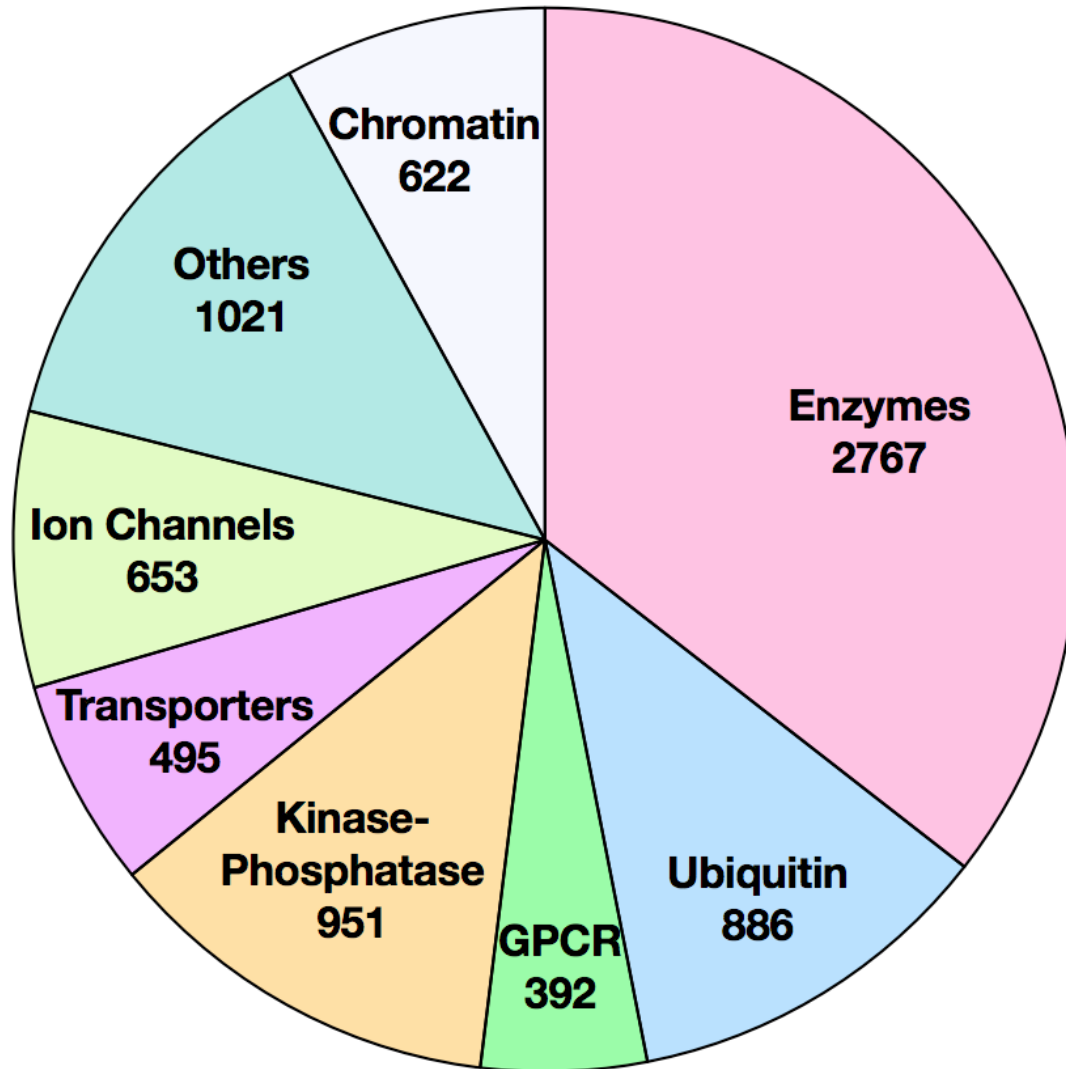
Pharmacological inhibition of PAKs decreases ATXN1 in mouse cerebellar granule neurons

