

LETTER TO THE EDITOR

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C9orf72 deficiency promotes motor deficits of a C9ALS/FTD mouse model in a dose-dependent manner



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G4C2 hexanucleotide repeat expansions in the first intron of *C9ORF72* are the most common cause of familial amyotrophic lateral sclerosis (ALS) and frontotemporal dementia (FTD) (collectively, C9ALS/FTD) [4, 6, 11, 14]. Haploinsufficiency (loss-of-function) of C9ORF72 protein is a key proposed disease mechanism which may act in parallel with gain-of-function mechanisms, including toxic RNAs from repeat transcription and dipeptide repeat proteins (DPRs) from repeat-associated non-AUG (RAN) translation [5, 9, 17]. However, the effect of *C9orf72* deficiency in the background of gain-of-function has not been examined in vivo. Neither heterozygous nor homozygous knockout (KO) of *C9orf72* in neurons leads to motor deficits in mice [8]. Recently, gain-of-function mouse models were generated using a *C9ORF72* bacterial artificial chromosome (BAC) from C9ALS/FTD patient DNA under the control of the endogenous regulatory elements. Interestingly, three out of four of these *C9-BAC* transgenic mice did not develop motor behavior deficits, even at advanced ages [7, 12, 13]. Since these *C9-BAC* mouse models contain elevated C9orf72 proteins from the endogenous mouse gene, we hypothesized that C9orf72 provides neuroprotective effects against motor deficits in *C9-BAC* mice.

To test this hypothesis and investigate the in vivo significance of C9orf72 haploinsufficiency, we crossed *C9orf72*^{+/-} mice with *C9-BAC* mice and examined the consequences of C9orf72 protein dose reduction (loss-of-function) in the background of *C9-BAC* (gain-of-function). We found that *C9orf72* loss and haploinsufficiency exacerbate motor behavior deficits in

