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Development and validation of SCACOMS, a composite scale for assessing disease progression and treatment effects in spinocerebellar ataxia

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CONCLUSIONS

- Using two natural history datasets, Spinocerebellar Ataxia Composite Scale (SCACOMS) was successfully developed and validated in subjects with spinocerebellar ataxia (SCA).
- SCACOMS detects ataxia-related functional decline in patients with SCA, and can support development of clinical trials with increased power to detect small but meaningful clinical impacts of disease modifying treatments.
- Analysis of troriluzole's treatment effect (BHV4157-206) using SCACOMS demonstrates a significant and meaningful reduction in disease progression, favoring troriluzole

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Researchers can submit requests for the CRC-SCA data at https://www.ataxia.org/crc-sca/academicresearch/ and EUROSCA data at https://www.eurosca.org.



OBJECTIVE

- genotype.

Statistical Analysis

- scale(s).

INTRODUCTION

SCA is a rare inherited disorder characterized by progressive ataxia affecting limb coordination, balance, and speech.^{1,2}

While there have been recent developments of disease modifying treatments (DMTs) targeting SCA pathology¹, many have yet to be tested in SCA patients and none have been approved for this population.

Due to heterogeneity of symptoms in SCA, clinical trial development can be challenging. Existing scales may not capture changes in response to DMTs with sufficient sensitivity over 1-year (a typical timeframe for interventional studies).³

A more sensitive measure of clinical decline derived from existing scales would allow for more robust study design and would enable DMTs to reach SCA patients more efficiently.

> To develop a composite scale using natural history datasets to assess disease progression and treatment effects more accurately in patients with SCA.

METHODS

Study Participants and Data

The primary analysis was conducted using natural history data from the Clinical Research Consortium for Spinocerebellar Ataxias (CRC-SCA; NCT01060371) and from the European Integrated Project on Spinocerebellar Ataxias (EUROSCA; NCT02440763), using patients with SCA3 genotype. Secondary analysis (not reported) repeated the analysis in all SCA

Candidate items for SCACOMS included: the Clinical Global Impression of change [CGI], 4 items from the Modified-functional Scale for the Assessment and Rating of Ataxia [f-SARA]) and Friedreich's ataxia rating scale – functional stage (FARS-FUNC) [CRC-SCA only]

Sensitivity of individual items to progression in the natural history datasets was assessed using partial least squares (PLS) regression, with responsive items objectively selected and summed to create SCACOMS; item weights reflected sensitivity to decline.

Scales were assessed for ability to detect disease progression using mean-to-standard deviation ratios (MSDR).

SCACOMS were applied to data from BHV4157-206, a 48-week study evaluating troriluzole in SCA subjects (NCT03701399) to measure treatment effects (between group differences in mean change from baseline, least squared mean [LSM]).

RESULTS

Baseline demographics of the natural history datasets are provided in Table 1.

Using PLS regression, the CGI, FARS-FUNC, and f-SARA gait, speech, and stance items were selected for inclusion (Table 2). Relative weightings of scale items are visualized in Figure 1.

Improved sensitivity (e.g., increased) MSDR) was observed for SCACOMS scales as compared to the original

Consistent results were found in the analysis of EUROSCA. With the absence of FARS-FUNC, the other items were distributed similarly, with CGI accounting for $\sim 1/2$ of the total score.

 Cross-validation between the natural. history sets yielded similar MSDRs, further supporting their validity.

Figure 1 Relative weightings (%) of scale items in SCACOMS for SCA3 patients, (A) **CRC-SCA and (B) EUROSCA models**



RESULTS

Table 1 Descriptive data for the CRC-SCA analytic dataset and EUROSCA analytic dataset used to derive SCACOMS

