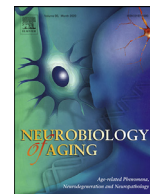


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## Neurobiology of Aging

journal homepage: [www.elsevier.com/locate/neuaging.org](http://www.elsevier.com/locate/neuaging.org)Deletion of the inflammatory S100-A9/MRP14 protein does not influence survival in hSOD1<sup>G93A</sup> ALS mice

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## ABSTRACT

Neuroinflammation is a hallmark of Amyotrophic Lateral Sclerosis (ALS) in hSOD1<sup>G93A</sup> mouse models where microglial cells contribute to the progressive motor neuron degenerative process. S100-A8 and S100-A9 (also known as MRP8 and MRP14, respectively) are cytoplasmic proteins expressed by inflammatory myeloid cells, including microglia and macrophages. Mainly acting as a heterodimer, S100-A8/A9, when secreted, can activate Toll-like Receptor 4 on immune cells, leading to deleterious proinflammatory cytokine production. Deletion of S100a9 in Alzheimer's disease mouse models showed a positive outcome, reducing pathology. We now assessed its role in ALS. Unexpectedly, our results show that deleting S100a9 in hSOD1<sup>G93A</sup> ALS mice had no impact on mouse survival, but rather accelerated symptoms with no impact on microglial activation and motor neuron survival, suggesting that blocking S100-A9 would not be a valuable strategy for ALS.

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## 1. Introduction

Amyotrophic Lateral Sclerosis (ALS) is the most common, adult onset, motor neuron disease leading to fatal paralysis. While microglial cells, the macrophages of the central nervous system are involved in disease progression, microglial cell pathways involved in motor neuron death remain largely unidentified (Beers et al., 2006; Boillée et al., 2006). S100-A8 (also known as MRP8) and S100-A9 (also known as MRP14 or Calgranulin B) are cytoplasmic proteins of the S100 calcium binding protein family, implicated in cytoskeleton remodeling, crucial for cell migration and phagocytosis. S100-A8 and S100-A9 are mainly expressed by myeloid cells and overexpressed during inflammation (Akiyama et al., 1994; Wang et al., 2018). Although some anti-inflammatory properties have been reported, they are secreted as danger-associated molecular pattern signals. They form a heterocomplex (S100-A8/A9 called calprotectin) inducing Toll-like Receptor-4 (TLR4) responses that leads to deleterious proinflammatory cytokine release (Donato

et al., 2013; Ehrchen et al., 2009; Vogl et al., 2007; Wang et al., 2018). Deleting S100a9 in Alzheimer's disease (AD) mouse models leads to increased capacity of microglia to phagocytose amyloid plaques and decreased memory impairment (Akiyama et al., 1994; Kim et al., 2014; Kummer et al., 2012). Although extracellular deposits are not present in ALS, secretion of misfolded mutant SOD1 or its release through extracellular vesicles are suspected to contribute to disease spreading (Silverman et al., 2016; Urushitani et al., 2006). In addition, S100-A8/A9 contributes to damage in focal ischemia through microglial recruitment and activation at a (sterile) inflammation site (Ziegler et al., 2009). Based on this data indicating a role of S100-A8/A9 in several models of neuroinflammation, and existing evidence suggesting disease contribution of neuroinflammatory reactions in ALS, we deleted S100a9 in hSOD1<sup>G93A</sup> ALS mice to uncover whether S100-A8/A9 would be involved in the mutant SOD1-mediated ALS toxicity and could represent a potential microglial target to increase motor neuron survival.

