

1 **Elevated serum creatine kinase in the early stage of sporadic amyotrophic lateral**
2 **sclerosis**

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21 Key words: amyotrophic lateral sclerosis, creatine kinase, biomarker

22 Acknowledgements: This work was funded by a Grant-in-Aid (KAKENHI) from the Ministry of
23 Education, Culture, Sports, Science, and Technology of Japan (No. 17H04195); grants from the
24 Japan Agency for Medical Research and Development (Nos. 17ek0109221h0001 and
25 18ek0109221h0002); a grant from the Naito Foundation; and a grant from the Hori Sciences
26 and Arts Foundation.

27 Author Contributions: D.I., A.Ha., and M.K. conceived and designed the study. D.I., A.Ha.,
28 Y.H., S.Y., Y.K., and H.M. contributed to the acquisition of clinical data. D.I., M.I., and Y.I.
29 performed animal experiments. D.I. and M.K. performed analysis and interpretation of the data.
30 A.Hi. performed statistical analysis. D.I. drafted the manuscript, and A.Ha. and M.K. revised it
31 for intellectual content.

32 Potential conflict of interest: The authors have no relevant conflicts of interest to report.

33 **ABSTRACT**

34 **Objective:** To assess the changes of muscle-related biomarkers at the early stage of
35 amyotrophic lateral sclerosis, and to confirm these findings in an experimental animal model.

36 **Methods:** Thirty-nine subjects with sporadic amyotrophic lateral sclerosis and 20 healthy
37 controls were enrolled and longitudinally evaluated. We evaluated serum creatine kinase and
38 creatinine levels and appendicular lean soft tissue mass using dual X-ray absorptiometry. The
39 levels of biomarkers at early ALS stages were estimated using linear mixed models with
40 unstructured correlation and random intercepts. We also analyzed the longitudinal changes of
41 serum creatine kinase and creatinine, together with the mRNA levels of acetylcholine receptor
42 subunit γ (*Chrn3*) and muscle-associated receptor tyrosine kinase, markers of denervation, in
43 the gastrocnemius muscle of superoxide dismutase 1 (SOD1)^{G93A} transgenic mice, an animal
44 model of amyotrophic lateral sclerosis.

45 **Results:** The estimated levels of creatine kinase were higher in subjects with amyotrophic
46 lateral sclerosis at the early stage than in healthy controls, although the estimated appendicular
47 lean soft tissue mass and creatinine levels were equivalent between both groups, suggesting
48 that the elevation of creatine kinase precedes both muscular atrophy and subjective motor
49 symptoms in sporadic amyotrophic lateral sclerosis. In SOD1^{G93A} mice, the serum levels of
50 creatine kinase were elevated at 9 weeks of age (peri-onset) when *Chrn3* started to be up-
51 regulated, and were then down-regulated at 15 weeks of age, consistent with the clinical data
52 from patients with sporadic amyotrophic lateral sclerosis.

53 **Interpretation:** Creatine kinase elevation precedes muscular atrophy and reflects muscle
54 denervation at the early stage.

55 Introduction

56 Amyotrophic lateral sclerosis (ALS) is a fatal neurodegenerative disease caused by the
57 selective loss of upper and lower motor neurons. ALS causes rapidly progressive muscle
58 weakness and atrophy, leading to death in approximately 3–5 years from onset, chiefly due to
59 respiratory failure [1, 2]. Mutations in genes such as chromosome 9 open reading frame 72
60 (*C9orf72*), TAR DNA binding protein, fused in sarcoma, and superoxide dismutase 1 (*SOD1*)
61 have been identified in 50–70% of familial cases, as well as in a smaller population of sporadic
62 cases, suggesting that several biological mechanisms are involved in the pathophysiology of
63 ALS, e.g., RNA metabolism, protein homeostasis, nucleocytoplasmic trafficking, and
64 neuroinflammation [3–5]. These insights have inspired a myriad of attempts to develop disease-
65 modifying therapies for this devastating disease; nevertheless, most agents that have shown
66 promise in animal studies have failed to demonstrate clear efficacy in clinical trials [6, 7]. This
67 failure in translation may be attributable to various clinical and biological factors, among which
68 disease progression before the onset and early stages of neurological symptoms is a key issue
69 for the successful development of a cure for ALS.

70 It is now widely accepted that pathological changes begin long before the clinical symptoms
71 manifest, indicating the preclinical progression of neurodegenerative processes [8–10]. This
72 hypothesis is explicitly emphasized in Alzheimer's disease, in which amyloid β deposition, a
73 component of senile plaques, precedes brain volume loss and cognitive decline by a couple of
74 decades [8]. Similar changes of biomarkers have been reported in most neurodegenerative
75 diseases, including Huntington's disease, Parkinson's disease, and spinal and bulbar muscular
76 atrophy, another adult-onset motor neuron disease [11, 12]. These findings provide a theoretical
77 basis for the development of preventive therapies that target the preclinical phase of
78 neurodegenerative disorders [13, 14].