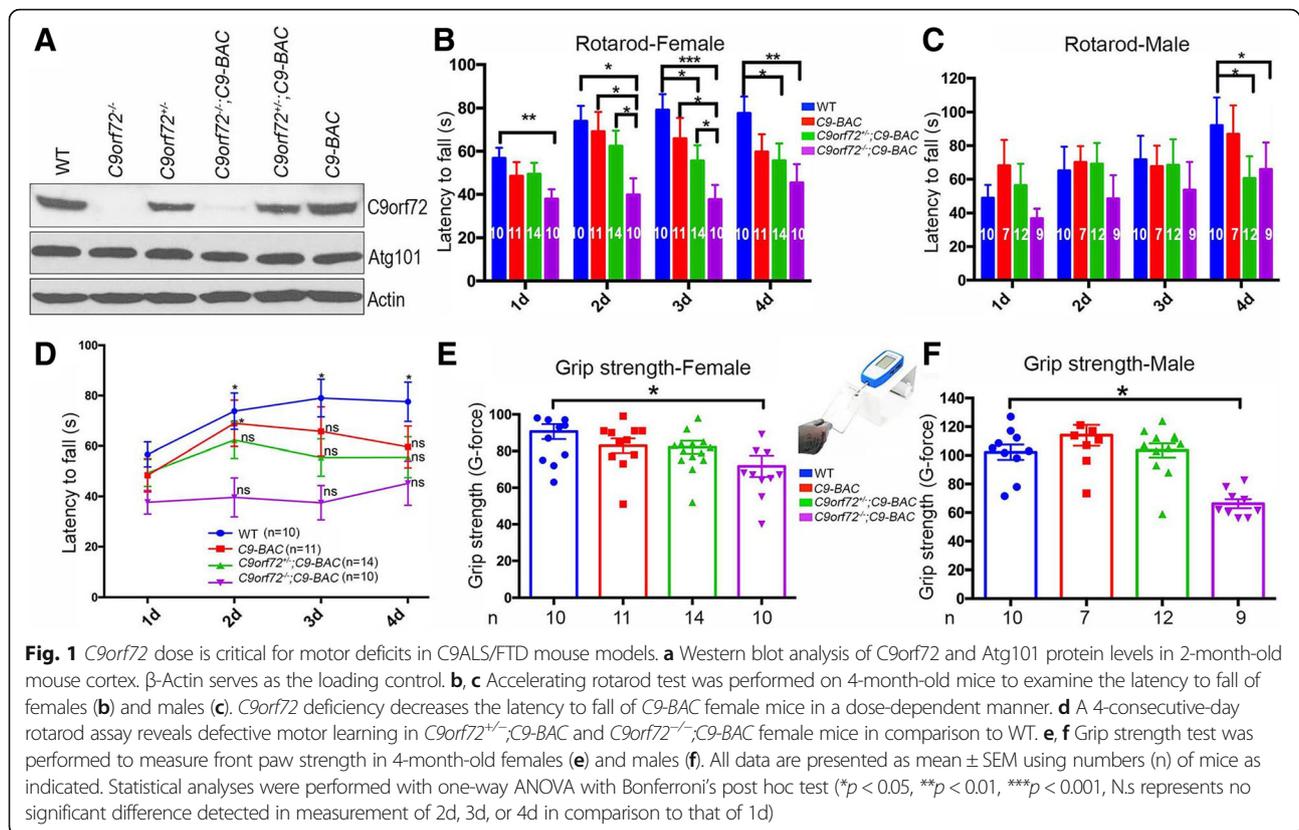
a dose-dependent manner, and this occurs early in the course of pathogenesis (4 months of age). Among the four published *C9-BAC* mouse models, we selected the one with motor deficits (we refer to this *C9orf72 BAC*^{Tg/+} model as the *C9-BAC* line here) [10]. To reduce C9orf72 protein levels at different doses, we crossed *C9orf72*^{+/-} and *C9-BAC* mice for two generations. We isolated proteins from brain tissues and confirmed the expected C9orf72 protein dose reduction (Fig. 1a, Additional file 1: Figure S1A). The unchanged protein level of Atg101, which is associated with the C9orf72/Smcr8 complex based on our previous study [16], suggests the specificity of C9orf72 reduction (Fig. 1a, Additional file 1: Figure S1A).

To study effects of *C9orf72* deficiency on the motor behaviors of *C9-BAC* mice, we monitored a cohort of mice [20 WT (10 females + 10 males), 18 *C9-BAC* (11 females + 7 males), 26 *C9orf72*^{+/-};*C9-BAC* (14 females + 12 males), and 19 *C9orf72*^{-/-};*C9-BAC* (10 females + 9 males)]. We excluded *C9orf72*^{+/-} and *C9orf72*^{-/-} mice for the following reasons: *C9orf72* heterozygous and homozygous KO mice exhibited no neurodegeneration and motor deficits based on previous studies [8]; complete deletion of *C9orf72*, which does not occur in C9ALS/FTD patients, led to autoimmune disorders and reduced survival in mice [1], which may complicate large-scale behavior and survival studies. We found that there were no significant differences among the four tested groups in their survival around 4 months, when behaviors were assessed. They also exhibited similar body weights, taking the sex of the mice into account (Additional file 1: Figure S1B-1C). To examine their general anxiety levels, we performed an open field test [3]. *C9-BAC* mice with different C9orf72 levels behaved similarly in total distance traveled,

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distance traveled in the center, and time spent in the center (Additional file 1: Figure S1E-1G).

We next examined their motor coordination and balance using an accelerating (4–40 rpm in 5 min) rotarod test. Mice were given five trials per day, with an inter-trial interval of 20 min, for 4 consecutive days. A *C9orf72* dose-dependent decrease in latency to fall was detected in *C9-BAC* female mice (Fig. 1b), and in *C9-BAC* male mice on day 4 of the rotarod assay (Fig. 1c). These results suggest that motor coordination is sensitive to *C9orf72* protein levels in *C9-BAC* mice. We further analyzed motor learning in female mice. WT mice exhibited an increase in latency to fall over the course of 4 consecutive days, indicating active motor learning (Fig. 1d). Latency to fall of *C9-BAC* mice was increased on day 2 but dropped on days 3 and 4 (Fig. 1d). Importantly, there was no increase in latency to fall from day 1 to day 4 in *C9orf72*^{+/-};*C9-BAC* and *C9orf72*^{-/-};*C9-BAC* animals (Fig. 1d). These results suggest that *C9orf72* deficiency impaired motor coordination and motor learning of *C9-BAC* mice in a dose-dependent manner.

