

Supplemental Table 1. Frequency of full repeat C9orf72 expansions in SCA patients and normal American population.

|  | Normal American population | SCA1 | SCA2 | SCA3 | SCA6 |
| :--- | :--- | :--- | :--- | :--- | :--- |
| C9orf72 expansions $\geq$ 31, $\mathbf{n}$ | 1659 | 48 | 56 | 108 | 58 |
| C9orf72 expansions $\leq$ 30, $\mathbf{n}$ | 0 | 3 | 2 | 1 | 1 |
| Total, $\mathbf{n}$ | 1659 | 51 | 58 | 109 | 59 |
| $\boldsymbol{p}$ value $^{\mathrm{a}}$ | - | 0.000 | $0.001^{\mathrm{a}}$ | $0.062^{\mathrm{a}}$ | $0.034^{\mathrm{a}}$ |

American normal controls are from the reference 5.
${ }^{a}$ Fisher exact test


Supplemental Table 2. Baseline features of SCA participants with normal ( $\leq 7$ ) and intermediate (8-30) C9orf72 repeat expansions

|  | $\begin{aligned} & \text { SCA } 1 \\ & n=48 \end{aligned}$ |  | $p$-value ${ }^{\text {d }}$ | $\begin{aligned} & \text { SCA } 2 \\ & \mathrm{n}=56 \end{aligned}$ |  | $p$-value ${ }^{\text {d }}$ | $\begin{aligned} & \text { SCA } 3 \\ & n=108 \end{aligned}$ |  | $p$-value ${ }^{\text {d }}$ | $\begin{aligned} & \text { SCA } 6 \\ & \mathrm{n}=58 \\ & \hline \end{aligned}$ |  | $p$-value ${ }^{\text {d }}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Normal | Intermediate |  | Normal | Intermediate |  | Normal | Intermediate |  | Normal | Intermediate |  |
| Sample size, n (\%) | 28 (58.3) | 20(41.7) |  | 35 (62.5) | 21 (37.5) |  | 68 (63.0) | 40 (37.0) |  | 31 (53.4) | 27 (46.6) |  |
| $\begin{aligned} & 0 \text { Age at first } \\ & 1 \text { visit, } y \end{aligned}$ | $49.8 \pm 14.4$ | $50.3 \pm 11.5$ | $0.888^{\text {b }}$ | $51.0 \pm 14.0$ | $47.3 \pm 12.5$ | $0.329^{\text {b }}$ | $53.0 \pm 12.1$ | $48.6 \pm 12.3$ | $0.075^{\text {b }}$ | $68.2 \pm 10.0$ | $62.1 \pm 12.2$ | $0.038^{\text {b }}$ |
| $\begin{aligned} & 3 \\ & 4 \\ & 5: \text { Gender, M } \end{aligned}$ | 17: 11 | 10:10 | $0.461{ }^{\text {a }}$ | 17:18 | 13:8 | $0.333^{\text {a }}$ | 30:38 | 21:19 | $0.399^{\text {a }}$ | 16:15 | 14:13 | $0.986^{\text {a }}$ |
| 6 Age of 7 onset, $y$ 8 9 0 | $39.9 \pm 12.9$ | $39.7 \pm 9.5$ | $0.943^{\text {b }}$ | $36.6 \pm 13.5$ | $35.0 \pm 10.9$ | $0.646^{\text {b }}$ | $\begin{gathered} 39.9 \pm 12.5 \\ \text { Median }= \\ 40.0 \end{gathered}$ | $\begin{gathered} 37.2 \pm 11.6 \\ \text { Median }= \\ 38.5 \end{gathered}$ | $0.339^{\text {c }}$ | $52.7 \pm 10.3$ | $49.7 \pm 11.0$ | $0.290^{\text {b }}$ |