79 As for ALS, a few markers have been identified that show changes in gene mutation carriers at
80 the preclinical stage. For instance, dipeptide repeat proteins are detectable in the cerebrospinal
81 fluid of asymptomatic *C9orf72* mutation carriers [15]. In addition, alterations of microRNA
82 expression profiles in cerebrospinal fluid have also been reported in *C9orf72* mutation carriers
83 [16]. It is also reported that short-interval intracortical inhibition, an index of cortical
84 hyperexcitability in electrophysiological studies, is affected in asymptomatic *SOD1* mutation
85 carriers [17]. Furthermore, a decrease in the estimated number of motor units, another
86 electrophysiological marker of motor neuron degeneration, is detectable in asymptomatic *SOD1*

87 mutation carriers [18]. However, there is no report on the preclinical changes of biomarkers in
88 sporadic ALS, except for scattered studies on altered body fat and lipid metabolism as risk
89 factors for ALS [19, 20].

90 Here, we conducted a longitudinal study to estimate the changes of biomarkers at the very
91 early stage of ALS by analyzing clinical data from sporadic patients at the early stage. Our
92 results indicate the elevation of the serum levels of creatine kinase (CK) in the early stage of
93 ALS, which was reproduced in G93A mutant SOD1 (SOD1^{G93A}) mice, an animal model of ALS.

94

95 **Materials and Methods**

96 **Ethics and consent**

97 This study was conducted according to the Declaration of Helsinki, the Ethics Guidelines for
98 Human Genome/Gene Analysis Research, and the Ethical Guidelines for Medical and Health
99 Research Involving Human Subjects endorsed by the Japanese government. This study was
100 approved by the Ethics Review Committee of Nagoya University Graduate School of Medicine
101 (Nos. 2013-0035 and 2015-0041), and all participants gave written informed consent before
102 participation.

103 **Participants**

104 Subjects who were clinically diagnosed with the revised El Escorial Criteria of definite, probable,
105 or possible ALS were recruited consecutively. The principal inclusion criteria were no family
106 history and disease duration of ≤ 2 years at the time of enrollment. Subjects who had severe
107 complications such as malignancy, heart failure, or renal failure were excluded from this study.
108 Age- and sex-matched healthy controls were also recruited during the same period as the ALS
109 patients. Subjects with sporadic ALS were assessed during hospitalization at the initial
110 evaluation and followed up every 6 months at our outpatient clinic as long as they were able to
111 continue to attend out-patient clinic. Healthy controls were followed up every 6 or 12 months at
112 the outpatient clinic.

113 Subjects with ALS who were evaluated twice or more were analyzed longitudinally. Results of
114 DXA which were performed within 1 week after a radiological examination with contrast agents
115 or radioisotopes were excluded as they influence the results of dual-energy X-ray
116 absorptiometry (DXA).

117 We also retrospectively analyzed serum CK before the onset, which had been measured for the
118 purpose of medical practice, in patients with or without follow-up data. All study subjects were

119 Japanese and observed at the Nagoya University Hospital between May 2013 and March 2018.

120 **Definition of disease onset and onset site**

121 Disease onset was defined as the time point when the subject felt weakness of any body part.

122 Subjects with ALS were classified into two groups, limb-onset type and bulbar-onset type,
123 according to the site where they first felt weakness. The limb-onset type included subjects who
124 felt weakness in either of the four extremities first, while the bulbar-onset type included subjects
125 who felt dysarthria or dysphagia first. There were no patients with respiratory-onset type ALS.

126 **Outcome measures**

127 Disease severity was assessed with the Japanese version of the revised ALS Functional Rating
128 Scale (ALSFRS-R), a validated questionnaire-based functional rating scale for ALS [21, 22].

129 Venous blood samples were collected in the supine position after more than 12 h of fasting and
130 just after waking up during hospitalization. At the outpatient clinic, they were collected while the
131 subjects were in the sitting or supine position after more than 12 h of fasting. We measured the
132 following serological indices: CK, creatinine, cystatin C, albumin, blood glucose, hemoglobin
133 A1c, low-density lipoprotein cholesterol (LDL-C), high-density lipoprotein cholesterol (HDL-C),
134 aspartate aminotransferase (AST), alanine aminotransferase, gamma-glutamyl transpeptidase,
135 alkaline phosphatase (ALP), total bilirubin (T-bil), lactate dehydrogenase, uric acid, uric nitrate,
136 white blood cell count, hemoglobin, and platelet count. The levels of serum CK were log
137 transformed before analysis as they had a non-normal distribution.

138 Body composition was assessed with DXA using fan-beam technology (Discovery A; Hologic,
139 Inc., Bedford, MA). DXA has been utilized frequently to evaluate body composition including
140 bone mineral content, fat mass, and lean body mass [23]. We calculated the sum of
141 appendicular lean soft tissue (ALST) mass as an index of skeletal muscle mass [24].

142 **ALS model animal**

143 Transgenic mice overexpressing the human *SOD1* gene carrying the G93A mutation, *SOD1*^{G93A}
144 mice, were purchased from the Jackson Laboratory (Bar Harbor, ME) and maintained as
145 hemizygotes by mating transgenic males with B6/SJLF1 females [25]. Mutant male mice and
146 their male littermate were sacrificed, and their venous blood was collected by puncturing the
147 right ventricle using a 26G needle under pentobarbital anesthesia at the age of 5, 9, and 15
148 weeks: the pre-onset, peri-onset, and symptomatic stage, respectively [26]. Serum CK and
149 creatinine levels were measured with L-Type CK (Wako, Japan) and an enzymatic method,
150 respectively, at Oriental Yeast Co., Ltd. (Tokyo, Japan). Hemolyzed samples were excluded

151 from the analysis. All of the animal experiments were performed in accordance with the National
152 Institutes of Health Guide for the Care and Use of Laboratory Animals and under the approval of
153 the Nagoya University Animal Experiment Committee (No. 29170).