To examine motor strength, we measured forearm grip strength and found that it was significantly reduced in both male and female *C9orf72*^{+/-};*C9BAC* animals compared to other genotypes (Fig. 1e, f). Lastly, we measured the maximal speed at which each animal fell from the rotarod device. Results showed that

C9orf72 deficiency, in a dose-dependent manner, decreased the maximum speed at which *C9-BAC* mice fell (Additional file 1: Figure S1H, S1I), which is consistent with the data on their latency to fall.

The rotarod assay revealed more evident motor impairment in female mice than in male mice. This could be due to toxic gain-of-function since *C9-BAC* female mice exhibited earlier and more pronounced abnormalities than male mice [10]. It will be important to examine using similar cohorts of mice whether motor neurons (MNs) degenerate or reduce in number in a *C9orf72* dose-dependent manner and whether these deficits correlate with the observed motor behavior deficits. Future studies should also investigate whether *C9orf72* exhibits dose-dependent effects in the three other *C9-BAC* mouse models [7, 12, 13]. It will be informative to examine the effects of *C9orf72* deficiency in the background of adeno-associated virus (AAV)-mediated G4C2 repeat expression [2]. Our study indicates that *C9orf72* haploinsufficiency contributes to disease onset in a mouse model by exacerbating the pathogenic effects of RNA/DPR-mediated neurotoxicity. Together with a recent report on patient iPSC-derived MNs [15], this study suggests indeed that we should focus more on the combination of loss- and toxic gain-of-function. Together, for the first time, our mouse genetic studies showed that *C9orf72* loss or haploinsufficiency in a gain-of-function mouse model of

C9ALS/FTD exacerbate motor behavior deficits in a dose-dependent manner, demonstrating the importance of C9orf72 haploinsufficiency in vivo.

Additional file

Additional file 1: Figure S1. Characterization of C9-BAC mice with C9orf72 dose reduction. (A) Quantification of C9orf72/Atg101 protein levels. Data are presented as mean \pm SEM from three independent experiments. (B, C) Body weight of female (B) and male (C) mice at 4 months of age. (D-G) Open field test was performed on 4-month-old mice to examine the total distance traveled (E), distance traveled in the center (F), and percentage of time spent in the center (G). (H, I) Quantification of the maximal g force from five trials of rotarod assay. C9orf72 deficiency decreases the maximal g force of C9-BAC female mice in a dose-dependent manner. All data are presented as mean \pm SEM using numbers (n) of mice as indicated. Statistical analyses were performed with one-way ANOVA with Bonferroni's post hoc test (* $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, n.s represents no significant difference detected). (PDF 2855 kb)

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Authors' contributions

QS, QC, CL performed all behavior studies and statistical analyses. MY and WZ helped with the manuscript writing. J-FC designed and interpreted the experiments and wrote the manuscript. All authors read and approved the final manuscript.

Competing interests

The authors declare that they have no competing interests.