| $\begin{aligned} & 1 \text { Disease } \\ & 2 \text { duration, } y \\ & 3 \\ & 4 \end{aligned}$ | $9.9 \pm 8.3$ <br> Median $=$ $8.0$ | $\begin{gathered} 10.6 \pm 6.3 \\ \text { Median }=7.5 \end{gathered}$ | $0.508^{\text {c }}$ | $15.0 \pm 9.1$ | $12.3 \pm 6.0$ | $0.245^{\text {b }}$ | $\begin{gathered} 13.4 \pm 8.0 \\ \text { Median }= \\ 12.0 \end{gathered}$ | $\begin{gathered} 11.6 \pm 7.9 \\ \text { Median = } 11.0 \end{gathered}$ | $0.204^{\text {c }}$ | $\begin{gathered} 15.5 \pm 11.8 \\ \text { Median }= \\ 12.0 \end{gathered}$ | $\begin{gathered} 12.3 \pm 9.7 \\ \text { Median }=9.0 \end{gathered}$ | $0.285^{\text {c }}$ |
| $\begin{aligned} & 5 \text { Expanded } \\ & 6 \text { CAG } \\ & 7 \text { repeats } \\ & 9 \end{aligned}$ | $46.4 \pm 4.1$ <br> Median = <br> 47.0 | $\begin{gathered} 46.6 \pm 5.0 \\ \text { Median }= \\ 46.5 \end{gathered}$ | $0.966^{\text {c }}$ | $\begin{gathered} 40.1 \pm 3.1 \\ \text { Median }= \\ 39.0 \end{gathered}$ | $\begin{gathered} 40.9 \pm 3.7 \\ \text { Median }= \\ 40.0 \end{gathered}$ | $0.298{ }^{\text {c }}$ | $\begin{gathered} 70.4 \pm 3.9 \\ \text { Median }= \\ 71.0 \end{gathered}$ | $\begin{gathered} 72.0 \pm 4.3 \\ \text { Median }= \\ 72.0 \end{gathered}$ | $0.111^{\text {c }}$ | $\begin{gathered} 22.4 \pm 1.1 \\ \text { Median }= \\ 22.0 \end{gathered}$ | $\begin{gathered} 22.3 \pm 0.7 \\ \text { Median }= \\ 22.0 \end{gathered}$ | $0.656{ }^{\text {c }}$ |
| $\begin{aligned} & 0 \\ & 1 \text { SARA } \\ & 2 \text { scores } \end{aligned}$ | $\begin{gathered} 12.6 \pm 8.4 \\ \text { Median }= \\ 12.25 \end{gathered}$ | $\begin{gathered} 15.9 \pm 7.7 \\ \text { Median }= \\ 14.25 \end{gathered}$ | $0.177^{\text {c }}$ | $17.5 \pm 6.9$ | $17.7 \pm 7.3$ | $0.939^{\text {b }}$ | $\begin{gathered} 15.3 \pm 8.2 \\ \text { Median }= \\ 14.5 \end{gathered}$ | $\begin{gathered} 15.4 \pm 10.1 \\ \text { Median }= \\ 13.25 \end{gathered}$ | $0.929^{\text {c }}$ | $16.1 \pm 7.1$ | $14.3 \pm 7.8$ | $0.343^{\text {b }}$ |
| $\begin{aligned} & 4 \\ & 5 \text { PHQ-9 } \\ & 6 \text { scores } \\ & 7 \end{aligned}$ | $\begin{gathered} 5.5 \pm 6.3 \\ \text { Median }= \\ 3.0 \end{gathered}$ | $\begin{gathered} 8.0 \pm 7.1 \\ \text { Median }=7.5 \end{gathered}$ | $0.178{ }^{\text {c }}$ | $\begin{gathered} 5.2 \pm 5.0 \\ \text { Median }=4.0 \end{gathered}$ | $\begin{gathered} 5.4 \pm 5.7 \\ \text { Median }=3.0 \end{gathered}$ | $0.931{ }^{\text {c }}$ | $\begin{gathered} 8.0 \pm 6.1 \\ \text { Median }= \\ 6.0 \end{gathered}$ | $\begin{gathered} 5.7 \pm 4.1 \\ \text { Median }=5.5 \end{gathered}$ | $0.101^{\text {c }}$ | $\begin{gathered} 6.7 \pm 6.0 \\ \text { Median }= \\ 6.0 \end{gathered}$ | $\begin{gathered} 7.7 \pm 6.4 \\ \text { Median }=6.0 \end{gathered}$ | $0.594^{\text {c }}$ |

[^0]${ }^{\text {c }} 2$ independent samples Mann-Whitney $U$ test
${ }^{d}$ A Bonferroni correction was made to adjust for multiple comparisons, $\mathrm{p}<0.007$ were considered significant ( 7 tests in each subtypes)