154 **Quantitative real-time PCR of murine muscle**

155 At the age of 5, 9, and 15 weeks, gastrocnemius muscles from mutant SOD1^{G93A} mice and their
156 littermates were dissected and snap-frozen with powdered CO₂ in acetone. Total RNA was
157 extracted from mouse skeletal muscles using TRIzol and a PureLink™ RNA Mini Kit (Invitrogen,
158 Carlsbad, CA). The extracted RNA was reverse-transcribed into first-strand cDNA with ReverTra
159 Ace reverse transcriptase (TOYOBO Co., Osaka, Japan). Quantitative real-time PCR was
160 performed using KOD SYBR® qPCR Mix (TOYOBO Co., Osaka, Japan), and the product was
161 detected using the CFX96™ real-time system (Bio-Rad Laboratories, Hercules, CA). The
162 reaction conditions were 98°C for 2 min, followed by 40 cycles of 10 s at 98°C, 10 s at 60°C,
163 and 30 s at 68°C. The expression levels of acetylcholine receptor subunit γ (*Chrn γ*) and muscle-
164 associated receptor tyrosine kinase (*Musk*) in gastrocnemius muscles were measured as
165 markers of muscle denervation [27]. The expression of glyceraldehyde-3-phosphate
166 dehydrogenase (*Gapdh*) was also quantified and used as an internal standard control. The
167 primers used were 5'-AGCCTCCCCAGCCATCCAGG-3' and 5'-
168 AGCCTCCCCAGCCATCCAGG-3' for *Chrn γ* , 5'-ATCACCACGCCTCTTGAAAC-3' and 5'-
169 TGTCTTCCACGCTCAGAATG-3' for *Musk* [27], and 5'-GAATTTGCCGTGAGTGGAGT-3' and
170 5'-CGTCCCGTAGACAAAATGGT-3' for *Gapdh*.

171 **Statistical analysis**

172 Linear mixed models with unstructured correlation and random intercepts were applied to
173 estimate the average trajectories of each biomarker [28]. Polynomial basis functions were
174 included to incorporate quadratic smoothing. Estimated values and 95% confidence intervals
175 are shown from 0 to +50 months relative to symptom onset at 5-months intervals in subjects
176 with ALS.

177 The chi-square test, unpaired t-test, and Mann-Whitney U test were used for the comparison of
178 variables between two groups, and Pearson's correlation coefficient was used for analyzing
179 correlations among parameters. A partial correlation was performed to determine the
180 relationship between ALSFRS-R and log CK while controlling for age, disease duration, and
181 ALST mass. $P < 0.05$ was considered to be significant, and correlation coefficients (r) were
182 interpreted as follows; greater than 0.8 was very strong, 0.6-0.8 was moderately strong and 0.3-

183 0.5 was fair [29]. All data are presented as the mean \pm standard deviation unless stated
184 otherwise.

185 Statistical Package for the Social Sciences 25.0J software (IBM Japan, Tokyo, Japan) and SAS
186 version 9.4 (SAS Institute, Inc., Cary, NC) were used to perform all statistical analyses.

187

188 **Results**

189 **Participants**

190 A total of 81 subjects with ALS and 20 age- and sex-matched healthy controls were recruited.

191 As a result, we analyzed the longitudinal data from 39 patients with sporadic ALS and 20
192 healthy controls. Four subjects with ALS could not undergo longitudinal evaluation of DXA
193 although they underwent other longitudinal evaluation including blood test. The data prior to the
194 initial evaluation from 8 subjects with ALS were also analyzed retrospectively (Fig 1).

195 **Baseline characteristics**

196 Mean age at the initial evaluation, sex ratio, and body mass index were equivalent between
197 both groups (Table 1). The mean disease duration was 12.6 ± 5.3 months, indicating that the
198 subjects with ALS in our study were early cases. The clinical backgrounds of the limb-onset and
199 bulbar-onset groups were equivalent.

200 **Blood tests and body composition**

201 The baseline levels of serum creatinine and ALST mass, which reportedly correlate with whole
202 muscle mass [24, 30], were lower in subjects with ALS than in healthy controls. In contrast, the
203 baseline levels of serum CK were higher in subjects with ALS (Table 2). Both male and female
204 subjects had similar results (Supplemental table 1). As for the onset sites of ALS, the baseline
205 levels of CK tended to be higher and those of creatinine were lower in limb-onset type ALS
206 compared to the bulbar-onset group (CK, 262.5 ± 257.2 vs. 135.6 ± 79.7 U/L, respectively, $p =$
207 0.105 ; creatinine, 0.62 ± 0.15 vs. 0.80 ± 0.27 mg/dL, respectively, $p = 0.013$).

208 We followed up the subjects with ALS and healthy controls for 8.2 ± 4.2 months and 11.4 ± 3.8
209 months, respectively. The annualized differences of those parameters between the initial and
210 final evaluation are shown in Table 2. The levels of CK, creatinine, and ALST mass decreased
211 longitudinally in subjects with ALS, although they rarely changed in healthy controls. These
212 changes were also common in both male and female subjects (Supplemental table 1). These
213 results suggest that ALST mass in DXA could be a biomarker reflecting muscular atrophy in ALS
214 patients. In addition, the levels of LDL-C, AST, ALP, and T-bil increased longitudinally in subjects

215 with ALS, although they rarely changed in healthy controls (Table 2).