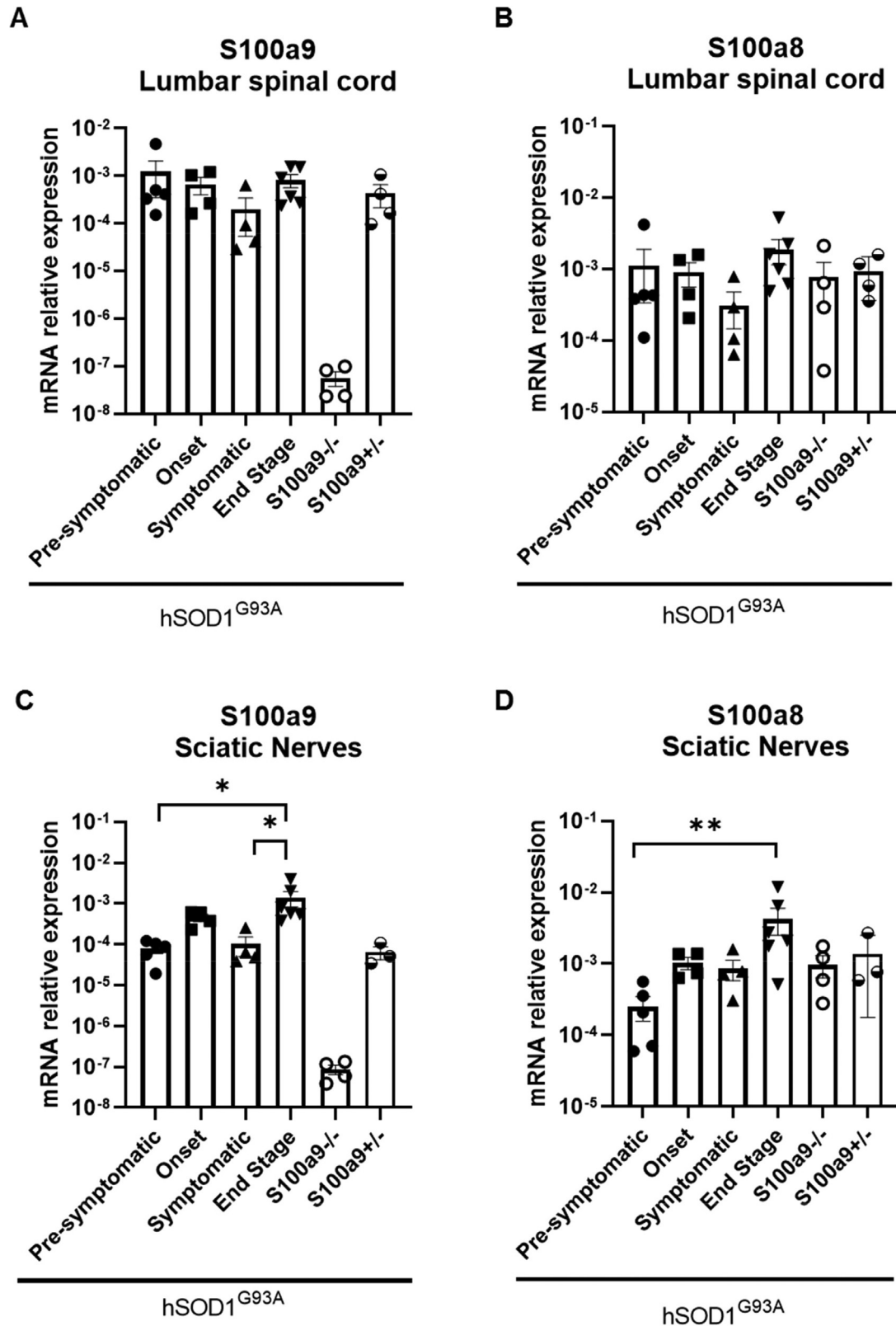
## 2. Methods

See Supplementary Material for extended methods. Mice from the three different genotypes (hSOD1<sup>G93A</sup>:S100a9+/+, hSOD1<sup>G93A</sup>:S100a9+/-, hSOD1<sup>G93A</sup>:S100a9-/-) were true litter-

