

Features of ATXN2 Expression Control Daniel R. Scoles, Lance T. Pflieger, Steven T. Hansen, Khanh K. Thai, Stefan-M. Pulst Department of Neurology, University of Utah, Salt Lake City

The presence of the CAG repeat supports *ATXN2-luc* expression, but if it is highly expanded protein abundance is reduced

Introduction

SCA2 is an autosomal dominant cerebellar ataxia characterized by progressive degeneration of the cerebellum and brain stem. SCA2 is caused by polyglutamine expansion in the SCA2 protein ataxin-2, causing gain-of-function. We hypothesize that reduction of *ATXN2* expression will be therapeutic for SCA2, based on reversible phenotypes in polyQ models and lack of neurodegeneration in SCA2 knockout mice.

Objectives

To characterize features of *ATXN2* that might be exploited therapeutically for treating SCA2.

ATXN2-luc reporter construct

ATXN2 Promo	AT ter 5'	WN2 UTR	Luciferase	<i>ATXN2</i> 3'-UTR	ATXN2 Downstre	eam
Ť	1	1			↑	Ť
-1062	+1	ATO	6	3' -UT	R end at	3' -UTR end at
	(+643)		3)	598 bp		1012 bp
		•	-	after s	stop	after stop

Similar to endogenous *Atxn2*, *ATXN2luc* is expressed in cerebellum (CB) and olfactory bulb (OB)



Wildtype Transgenic Wildtype Transgenic Wildtype Transgenic Images display photons/sec/cm²/sr determined by *in vivo* imaging on a Xenogen IVIS 100.



Not shown is nearly identical data for full-length ATXN2 with varying CAG repeats.

Deletion of an ETS element in the ATXN2 promoter inhibited ATXN2-luc expression



The ATXN2 ETS site has 100% match with the ETS consensus and when mutated in ATXN2-luc expression was reduced

	+87 <u>ETS</u> +109		CMINO ATANO COT ON
ATXN2	3'-ACCCCTCCGACTTCCGGTAAAGA-3' 3'-TGGGGAGGC <u>TGAAGGCC</u> ATTTCT-5'	Luciforaça	
mCGA	5'-ACCCCTCCGACTCGAGGTAAAGA-3' 3'-TGGGGAGGCTGA <u>GCT</u> CCATTTCT-5'	Lucherase	
D14	5'-ACCCCTCGA-3' 3'-TGGGGAGCT-5'	Actin	
			Western Blot

5 bp deletions in *ATXN2* reveal lowest expression with ETS core elimination

TTCTGCTTCCGTCTGACCCCTCCGACTTCCGGTAAAGAGTCCCTATCCGCACCTCCGC TTCTGCTTCCGTCTGACCCCT______GTAAAGAGTCCCTATCCGCACCTCCGC TTCTGCTTCCGTCTGACCCCTCCGACTTCCGGTAAAGAGTCCCTATCCGCACCTCCGC TTCTGCTTCCGTCTGACCCCT_____GTAAAGAGTCCCTATCCGCACCTCCGC TTCTGCTTCCGTCTGACCCCTCCGACTTCCG TTCTGCTTCCGTCTGACCCCTCCGACTTCCGGTAAAGAGTCCCTATCCGCACCTCCGC TTCTGCTTCCGTCTGACCCCTCCGACTTCCGGTAAAGAGTCCCTATCCGCACCTCCGC TTCTGCTTCCGTCTGACCCCTCCGACTTCCGGTAAA_____CCTATCCGCACCTCCGC TTCTGCTTCCGTCTGACCCCTCCGACTTCCGGTAAAGAGTC_____CCGCACCTCCGC

* 58 bp *ATXN2* promoter fragments placed between a 75 bp minimal prolactin promoter and luciferase



ETS1 binds the ETS element in ATXN2 EMSA Supershift ChIP-PC



SH-SY5Y

ETS1 overexpression increased ataxin-2



SH-SY5Y

ETS1 underexpression decreased ataxin-2



Immunofluorescent labeling showed ETS1 overexpression increased endogenous ataxin-2 expression in SH-SY5Y cells



ETS1 transfected cells (red) make more ataxin-2 (green)



Conclusions

- ATXN2-luc expression mimicked predictions for endogenous Atxn2 (seen in Allen Brain Atlas), suggesting this ATXN2 fragment possesses promoter elements required for tissue-specific expression.
- Ets1 in the Allen Brain Atlas shows high expression in Purkinje cells, suggesting ETS1 may support ATXN2 expression in this cell type.



Ets1 in situ hybridization. Reproduced with permission of the Allen Mouse Brain Atlas [Internet]. Seattle (WA): Allen Institute for Brain Science. ©2009. Available from: http://mouse.brainmap.org.

- The presence of the CAG repeat increased protein abundance without increasing transcription while very long CAG tracts in *ATXN2* reduced protein abundance. We hypothesize that support of *ATXN2* expression may involve RNA binding proteins binding the CAG repeat or increased ataxin 2 stability, and reduced protein abundance when CAG is expanded may reflect RNA or protein aggregation.
- The ETS element supports the majority of ATXN2 expression, therefore ETS1 may represent a therapeutic target for SCA2.
- ATXN2 expression might be targeted by decoy oligonucleotides containing the ATXN2 ETS element and short flanking sequences.

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