216 **Correlation between muscle-related biomarkers and clinical indices**

217 ALSFRS-R was weakly correlated with log serum CK, but not with ALST mass or serum
218 creatinine, at the initial evaluation (Fig 2A–C, Supplementary Table 2). The correlation between
219 log CK and ALSFRS-R at baseline was still significant after being adjusted for age, disease
220 duration, and ALST mass (partial correlation coefficient = 0.453, $p = 0.006$). Furthermore, the
221 annualized differences (shown as delta) of all of these muscle-related markers were correlated
222 with those of ALSFRS-R (Fig 2D–F). However, log serum CK, ALST mass, and serum creatinine
223 levels at the initial evaluation were not correlated with the longitudinal change of motor function
224 (data not shown). These results suggest that serum CK levels reflect the motor function of each
225 individual, although they do not indicate the prognosis of ALS.

226 **Estimation of muscle-related biomarkers at the early stage of ALS**

227 Muscle-related biomarkers at early ALS stages and average trajectories were estimated using a
228 linear mixed model with quadratic smoothing (Fig 3, Table 3). The estimated values of serum
229 CK at early ALS stages were higher in ALS subjects than in controls, though estimated serum
230 creatinine and ALST mass were equivalent between both groups in this statistical model. These
231 results suggest that, at the early stage of ALS, serum CK levels are elevated, while there
232 appears to be little muscular atrophy.

233 **Retrospective analysis of serum CK before disease onset**

234 For 8 male subjects with sporadic ALS, their serum CK levels before clinical onset had been
235 recorded as a part of medical practice independent of ALS. In these subjects, the levels of
236 serum CK started to be elevated before onset, reached their maximum approximately around
237 onset, and then decreased after onset (Fig 4).

238 **Muscle-related biomarkers in an animal model of ALS**

239 The results of the clinical part of the present study led us to conduct biomarker analysis in an
240 animal model of ALS. We examined the changes of muscle-related serum markers, CK and
241 creatinine, in SOD1^{G93A} transgenic mice. We also measured the mRNA levels of *Chrn*g and
242 *Musk* in the gastrocnemius muscles of the mice, as they are reportedly elevated with
243 denervation [27]. At week 5, the levels of all indices were equivalent between mutant SOD1
244 mice and their wild-type littermates (Fig 5A–D). At week 9, the timing of onset, serum CK levels
245 were elevated and *Chrn*g mRNA expression was up-regulated significantly in mutant SOD1
246 mice compared with wild-type mice (Fig 5A and C). Although not statistically significant, *Musk*

247 was also up-regulated in SOD1^{G93A} transgenic mice at week 9 (Fig 5D). At the symptomatic
248 stage, week 15, serum CK levels were decreased in mutant SOD1 mice, despite the continued
249 up-regulation of *Chrng* mRNA (Fig 5A and C). These results suggest that serum CK starts to be
250 elevated at a very early stage when muscle denervation emerges, and declines with disease
251 progression, consistent with the results of our clinical study.

252

253 **Discussion**

254 In the present study, we found that the estimated level of serum CK was elevated at the onset
255 of motor symptoms in patients with sporadic ALS. Our study also indicated that other muscle-
256 related markers, i.e., serum creatinine and ALST mass in DXA, were not substantially altered at
257 the early stage, although they declined with disease progression during the symptomatic phase
258 of the disease. These results indicate that serum CK is a potential biomarker that reflects the
259 progression of ALS at the early stage.

260 Several studies have documented the elevation of serum CK in ALS [31–37]. As for the
261 pathophysiology underlying this phenomenon, an association between active denervation and
262 the elevation of serum CK was demonstrated in a previous electromyographic study [38].
263 Increased serum CK levels are associated with muscle cramp in patients with ALS [33] and
264 Guillain-Barre syndrome [39]. Furthermore, denervation reportedly induces membrane instability
265 in muscle tissue and leakage of CK into blood *in vivo* [40]. Taken together, the elevation of
266 serum CK at the early stage in ALS appears to be caused by membrane instability or the
267 destruction of muscle tissue due to the denervation and hyperexcitability of motor neurons.

268 We also revealed that mutant SOD1 mice had increased serum CK levels, together with up-
269 regulation of *Chrng*, at the onset of motor dysfunction. *Chrng* encodes the gamma subunit of the
270 acetylcholine receptor protein, which is up-regulated upon muscle denervation, together with
271 *Musk*. Given a previous report on denervation and compensative reinnervation in skeletal
272 muscle of mutant SOD1 mice at their preclinical stage [41], our findings suggest that the
273 elevated levels of CK likely reflect denervation at the preclinical stage of ALS.

274 However, it is also possible that other factors are involved in the elevation of serum CK.
275 Several reports have suggested that serum CK can be elevated by energetic compensatory
276 changes of muscle metabolism, as CK is an enzyme that phosphorylates creatine for the
277 contraction of muscle fibers [30, 36]. The level of serum CK is also influenced by physical
278 activity [42]. Although ALS is classified as a motor neuron disease, a primary change of muscle

279 is also suggested [43]. For example, microRNA-206, which regulates histone deacetylase 4
280 (HDAC4), is up-regulated in skeletal muscle and has a protective effect on disease progression
281 in a mouse model of ALS [44]. The relative mRNA expression level of HDAC4 in skeletal muscle
282 of ALS patients reportedly correlates with disease progression [45]. Alternatively, the elevation
283 of CK might reflect the difference in the proportion of type 1 and 2 muscle fibers affected. In
284 mutant SOD1 mice, the selective vulnerability of fast-fatigable motor neurons, which innervate
285 type 2b muscle fibers, is observed in the pre-symptomatic stage [41]. Given that the enzyme
286 activity of CK is higher in type 2 fibers [46], the proportion of affected myofiber types might be
287 another determinant of CK elevation and motor function at the early stage of ALS.