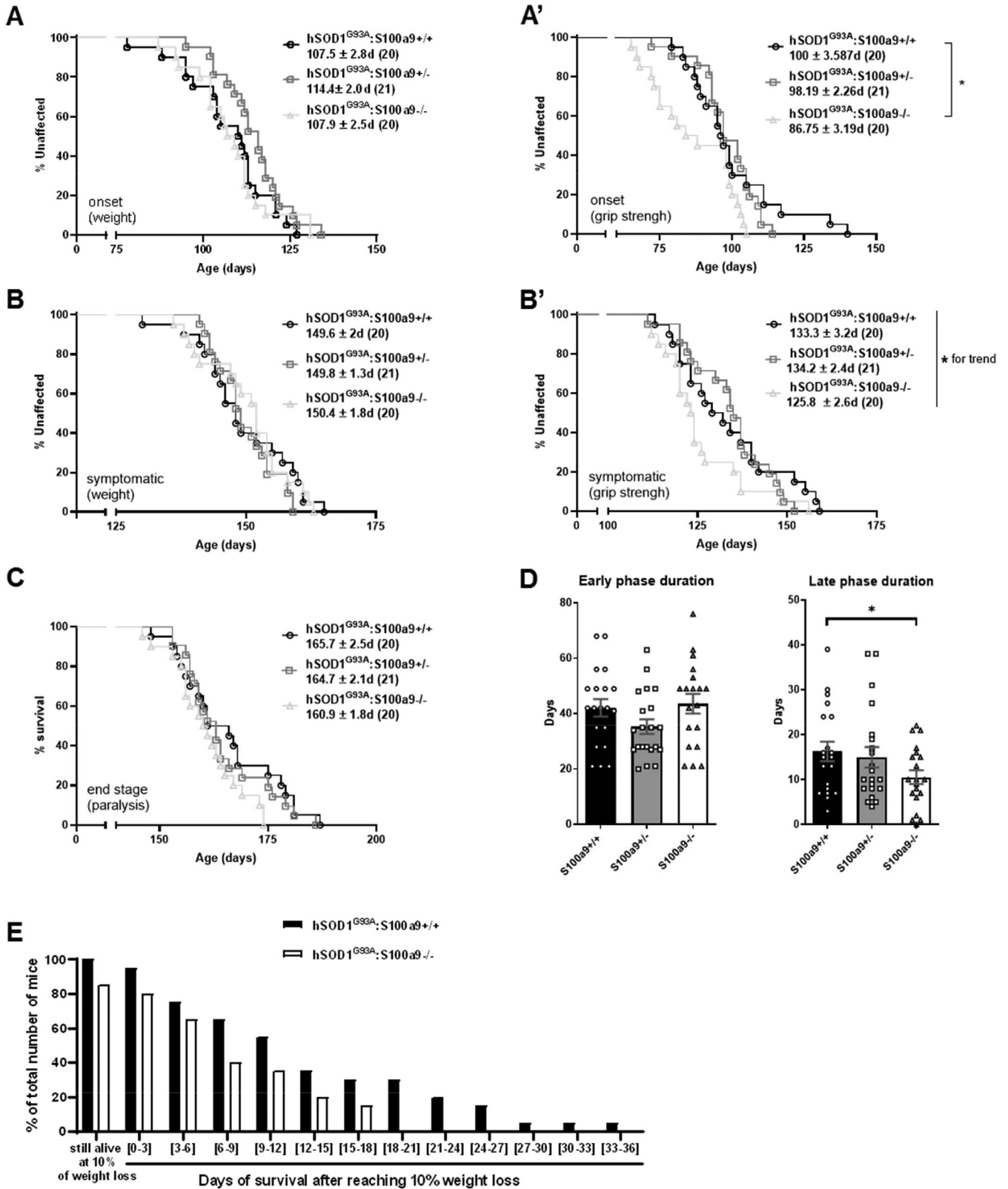
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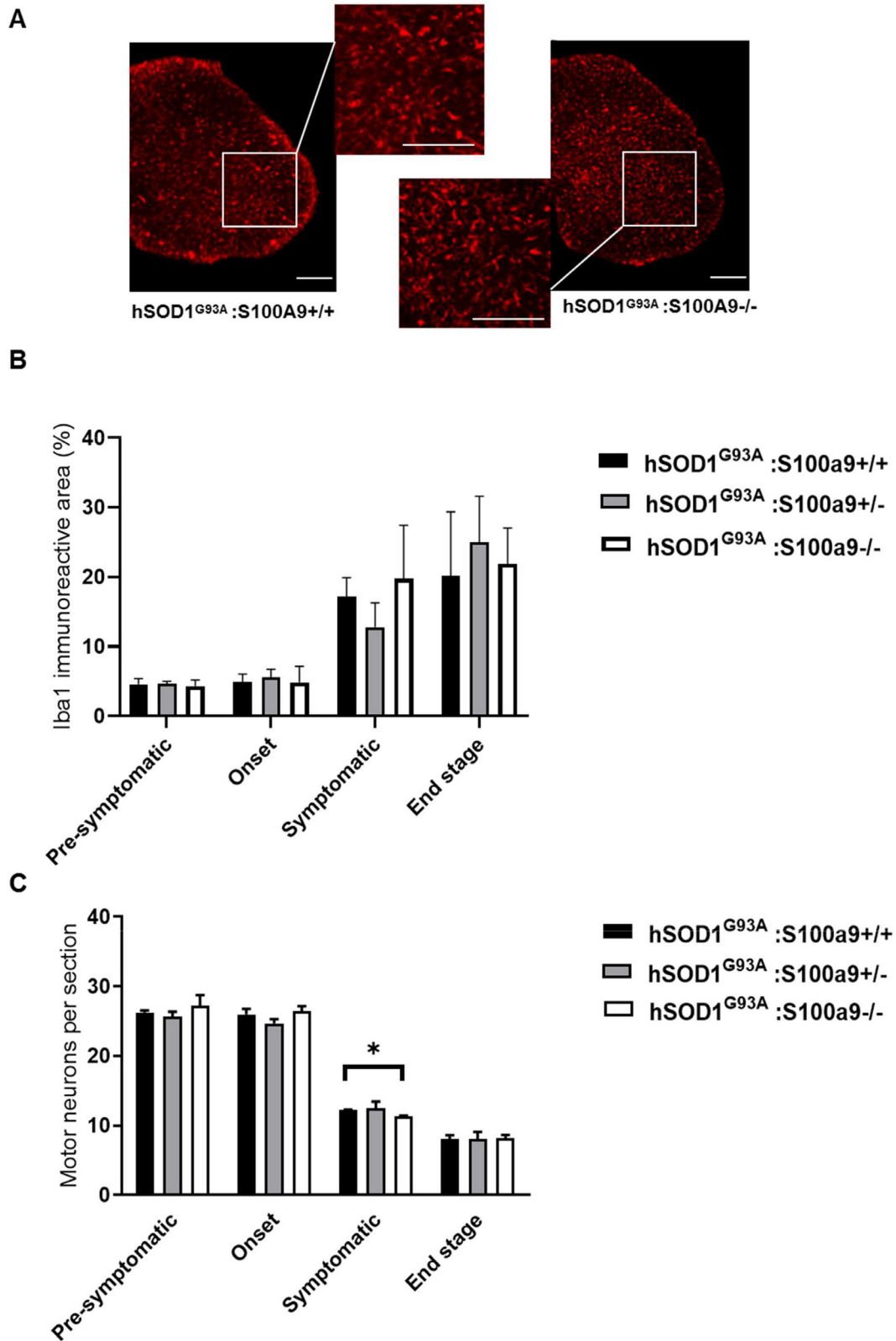
# These authors contributed equally.