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References

- Burberry A, Suzuki N, Wang J-Y, Moccia R, Mordes DA, Stewart MH et al (2016) Loss-of-function mutations in the C9ORF72 mouse ortholog cause fatal autoimmune disease. *Sci Transl Med* 8:347ra93–347ra93
- Chew J, Gendron TF, Prudencio M, Sasaguri H, Zhang Y-J, Castanedes-Casey M et al (2015) Neurodegeneration. C9ORF72 repeat expansions in mice cause TDP-43 pathology, neuronal loss, and behavioral deficits. *Science* 348:1151–1154
- Crawley JN (1999) Behavioral phenotyping of transgenic and knockout mice: experimental design and evaluation of general health, sensory functions, motor abilities, and specific behavioral tests. *Brain Res* 835:18–26
- DeJesus-Hernandez M, Mackenzie IR, Boeve BF, Boxer AL, Baker M, Rutherford NJ et al (2011) Expanded GGGGCC hexanucleotide repeat in noncoding region of C9ORF72 causes chromosome 9p-linked FTD and ALS. *Neuron* 72:245–256
- Gao F-B, Almeida S, Lopez-Gonzalez R (2017) Dysregulated molecular pathways in amyotrophic lateral sclerosis-frontotemporal dementia spectrum disorder. *EMBO J* 36:2931–2950
- Gijselsinck I, Van Langenhove T, van der Zee J, Sleegers K, Philtjens S, Kleinberger G et al (2012) A C9orf72 promoter repeat expansion in a Flanders-Belgian cohort with disorders of the frontotemporal lobar degeneration-amyotrophic lateral sclerosis spectrum: a gene identification study. *Lancet Neurol* 11:54–65
- Jiang J, Zhu Q, Gendron TF, Saberi S, McAlonis-Downes M, Seelman A et al (2016) Gain of toxicity from ALS/FTD-linked repeat expansions in C9ORF72 is alleviated by antisense oligonucleotides targeting GGGGCC-containing RNAs. *Neuron* 90:535–550
- Koppers M, Blokhuis AM, Westeneng H-J, Terpstra ML, Zundel CAC, Vieira de Sá R et al (2015) C9orf72 ablation in mice does not cause motor neuron degeneration or motor deficits. *Ann Neurol* 78:426–438
- Ling S-C, Polymenidou M, Cleveland DW (2013) Converging mechanisms in ALS and FTD: disrupted RNA and protein homeostasis. *Neuron* 79:416–438
- Liu Y, Pattamatta A, Zu T, Reid T, Bardhi O, Borchelt DR et al (2016) C9orf72 BAC mouse model with motor deficits and neurodegenerative features of ALS/FTD. *Neuron* 90:521–534
- Majounie E, Renton AE, Mok K, Doppler EGP, Waite A, Rollinson S et al (2012) Frequency of the C9orf72 hexanucleotide repeat expansion in patients with amyotrophic lateral sclerosis and frontotemporal dementia: a cross-sectional study. *Lancet Neurol* 11:323–330
- O'Rourke JG, Bogdanik L, Muhammad AKMG, Gendron TF, Kim KJ, Austin A et al (2015) C9orf72 BAC transgenic mice display typical pathologic features of ALS/FTD. *Neuron* 88:892–901
- Peters OM, Cabrera GT, Tran H, Gendron TF, McKeon JE, Metterville J et al (2015) Human C9ORF72 Hexanucleotide expansion reproduces RNA foci and dipeptide repeat proteins but not neurodegeneration in BAC transgenic mice. *Neuron* 88:902–909
- Renton AE, Majounie E, Waite A, Simón-Sánchez J, Rollinson S, Gibbs JR et al (2011) A hexanucleotide repeat expansion in C9ORF72 is the cause of chromosome 9p21-linked ALS-FTD. *Neuron* 72:257–268
- Shi Y, Lin S, Staats KA, Li Y, Chang W-H, Hung S-T et al (2018) Haploinsufficiency leads to neurodegeneration in C9ORF72 ALS/FTD human induced motor neurons. *Nat Med* 24:313–325
- Yang M, Liang C, Swaminathan K, Herrlinger S, Lai F, Shiekhattar R et al (2016) A C9ORF72/SMCR8-containing complex regulates ULK1 and plays a dual role in autophagy. *Sci Adv* 2:e1601167–e1601167
- Zu T, Pattamatta A, LPW R (2018) Repeat-Associated Non-ATG Translation in Neurological Diseases. *Cold Spring Harb Perspect Biol* 10:a033019

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