288 The decrease of serum CK levels with disease progression both in patients and animals in the
289 present study appears to be associated with the loss of muscle volume, given the progressive
290 decline in serum creatinine levels and ALST mass in DXA of sporadic ALS patients. Serum
291 creatinine levels and ALST in DXA are both regarded as clinical biomarkers for the skeletal
292 muscle mass of patients with neuromuscular disorders including ALS [24]. However, it is of note
293 that the estimated values of these indices were not decreased at the onset of sporadic ALS,
294 suggesting that muscle volume is not an appropriate biomarker at the early stage, although they
295 can be used to monitor disease progression at the symptomatic stage of ALS. This view is in
296 accordance with a previous observation that there is little muscle atrophy at disease onset when
297 motor neuron loss is supposed to have already begun [43].

298 There are some reports describing a change of lipid metabolism in ALS [20, 47, 48]. A
299 preclinical study of Swedish ALS patients demonstrated elevation of LDL-C and HDL-C 10
300 years before the disease onset, being more conspicuous in female [20]. However, our cohort
301 study did not show a change of HDL-C or LDL-C in ALS patients at the baseline, although
302 prospective evaluation showed a longitudinal increase in LDL-C levels. The disagreement
303 between our study and previous report may be explained by several factors. Our study
304 population was relatively male dominant. The level of LDL-C in our study tended to be higher in
305 14 female patients than in the male patients (data not shown), and this might have influenced
306 the results. Furthermore, a difference in genetic background might be another explanation for
307 the discrepancy of cholesterol, considering that the proportion of sporadic ALS patients who
308 carry the hexanucleotide repeat expansion of C9orf72 is 0.4% in Japan, which is rather small
309 compared to patients in Western countries (3–21%) [49]. As for the longitudinal elevation of
310 LDL-C, it was accompanied by longitudinal elevations of AST, ALP, and T-bil. This might reflect

311 dyslipidemia and steatosis of liver, both of which have been observed during progression of ALS
312 [47, 48] . The present study has several limitations. First, as the number of subjects with ALS in
313 the present study was small, selection bias could not be excluded. Secondly, the estimation of
314 values was based on a statistical model, even though we confirmed that estimation using
315 retrospective data from patients was in agreement with the animal experimental data. Moreover,
316 we did not analyze comprehensive changes of metabolism in ALS patients. More detailed
317 analysis should be performed focused on creatine and its related metabolism to discuss the
318 changes of creatinine and CK. A further large-scale study is required to assess the preclinical
319 changes in subjects with ALS.

320

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- 440

441 **Figure legends**

442 **Figure 1. Flowchart of study population enrollment**

443 Flowchart describing the enrollment and exclusion of the study population. Data before the
444 onset were retrospectively analyzed in 4 subjects without follow-up data^a and 4 subjects on
445 whom longitudinally analysis was performed^b. Longitudinal data of body compositions was not
446 available in 4 subjects with ALS.

447 ALS, amyotrophic lateral sclerosis; DXA, dual X-ray absorptiometry.

448

449 **Figure 2. Correlation between ALSFRS-R and muscle-related biomarkers**

450 A–C. Scatter plots of ALSFRS-R and muscle related biomarkers [A, logCK (n=39); B, ALST
451 mass (n=35); C, Cr (n=39)] at the initial evaluation. D–E. Scatter plots of the annualized
452 differences of ALSFRS-R and muscle related biomarkers [D, Δ logCK (n=39); E, Δ ALST mass
453 (n=35); F, Δ Cr (n=39)]. Significant correlation coefficients and p-values are annotated.

454 ALS, amyotrophic lateral sclerosis; ALSFRS-R, revised ALS Functional Rating Scale; ALST,
455 appendicular lean soft tissue; CK, creatine kinase; Cr; creatinine; Δ (delta) is the annualized
456 difference of each parameter between the initial and final evaluation.

457

458 **Figure 3. Raw data and estimated average trajectories of muscle-related biomarkers**

459 A–C. Raw data of muscle-related biomarker in subjects with ALS [A, LogCK (n=39); B, ALST
460 mass (n=35); C, Cr (n=39)]. D–F. Estimated average trajectories and their confidence intervals
461 in subjects with ALS. G–L. Raw data and average trajectories in HCs (n=20). The date of the
462 initial evaluation of HCs was set at the mean disease duration (12.6 months) of subjects with
463 ALS. M–O. Merged average trajectories of both groups. Blue lines indicate the data of the
464 subjects for whom retrospective data of serum CK before onset was also analyzed (Patients
465 ALS1, 2, 3, and 4 in Fig 4). Red lines and red dotted lines demonstrate average trajectories and
466 their 95% confidence intervals in subjects with ALS. Black trajectories and black dotted lines
467 demonstrate average trajectories and their 95% confidence intervals of HCs.

468 ALS, amyotrophic lateral sclerosis; ALST, appendicular lean soft tissue; CK, creatine kinase; Cr,
469 creatinine; HC, healthy control; mo, months.