Supplemental Table 3. Longitudinal SARA and PHQ-9 scores of normal and intermediate C9orf72 repeat expansion in GEE models

|  | Regression coefficients of SARA score |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Age of first visit (yrs) | Sex | Expanded CAG repeats | Allele type of C9orf72 | Visit time | Allele type of C9orf72 <br> $\times$ Visit time |
|  | SCA1 | 0.62**** | 4.71** | 1.42**** | 2.66 | 0.95* | -0.04 |
| $\cdots$ | SCA2 | $0.42^{* * * *}$ | -1.40 | 1.87**** | 0.64 | 0.40 | -0.84 |
|  | SCA3 | 0.67 **** | -0.82 | $1.88{ }^{* * * *}$ | 0.14 | 0.37* | -0.02 |
|  | SCA6 | 0.11 | -1.68 | 0.71 | 2.12 | 0.23 | 0.96 |
|  | Regression coefficients of PHQ-9 score |  |  |  |  |  |  |
|  |  | Age of first visit (yrs) | Sex | Expanded CAG repeats | Allele type of C9orf72 | Visit time | Allele type of C9orf72 <br> $\times$ Visit time |
|  | SCA1 | 0.08 | -0.37 | 0.25 | 2.16 | 0.32 | -1.90*** |
|  | SCA2 | 0.01 | -1.35 | -0.12 | 1.54 | 1.00*** | -0.63 |
|  | SCA3 | 0.26*** | 2.41 | 0.66* | -5.49**** | -0.38 | 3.48**** |
|  | SCA6 | -0.12* | -2.65 | -1.23* | -1.01 | 0.30 | -1.72* |

Abbreviations: GEE = generalized estimating equation, PHQ-9 = The 9-item Patient Health Questionnaire, SARA = Scale for Assessment and Rating of Ataxia, SCA = Spinocerebellar Ataxia.
*p <0.05, **p < 0.01, ***p $<0.005,{ }^{* * * *} p<0.001$
Sex: women $=0$, men $=1$
Allele type of C9orf72: normal allele $=0$, intermediate allele $=1$
Visit time: time order every 6 month during the two-year follow up


# C9orf72 Repeat Expansions as Genetic Modifiers for Depression in Spinocerebellar Ataxias 

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The genetic interactions between pathological repeat expansions have been of major interests in neurodegenerative disorders. Recently, pathogenic C9orf72 repeat expansions, a main genetic cause of amyotrophic lateral sclerosis (ALS) and frontotemporal dementia and pathogenic ATXN2 repeat expansions, the causative gene for spinocerebellar ataxia (SCA) type 2 , are reported to coexist in a single family with ataxia. ${ }^{1}$ Therefore, this observation raises an interesting possibility that C9orf72 repeat/expansions could be genetic modifiers in CAG-repeat SCAs and might influence the disease progression.

Therefore, we studied 277 patients with SCA1, 2, 3 and 6 from the Clinical Research Consortium for Spinocerebellar Ataxias (CRC-SCA) cohort, ${ }^{2}$ and we determined the C9orf72 repeat length as previously described. ${ }^{3}$ The Scale for Assessment and Rating of Ataxia (SARA) and the 9-item Patient Health Questionnaire (PHQ-9) were used to measure the severity of ataxia and depression, respectively. We studied the rate of ataxia and depression progression using generalized estimating equation to test whether the intermediate repeats of C9orf72 were associated with ataxia or depression progression in SCAs. As described previously, full repeat expansions of C9orf72 were defined as $\geq 31$ hexanucleotide repeats whereas intermediate repeat expansions were 8-30. ${ }^{3-5}$

We identified seven patients (3 out of 51 SCA1; 2 out of 58 SCA2; 1 out 109 SCA3; 1 out of 59 SCA6) with pathogenic C9orf72 repeat expansions. None of the 7
cases had motor neuron disease, but they had various degrees of motor neuron signs
(Table 1). Compared to cognitively normal control (The original paper cites 1039 Europeans and 620 African American) population, ${ }^{5}$ the frequencies of expanded C9orf72 repeats in our cohort were significantly higher in SCA1, 2 , and 6 but not in SCA3 (Supplemental table 1). Forty percent of SCA patients carry intermediate C9orf72 repeat expansions, and the demographic and clinical features of SCA subjects with normal and intermediate alleles of C9orf72 are shown in Supplemental table 2. Intermediate C9orf72 repeat expansions did not influence the rate of ataxia progression but were associated with different rates of depressive symptom progression in SCA1, 3, and 6 (SCA1: $\beta=-1.90, p<0.005$; SCA3: $\beta=3.48$, $p<0.001$; SCA6: $\beta=-1.72, p<0.05$; Supplemental table 3).

In the present study, we identified patients of SCA1, 2, 3, and 6 who also carry pathogenic C9orf72 repeat expansions. Intermediate C9orf72 repeat expansions might influence the non-motor symptom (i.e. depression) progression in SCAs. Our study highlights the genetic interactions between repeat expansions.