**Fig. 1.** S100a8 and S100a9 mRNA expression levels are not modified in lumbar spinal cords throughout the disease, but increased at end stage in sciatic nerves of hSOD1<sup>G93A</sup> ALS mice. mRNA levels of S100a8 and S100a9 were measured in whole lumbar spinal cords and sciatic nerves at four different disease stages in hSOD1<sup>G93A</sup> ALS mice. In hSOD1<sup>G93A</sup>:S100a9<sup>+/-</sup> mice, 'Presymptomatic': 50 days of age, 'Onset': defined as the weight peak, in mean at the age of 107 days, 'Symptomatic stage': defined by 10% of weight loss, in mean at the age of 150 days, End stage: defined as complete hindlimb paralysis reached in mean at the age of 166 days, and in hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> and hSOD1<sup>G93A</sup>:S100a9<sup>+/-</sup> at onset. Whole lumbar spinal cord (A, B) or sciatic nerve (C, D) tissue mRNA levels for S100a9 (A, C) and S100a8 (B, D) were measured by reverse-transcription quantitative PCR and normalized to the housekeeping gamma-actin (Actg1) gene. Results are shown relative to Actg1 expression. Bars represent Means  $\pm$  SEM for  $n = 4$  mice at every stage (except at presymptomatic  $n = 5$ , end stage  $n = 6$  and hSOD1<sup>G93A</sup>:S100a9<sup>+/-</sup>  $n = 3$  mice). \*  $p < 0.05$ , \*\*  $p < 0.01$  (Kruskal-Wallis test followed by Dunnett's post-hoc test).