470

471 **Figure 4. Retrospective data of serum CK levels in 8 subjects with ALS**

472 Serum CK levels in 8 subjects with sporadic ALS, for whom serum CK levels had been
473 measured for the purpose of medical practice other than ALS. Age, sex, and the site of onset
474 were described above each graph. Vertical dotted lines demonstrate the timing of the first
475 symptom of weakness and arrow heads demonstrate the timing of the initial evaluation.
476 ALS, amyotrophic lateral sclerosis; CK, creatine kinase.

477

478 **Figure 5. Serum CK, creatinine, and mRNA expression of denervation markers in**
479 **SOD1^{G93A} transgenic mice and their littermates**

480 Serum CK (A), serum creatinine (B), and *Chrn*g (C) and *Musk* (D) mRNA levels in the
481 gastrocnemius muscles (wild-type, male, n = 5; SOD1^{G93A}, male, n = 5). *p < 0.05. Data are
482 presented as the mean ± standard error.

483 CK, creatine kinase; Cr, creatinine; *Chrn*g, acetylcholine receptor subunit γ ; *Gapdh*,
484 glyceraldehyde-3-phosphate dehydrogenase; mSOD1, SOD1^{G93A} transgenic mice; *Musk*,
485 muscle-associated receptor tyrosine kinase; N.S., not significant; WT, wild-type mice.

486

487 **Table 1. Baseline characteristics**

488

	Total ALS	HC	p-value	Bulbar	Limb	p-value
	(n = 39)	(n = 20)		(n = 12)	(n = 27)	
Sex (male%)	64.1	65	0.946	66.7	63.0	0.824
Age at the first evaluation (years)	66.4 ± 7.3	64.4 ± 6.9	0.314	69.6 ± 5.8	65.0 ± 7.6	0.071
Disease duration (months)	12.6 ± 5.3	N.A.	N.A.	12.0 ± 5.8	12.9 ± 5.2	0.788
Height (cm)	160.4 ± 7.9	162.9 ± 8.9	0.293	158.5 ± 7.3	161.3 ± 8.2	0.320
Body weight (kg)	56.7 ± 10.9	62.2 ± 11.0	0.071	54.6 ± 9.3	57.6 ± 11.6	0.428
Body mass index	21.9 ± 3.0	23.3 ± 2.8	0.081	21.6 ± 2.6	22.0 ± 3.1	0.685
ALSFRS-R	40.9 ± 4.7	47.8 ± 0.3	<0.001	40.5 ± 4.8	41.1 ± 4.8	0.730

489 ALS, amyotrophic lateral sclerosis; ALSFRS-R, revised Amyotrophic Lateral Sclerosis

490 Functional Rating Scale; Bulbar, bulbar-onset ALS; HC, healthy control; Limb, limb-onset ALS;

491 N.A., not available. Data represent mean ± standard deviation.

492

493 **Table 2. Baseline and changes of blood tests and body composition**

	Baseline			Change at 48 weeks		
	ALS	HC	p-value	ALS	HC	p-value
Blood test	ALS (n = 39)	HC (n = 20)		ALS (n = 39)	HC (n = 20)	
CK (U/L)	223.5 ± 225.0	107.4 ± 53.5	0.004	-77.0 ± 136.9	2.2 ± 49.7	0.002
log CK	2.21 ± 0.34	1.99 ± 0.20	0.008	-0.19 ± 0.31	0.004 ± 0.14	0.001
Cr (mg/dL)	0.68 ± 0.21	0.77 ± 0.16	0.100	-0.09 ± 0.14	0.02 ± 0.08	<0.001
CysC (mg/L)	0.96 ± 0.20	0.90 ± 0.14	0.221	0.09 ± 0.16	0.01 ± 0.08	0.017
UN (mg/dL)	15.8 ± 4.6	13.4 ± 2.2	0.010	0.36 ± 8.31	0.26 ± 2.60	0.941
UA (mg/dL)	5.1 ± 1.3	5.5 ± 1.0	0.328	-0.69 ± 1.53	0.28 ± 1.28	0.012
AST (U/L)	24.3 ± 7.3	25.6 ± 6.3	0.502	7.4 ± 24.7	-1.0 ± 4.3	0.044
ALT (U/L)	22.1 ± 10.8	21.1 ± 7.8	0.695	5.4 ± 27.7	-1.0 ± 6.4	0.178
T-Bil (mg/dL)	0.87 ± 0.36	0.92 ± 0.42	0.688	0.25 ± 0.49	-0.31 ± 0.22	0.001
ALP (U/L)	209.2 ± 61.6	242.1 ± 59.8	0.055	29.6 ± 80.9	-30.8 ± 45.9	0.003
γ-GTP (U/L)	34.2 ± 26.6	56.2 ± 57.8	0.120	17.7 ± -10.6	-10.6 ± 71.0	0.159
TP (g/dL)	7.0 ± 0.4	7.2 ± 0.3	0.065	0.17 ± 0.96	-0.09 ± 0.76	0.291
Alb (g/dL)	4.2 ± 0.3	4.4 ± 0.3	0.077	0.10 ± 0.81	-0.09 ± 0.43	0.354
LDH (U/L)	210.7 ± 42.8	185.4 ± 34.7	0.026	-19.8 ± 49.5	-15.0 ± 33.9	0.699
LDL-C (mg/dL)	119.4 ± 36.6	117.3 ± 27.9	0.822	23.0 ± 57.9	1.2 ± 19.7	0.043
HDL-C (mg/dL)	54.6 ± 15.9	59.7 ± 18.2	0.282	4.0 ± 18.3	0.85 ± 7.9	0.365
HbA1c (%)	5.9 ± 0.8	5.9 ± 0.4	0.935	-1.4 ± 1.9	0.10 ± 0.25	<0.001
WBC (10³/μL)	5.7 ± 2.1	5.8 ± 1.7	0.910	0.9 ± 3.0	-0.18 ± 1.36	0.072
Hb (g/dL)	14.6 ± 4.9	14.6 ± 1.4	0.970	-1.6 ± 10.1	-0.3 ± 1.1	0.571
Plt (10³/μL)	232.8 ± 69.5	222.3 ± 57.0	0.561	8.0 ± 75.6	-5.7 ± 26.1	0.313
Body composition	ALS (n = 39)^a	HC (n = 20)		ALS (n = 35)	HC (n = 20)	
ALST mass (kg)	16.65 ± 4.01	19.20 ± 4.5	0.030	-2.56 ± 2.27	-0.42 ± 1.22	<0.001
BMC (kg)	1.90 ± 0.39	1.94 ± 0.42	0.716	-0.09 ± 0.12	0.02 ± 0.12	0.002
Total fat mass(kg)	14.25 ± 4.12	15.21 ± 3.79	0.385	-0.97 ± 4.49	0.46 ± 1.65	0.094