The presence of CAG repeat expansions could interfere with the DNA repair process, ${ }^{6}$ which may destabilize C9orf72 repeat expansions and explain the co-existence of C9orf72 repeat expansions and expanded CAG repeats. Since cerebellar pathology could be found in C9orf72-linked ALS, the presence of C9orf72 repeat expansions might affect polyglutamine aggregates preferentially in the cerebellum or brainstem structures implicated in depression.

In conclusion, our study provides supporting evidence that repeat expansions of

C9orf72 may be genetic modifiers in SCAs, and perhaps ataxia patients in general. ${ }^{7}$ Therefore, the interplay of repeat expansions in two different loci may lead to diverse clinical phenotypes in degenerative cerebellar ataxia.

## Author contributions

Ms. Figueroa: study concept and design, statistical analysis and interpretation, writing the manuscript, critical revision of the manuscript for important intellectual content.

Dr. Gan: study concept and design, statistical analysis and interpretation, critical revision of the manuscript for important intellectual content.

Dr. Perlman: acquisition of data.
Dr. Wilmot: acquisition of data.

Dr. Gomez: acquisition of data.
Dr. Schmahmann: acquisition of data.
Dr. Paulson: acquisition of data.

Dr. Shakkottai: acquisition of data.

Dr. Ying: acquisition of data.
Dr. Zesiewicz: acquisition of data.
Dr. Bushara: acquisition of data.

Dr. Geschwind: acquisition of data.
Dr. Xia: acquisition of data.
Dr. Subramony: study concept and design, acquisition of data, study supervision.

Dr. Ashizawa: study concept and design, acquisition of data, critical revision of the
manuscript for important intellectual content, study supervision.
Dr. Pulst: study concept and design, acquisition of data, critical revision of the manuscript for important intellectual content, study supervision.

Dr. Kuo: study concept and design, acquisition of data, analysis and interpretation, critical revision of the manuscript for important intellectual content.

Disclosure: Dr. Zesiewicz has served as a clinical advisor for Steminent Biotherapeutics, and she has received travel reimbursement from the department of neurology at University of Southern Florida; has received travel reimbursement for a Biohaven Pharmaceuticals meeting. Dr. Zesiewicz has served on the editorial board for Neurodegenerative Disease Management and Tremor and other Hyperkinetic Movements, and has received research support for her division for approximately 20 clinical trials for Parkinson's disease, Friedreich's ataxia, and spinocerebellar ataxias. Dr. Zesiewicz's division is a site in a multi-site trial of Parkinson's disease patients with the LRRK2 mutation and is sponsored by the National Institutes of Health but funded by Emory University. The rest of authors report no conflicts of interest.

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Table 1. Demographic and clinical features of 7 SCA patients with full C9orf72 repeat expansions

|  |  | CAG Repeats | C9orf72 | Gender | Age of | SARA | Mental Status | Motor neuron deficits |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| ${ }_{7}^{6}$ umber <br> 8 | Type | Number <br> (Small/Large) | Repeats |  | Onset |  |  | Fasciculation | Weakness | Reflexes of Extremities | Plantar Reflex | Spasticity |
| $\begin{aligned} & 9 \\ & 10 \end{aligned}$ | 1 | $30 / 44$ | 5/>30 | Woman | 38 | 10.5 | MoCA 25/30 | Present | None | Hyperreflexia in biceps, patellar and Achilles | Extensor | Mild in lower and upper limbs |
| $\begin{aligned} & 11 \\ & 12 \\ & 13 \end{aligned}$ | 1 | 30/42 | 10/>30 | Woman | 45 | 0.5 | MoCA 30/30 | Present | None | Hyperreflexia in biceps, patellar and Achilles | Flexor | Not evaluated |
| 34 15 | 1 | 30/42 | 10/>30 | Woman | 52 | 8.5 | MoCA 25/30 | Present | None | Hyperreflexia in biceps, patellar and Achilles | Flexor | Mild in lower limbs |
| $\begin{aligned} & 16 \\ & 17 \\ & 18 \\ & 19 \end{aligned}$ | 2 | $22 / 39$ | 6/>30 | Man | 35 | 24 | Not evaluated | None | None | Areflexia in biceps, patellar and Achilles | Extensor | Moderate in lower limbs, mild in upper limbs |
| 20 21 22 23 | 2 | 22/39 | 2/>30 | Woman | 40 | 24.5 | Poor in serial 7s | Mild in tongue/ face and four limbs | Mild in four limbs | Areflexia in biceps, patellar and Achilles | Flexor | None |
| 28 25 26 27 | 3 | 28/70 | 2/>30 | Woman | 48 | 30.5 | Not evaluated | Moderate in tongue and face | Mild in four limbs | Areflexia in biceps, hyperreflexia in patellar and Achilles | Extensor | Moderate in four limbs |
| $\underline{8}$ | 6 | 11/23 | 8/>30 | Man | 59 | 12.5 | Not evaluated | None | None | None | None | None |