	CRC-SCA		EUROSCA			
Baseline characteristic	ALL SCA	SCA3	ALL SCA	SCA3		
	(n = 214)*	(n = 77)*	(n = 423)*	(n = 106)*		
Age, mean (SD) years	52.3 (13.4)	51.3 (12.5)	47.3 (12.7)	50.3 (11.5)		
Sex, % female	54.2	57.1	53.9	52.8		
Genotype, %						
SCA1	18.2	0.0	23.9	0.0		
SCA2	20.1	0.0	31.9	0.0		
SCA3	35.5	98.7	25.1	100.0		
SCA6	22.4	0.0	19.1	0.0		
SCA8	2.3	0.0	0.0	0.0		
SCA10	0.9	0.0	0.0	0.0		
SCA3 and SCA8	0.5	1.3	0.0	0.0		
f-SARA gait score, mean (SD)	1.5 (1.1)	1.4 (1.1)	1.6 (1.2)	1.9 (1.4)		
f-SARA stance score, mean (SD)	1.3 (1.1)	1.3 (1.1)	1.5 (1.3)	1.4 (1.3)		
f-SARA speech score, mean (SD)	0.7 (0.9)	0.4 (0.7)	1.1 (1.0)	0.9 (1.1)		
f-SARA sitting score, mean (SD)	0.5 (0.8)	0.5 (0.7)	0.8 (1.0)	0.9 (1.1)		
FARS functional stage, mean (SD)	2.8 (1.0)	2.9 (0.9)	NA	NA		
CGI, mean (SD)	NA	NA	NA	NA		
 Subjects in the analytic dataset were required to have baseline and either 12- or 24-months values on the measures of interest CGI, Clinical Global Impression - Global Improvement Scale; CRC-SCA, Clinical Research Consortium for the Study of Cerebellar Ataxia; EUROSCA, European Integrated Project on Spinocerebellar Ataxias; FARS, Friedreich ataxia rating scale; f-SARA, Modified functional Scale for the Assessment and Rating of Ataxia; NA, not available; SCA, spinocerebellar ataxia; SD, standard deviation 						

Table 2 VIP scores and PLS coefficients for SCA3 patients, CRC-SCA and **EUROSCA** models

tem SCA3 patients f-SARA gait FARS function f-SARA stand f-SARA spee f-SARA sittir CGI SCACOMS SCA3 patients f-SARA gait FARS function f-SARA stan f-SARA spee f-SARA sittir CGI SCACOMS MSDF

CGI, Clinical Global Impression - Global Improvement Scale; CRC-SCA, Clinical Research Consortium for the Study of Cerebellar Ataxia; EUROSCA, European Integrated Project on Spinocerebellar Ataxias; FARS, Friedreich ataxia rating scale; f-SARA, Modified functional Scale for the Assessment and Rating of Ataxia; MSDR, mean to standard deviation ratio; PLS, partial least squares; SCA, spinocerebellar ataxia; VIP, Variable Importance of Projection *The MSDR of changes in total f-SARA score was 0.4826.

	Item MSDR*	VIP	PLS weight	% contribution			
s, CRC-SCA dataset (N=77)							
score	0.4604	0.6417	4.5390	10.45			
onal stage	0.1688	0.4741	10.8056	24.87			
ce score	0.2683	1.0522	9.9743	22.95			
ech score	0.2614	0.4952	4.2284	9.73			
ng score	0.2538	Removed	0.0000				
	0.9125	1.7353	13.9060	32.00			
MSDR	0.9171						
s, EUROSCA dataset (N=106)							
score	0.2952	0.6718	2.5315	4.83			
onal stage							
ce score	0.5629	1.0763	13.5141	25.76			
ech score	0.2320	0.6486	10.7757	20.54			
ng score	0.3056	0.5132	2.0793	3.96			
	1.0742	1.645	23.562	44.91			
MSDR	1.1157						

- subjects, favoring troriluzole (Figure 2).
- subjects, Cohen's d of 0.59 (Figure 2).

Figure 2 SCACOMS outcomes in BHV4157-206 SCA3 subjects using the (A) CRC-SCA model and (B) EUROSCA model





DISCUSSION

- weeks.
- patients with SCA.

Examination of SCACOMS treatment effects at 48-weeks in BHV4157-206 SCA3 subjects (n=85) showed a statistically significant treatment effect in

The percentage of progression avoided was 80% for the CRC-SCA derived model, Cohen's d of 0.56; and 88% for the model derived from EUROSCA

% of disease progression avoided with troriluzole: 80% Treatment Placebo Troriluzol P value: 0.0030 Time (Weeks) % of disease progression avoided with troriluzole: 88% Treatment 🔸 Placebo ----- Troriluzole P value: 0.0041 Time (Weeks)

The overall MSDRs indicated that SCACOMS endpoints were highly sensitive to detecting progression and showed an improvement in responsiveness when compared to the original component scales.

Application of SCACOMS to analysis of data from the BHV4157-206 study, SCA3 genotype subpopulation, yielded compelling and consistent treatment effects across CRC-SCA and EUROSCA composite score models at 48-

This study highlights the potential usefulness of SCACOMS for clinical and research purposes. The availability of a scale that detects small but meaningful changes in disease progression and treatment effects will lead to more efficient trial design and support the development of novel DMTs for