**Fig. 2.** Deletion of S100a9 in ALS mice does not modify survival but slightly accelerates symptoms. (A–C) Kaplan-Meier analyses of ages when onset (weight peak (A)), or grip strength peak (A’), symptomatic stage (10% of weight loss (B), or 35% of grip strength loss (B’)) and end stage (complete hindlimb paralysis) (C) were reached, in hSOD1<sup>G93A</sup> (black), hSOD1<sup>G93A</sup>:S100a9<sup>+/-</sup> (gray) and hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> (light gray) mice. Means ± SEM are indicated with number of animals (gender balance) in brackets. \* *p* < 0.05 (log-rank test). (D left), Early disease duration corresponding to the mean of the difference between age at onset and age when reaching 10% of weight loss for each individual mouse. (D right), Late disease duration, corresponding to the mean of the difference between ages at 10% of weight loss and end stage for each individual mouse. hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> (black bars) *n* = 20 mice, hSOD1<sup>G93A</sup>:S100a9<sup>+/-</sup> (gray bars) *n* = 21 mice and hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> (white bars) *n* = 20 mice. Bars represent Means ± SEM. \* *p* < 0.05 (Student’s *t*-test). (E) Survival duration in days after reaching 10% weight loss. Bars represent the proportion of mice still alive within the indicated time frame. hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> (black bars) *n* = 20 mice, and hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> (white bars) *n* = 20 mice.



**Fig. 3.** Microglial activation and motor neuron survival are not affected by S100a9 deletion in hSOD<sup>G93A</sup> mice. (A) Representative pictures of lumbar spinal cord cross-sections stained against Iba1 reflecting microglial activation at onset. Right panel hSOD1<sup>G93A</sup>:S100A9<sup>+/+</sup> mouse, left panel: hSOD1<sup>G93A</sup>:S100A9<sup>-/-</sup> mouse. Scale bars, 100  $\mu$ m. (B) Microglial cell activation in lumbar spinal cord cross sections measured by anti-Iba1 fluorescent immunoreactive area. (C) Motor neuron numbers per lumbar spinal cord section. Bars represent Mean  $\pm$  SEM for n = 4 mice per stage and per genotype (except for n = 3 for hSOD1<sup>G93A</sup>:S100a9<sup>+/-</sup> at onset for motor neuron counts), gender balance. \*  $p < 0.05$  (mixed-effects analysis (ANOVA) followed by Tukey's post-hoc test).