494

495 Change at 48 weeks shows the annualized difference of each parameter between the initial and

496 final evaluation.

497 ^aLongitudinal data of body compositions was not available in 4 subjects with ALS.
498 Alb, albumin; ALP, alkaline phosphatase; ALS, amyotrophic lateral sclerosis; ALST,
499 appendicular lean soft tissue; ALT, alanine aminotransferase; AST, aspartate aminotransferase;
500 BMC, bone mineral content; CK, creatine kinase; Cr, creatinine; CysC, cystatin C; γ -GTP,
501 gamma glutamyl transpeptidase; Hb, hemoglobin; HbA1c, hemoglobin A1c; HC, healthy control;
502 HDL-C, high-density lipoprotein cholesterol; LDH, lactate dehydrogenase; LDL, low-density
503 lipoprotein cholesterol; Plt, platelet count; T-Bil, total bilirubin; TP, total protein; UA, uric acid;
504 UN, uric nitrate; WBC, white blood cell count. Data represent mean \pm standard deviation.

505 **Table 3. Estimated values of muscle-related biomarkers at onset**

	ALS		HC (n = 20)	
	Estimation	95% CI	Estimation	95% CI
Log CK	2.34 (n =39)	1.68–3.00	1.97	1.57–2.37
ALST mass (kg)	19.91 (n = 35) ^a	11.73–28.09	19.32	9.76–28.89
Cr (mg/dL)	0.80 (n = 39)	0.38–1.22	0.75	0.41–1.09

506

507 ^aLongitudinal data of body compositions was not available in 4 subjects with ALS.

508 The values were calculated using a linear mixed model with unstructured correlation and

509 random intercepts.

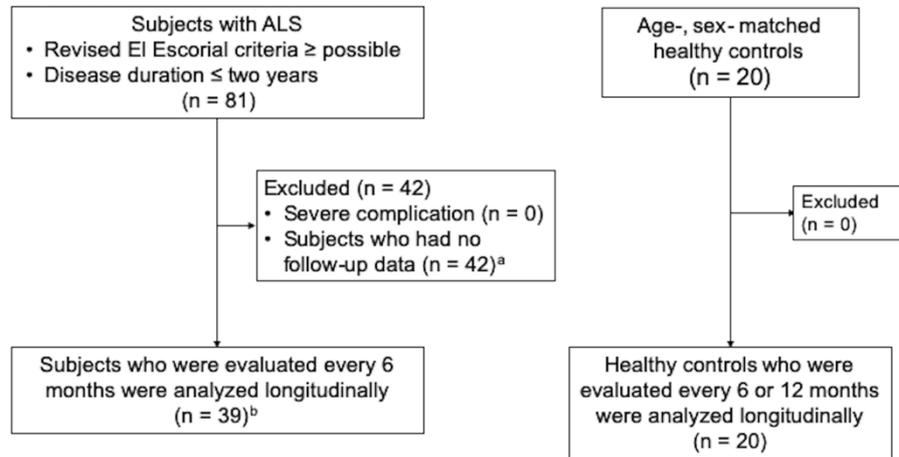
510 ALS, amyotrophic lateral sclerosis; ALST, appendicular lean soft tissue; CI, confidence interval;