29 Abbreviations: SCA: spinocerebellar ataxias, SARA: scale for ataxia rating and assessment, MoCA: Montreal Cognitive Assessment, Serial 7s: serial sevens subtraction test,

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|  | Normal American population | SCA1 | SCA2 | SCA3 | SCA6 |  |  |
| C9orf72 expansions $\geq \mathbf{3 1 , ~} \mathbf{n}$ | 1659 | 48 | 56 | 108 | 58 |  |  |
| C9orf72 expansions $\leq \mathbf{3 0}, \mathrm{n}$ | 0 | 3 | 2 | 1 | 1 |  |  |
| Total, $\mathbf{n}$ | 1659 | 51 | 58 | 109 | 59 |  |  |
| $\boldsymbol{p}$ value ${ }^{\mathrm{a}}$ | - | 0.000 | $0.001^{\mathrm{a}}$ | $0.062^{\mathrm{a}}$ | $0.034^{\mathrm{a}}$ |  |  |

American normal controls are from the reference 5.
${ }^{\text {a }}$ Fisher exact test



Abbreviations: PHQ-9= The 9-item Patient Health Questionnaire, SARA = Scale for Assessment and Rating of Ataxia, SCA = Spinocerebellar Ataxia.

Values represent mean $\pm$ standard deviation or number, and for variables with non-normal distribution, the median is reported as well.
${ }^{\text {a }}$ Chi-square test
${ }^{\mathrm{b}} 2$ independent samples t-test
${ }^{\text {c }} 2$ independent samples Mann-Whitney $U$ test
${ }^{d}$ A Bonferroni correction was made to adjust for multiple comparisons, $\mathrm{p}<0.007$ were considered significant ( 7 tests in each subtypes)

## Supplemental Table 3. Longitudinal SARA and PHQ-9 scores of normal and intermediate C9orf72 repeat expansion in GEE models

| Regression coefficients of SARA score |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Age of first visit (yrs) | Sex | Expanded CAG repeats | Allele type of C9orf72 | Visit time | Allele type of C9orf72 <br> $\times$ Visit time |
| SCA1 | 0.62**** | 4.71** | 1.42**** | 2.66 | 0.95* | -0.04 |
| SCA2 | $0.42^{* * * *}$ | -1.40 | 1.87**** | 0.64 | 0.40 | -0.84 |
| SCA3 | $0.67 * * * *$ | -0.82 | 1.88**** | 0.14 | 0.37* | -0.02 |
| SCA6 | 0.11 | -1.68 | 0.71 | 2.12 | 0.23 | 0.96 |
| Regression coefficients of PHQ-9 score |  |  |  |  |  |  |
|  | Age of first visit (yrs) | Sex | Expanded CAG repeats | Allele type of C9orf72 | Visit <br> time | Allele type of C9orf72 <br> $\times$ Visit time |
| SCA1 | 0.08 | -0.37 | 0.25 | 2.16 | 0.32 | -1.90*** |
| SCA2 | 0.01 | -1.35 | -0.12 | 1.54 | 1.00*** | -0.63 |
| SCA3 | 0.26*** | 2.41 | 0.66* | -5.49**** | -0.38 | $3.48^{* * * *}$ |
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Abbreviations: GEE = generalized estimating equation, PHQ-9 = The 9-item Patient Health Questionnaire, SARA = Scale for Assessment and Rating of Ataxia SCA = Spinocerebellar Ataxia.
*p <0.05, **p < 0.01, ***p < 0.005, ****p <0.001
Sex: women $=0$, men $=1$
Allele type of C9orf72: normal allele $=0$, intermediate allele $=1$
Visit time: time order every 6 month during the two-year follow up


[^0]:    Abbreviations: PHQ-9= The 9-item Patient Health Questionnaire, SARA = Scale for Assessment and Rating of Ataxia, SCA = Spinocerebellar Ataxia.
    Values represent mean $\pm$ standard deviation or number, and for variables with non-normal distribution, the median is reported as well.
    ${ }^{\text {a }}$ Chi-square test
    ${ }^{\mathrm{b}} 2$ independent samples t-test