mates, obtained through a two-step mating strategy between hSOD1<sup>G93A</sup> (Gurney et al., 1994) and S100a9<sup>-/-</sup> mice (Manitz et al., 2003), both on a C57BL/6 background. Mice were weighed and their grip-strength measured weekly from the presymptomatic stage (50 days of age) to disease end stage (complete hindlimb paralysis, around 165 days), with onset (weight or grip strength peak) and symptomatic (10% of weight loss or 35% of grip-strength loss (Chiot et al., 2020; Mesci et al., 2015)) stages determined retrospectively. At least 20 mice per genotype (gender balance) were used for the survival analysis. Lumbar spinal cords and sciatic nerves were collected at each time point from 3–4 mice per genotype (gender balance) to perform immunostainings for microglial cells (Iba1), to stain motor neurons for quantification (Nissl staining, cresyl-violet-acetate) and for RT-qPCR analysis of S100a8 and S100a9 (with Actg1, gamma-actin, as a normalizer).

### 3. Results

#### 3.1. S100a8 and S100a9 mRNA expression levels are not modified in lumbar spinal cords but are increased at disease end stage in sciatic nerves of ALS mice

S100a8 and S100a9 mRNAs being enriched in macrophages/microglia (Zhang et al., 2014), we measured S100a8 and S100a9 mRNA expression in lumbar spinal cord and sciatic nerve tissues of ALS mice, which shows increased microglial and peripheral nerve macrophage activation, throughout the disease. S100a8 and S100a9 mRNA levels were stable in the spinal cord (with the trend toward downregulation at the symptomatic stage, (Fig. 1A and B)). In sciatic nerves, S100a8 and S100a9 mRNA levels increased at disease end stage compared to the presymptomatic stage (Fig. 1C and D).

#### 3.2. Deleting S100a9 in ALS mice does not modify hSOD1<sup>G93A</sup> ALS mouse survival but slightly accelerates the appearance of disease symptoms

To determine the role of the S100-A8/A9 complex on the ALS disease course (and since S100a8 KO mice are embryonic lethal (Passey et al., 1999)), we deleted S100a9 in hSOD1<sup>G93A</sup> mice. While (littermate) hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> and hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> mice had similar maximum weight and grip-strength (Supplementary Fig. 1), and hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> mice reached disease onset, defined by weight peak, at the same age than hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> mice (Fig. 2A), there was a trend that hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> mice reached onset, defined by grip-strength peak, slightly earlier than hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> mice (Fig. 2A'). The same trend was also observed for the symptomatic stage (Fig. 2B and B'). However, hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> and hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> mice reached disease end stage both at the same age (Fig. 2C). However, the duration of the late phase (defined by weight) was shorter in hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> compared to hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> mice (Fig. 2D). Indeed, disease was accelerated from the symptomatic stage (defined by 10% of weight loss) onward (Fig. 2E). Of note, mice with deletion of only one copy of S100a9 (hSOD1<sup>G93A</sup>:S100a9<sup>+/-</sup>), showed similar results as hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> mice.

#### 3.3. Microglial activation and motor neuron survival are not affected by S100a9 deletion in hSOD1<sup>G93A</sup> mice

Microglial activation measured by Iba1+ immunoreactive area in the lumbar spinal cord (Fig. 3A) was increased in symptomatic hSOD1<sup>G93A</sup> mice, but regardless of S100a9 expression, without difference between hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> and hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup>

mice (Fig. 3A and B). In addition, while motor neurons progressively degenerated throughout the disease, motor neuron numbers were similar in hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> and hSOD1<sup>G93A</sup>:S100a9<sup>+/+</sup> mice at equivalent disease stages, (except for a minor difference at the symptomatic stage; Fig. 3C), showing no significant impact of S100a9 deletion on pathological hallmarks of the disease.