panPAK inhibitor:
PF3758309



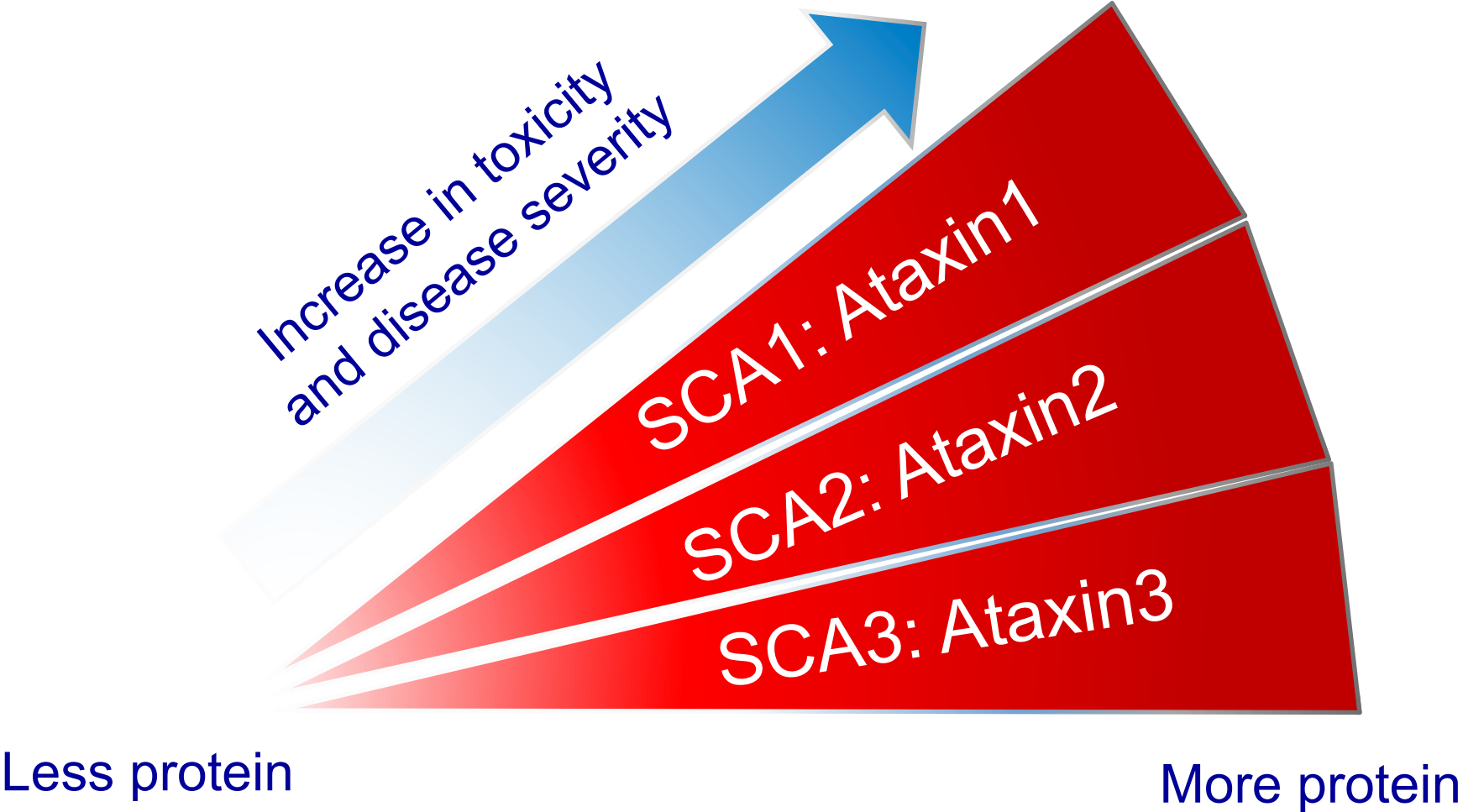
*p<0.05

The druggable genome is composed of eight sub-libraries



7,787 genes x 10
shRNAs/gene=
77,870 shRNAs

Elevated protein levels are a common theme among spinocerebellar ataxias



Conclusions

- Understanding disease mechanism and function of disease-driving proteins are crucial for designing safe interventions
- ASOs and modulators of ATXN1 provide a viable path to develop interventions for SCA1
- Key to any therapeutic intervention is long-term safety as therapy will be required for decades

Acknowledgments

Zoghbi Lab Members

Past

John Fryer, PhD
Paymaan Jafar-Nejad, MD
Jeehye Park, PhD

Current

Carolyn Adamski, PhD
Vitaliy Bondar
Laura Lavery, PhD
Won-Seok Lee
V. Alessandro Gennarino, PhD
Maxime Rousseaux, PhD

Patients & their families

Lab of Harry T. Orr, PhD

Lisa Duvick
Sara Lagalwar
Nissa Mollema
Jill Friedreich

Lab of Juan Botas, PhD

Ismael Alramahi, PhD

Zhandong Liu, PhD

Hyun-Hwan Jeong, PhD

Thomas (Trey) Westbrook, PhD

Steve Elledge PhD

HHMI, HHMI-CIA, NINDS, IDDRRC, and Belfer NDC



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