### 4. Discussion

While our hypothesis was that deletion of S100a9 would have a positive outcome on mutant SOD1-mediated ALS disease in mice, here, we show that, if at all, it rather had a negative effect, by marginally accelerating onset and the late disease phase. In addition, since S100-A9 was reported to be mainly expressed by innate immune cells, we expected that microglial cell reactivity, in the context of the disease, would be overall reduced in hSOD1<sup>G93A</sup>:S100a9<sup>-/-</sup> mice, what we, however, did not find. Several inflammatory properties mediated by S100-A9 made it a good target to be modulated in ALS, including its ability to induce chemotaxis of leukocytes and adherence of neutrophils, to activate inflammatory cells through TLR4 binding and MyD88/NFκB pathway activation, leading to proinflammatory cytokine release and production of reactive oxygen species (Ryckman et al., 2003; Vogl et al., 2007). However, in our study, deletion of S100a9 did not impact overall microglial activation or motor neuron survival in mutant SOD1 ALS mice.

In AD, a recent study showed that S100-A8 aggregated prior to amyloid plaque formation in APP transgenic mice (Lodeiro et al., 2017). Moreover, the S100-A8/A9 complex is highly expressed in microglial cells surrounding amyloid plaques in AD postmortem brain tissues and S100-A9 is overexpressed in cerebrospinal fluid of AD patients (Ha et al., 2010; Kim et al., 2014; Kummer et al., 2012). In addition, the S100-A8/A9 complex drives inflammation after ischemia, emphasizing its role in several inflammatory conditions (Ziegler et al., 2009). Our results suggest that, at least in mouse models, deletion of S100a9 has rather opposite effects in ALS than in AD and ischemia models.

In conclusion, although S100-A9 can play an important role in inflammation, is mainly expressed by phagocytic cells and is a microglial candidate to target in certain models with neuroinflammation, we showed that deletion of S100a9 in mutant SOD1 ALS mice did not alleviate but rather slightly exacerbated the disease. Therefore, this suggests that the S100-A8/A9 pathway does not contribute to mutant SOD1 mediated ALS toxicity and, at least based on this mouse model, its blockage does not represent a promising strategy to slow ALS.

#### Disclosure statement

The authors declare no conflict of interest.

#### Data contained in the manuscript

The authors declare that the data contained in the manuscript being submitted have not been previously published, have not been submitted elsewhere and will not be submitted elsewhere while under consideration at *Neurobiology of Aging*.

#### Animals

All animal procedures (including end stage definition) were performed in accordance with the guidelines for care and use of experimental animals of the European Union and approved by the ethics committee for animal experimentation n°5 in Ile-de-France to the UMS28, Centre d'expérimentation fonctionnelle, Paris, France.

## Authors approval

All author's have reviewed the content of the manuscript being submitted, approved its contents and validated the accuracy of the data.

## Authors' contributions

**Matthieu Ribon:** Investigation, Validation, Formal Analysis, Visualization, Writing Original draft; **Céline Leone:** Investigation, Validation, Formal Analysis; **Aude Chiot:** Investigation, Validation, Writing – Review & Editing; **Félix Berriat:** Investigation, Writing – Review & Editing; **Martine Rampanana:** Investigation, Visualization; **Julie Cottin:** Investigation, Visualization; **Delphine Bohl:** Writing – Review & Editing; **Stéphanie Millecamps:** Writing – Review & Editing; **Christian S. Lobsiger:** Conceptualization, Writing – Review & Editing; **Michael T. Heneka:** Conceptualization, Resources, Funding Acquisition, Writing – Review & Editing; **Séverine Boillée:** Supervision, Conceptualization, Funding Acquisition, Writing – Review & Editing.

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## Supplementary materials

Supplementary material associated with this article can be found, in the online version, at doi:10.1016/j.neurobiolaging.2021.01.015.

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