511 CK, creatine kinase; Cr, creatinine; HC, healthy controls

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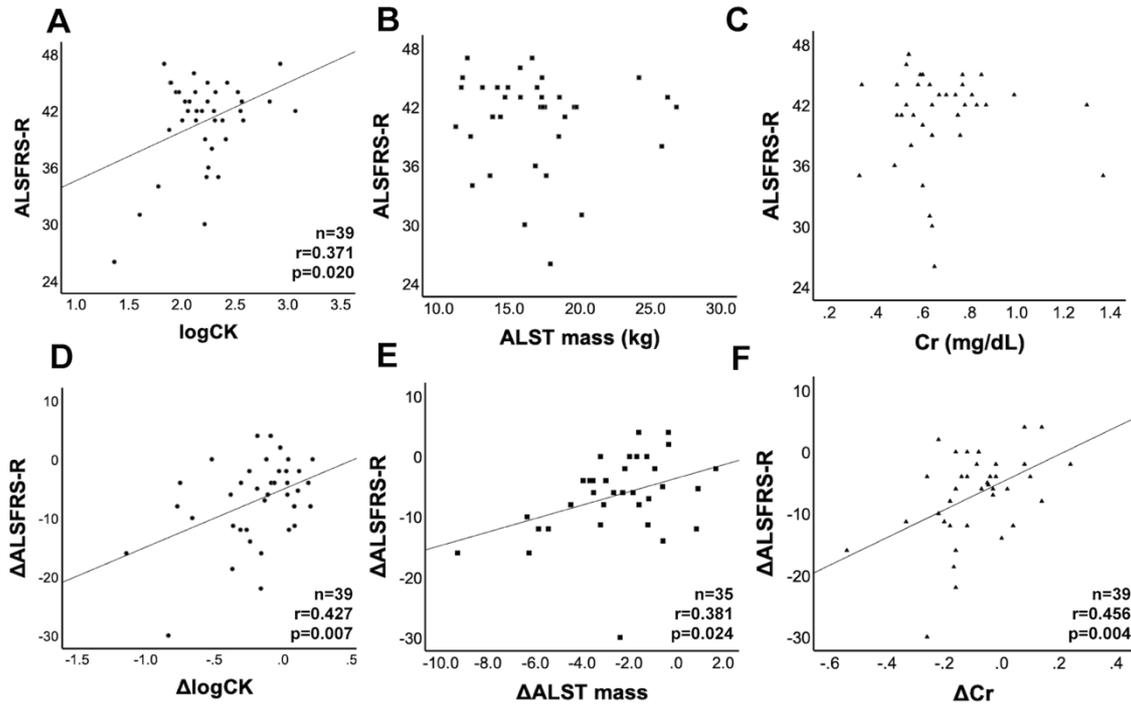
514 Figure 1



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516

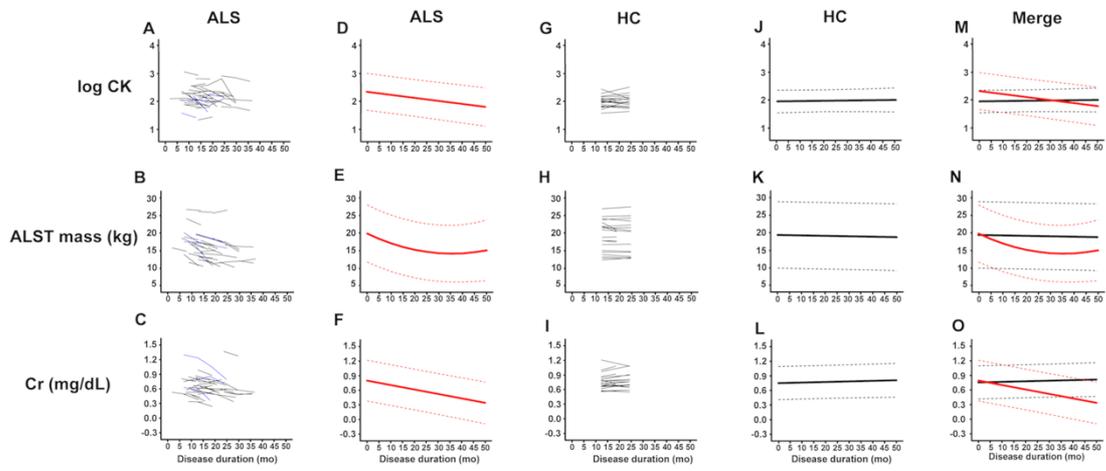
517 Figure 2



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520 Figure 3

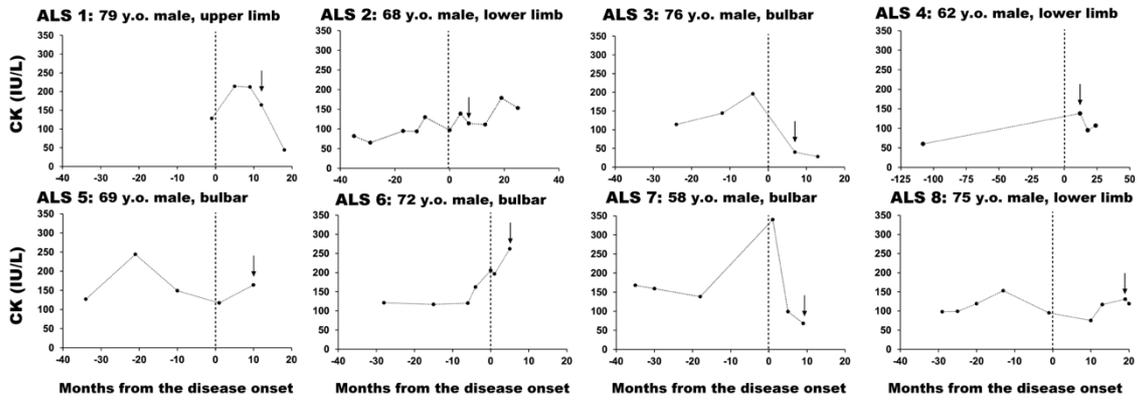


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524 Figure 4



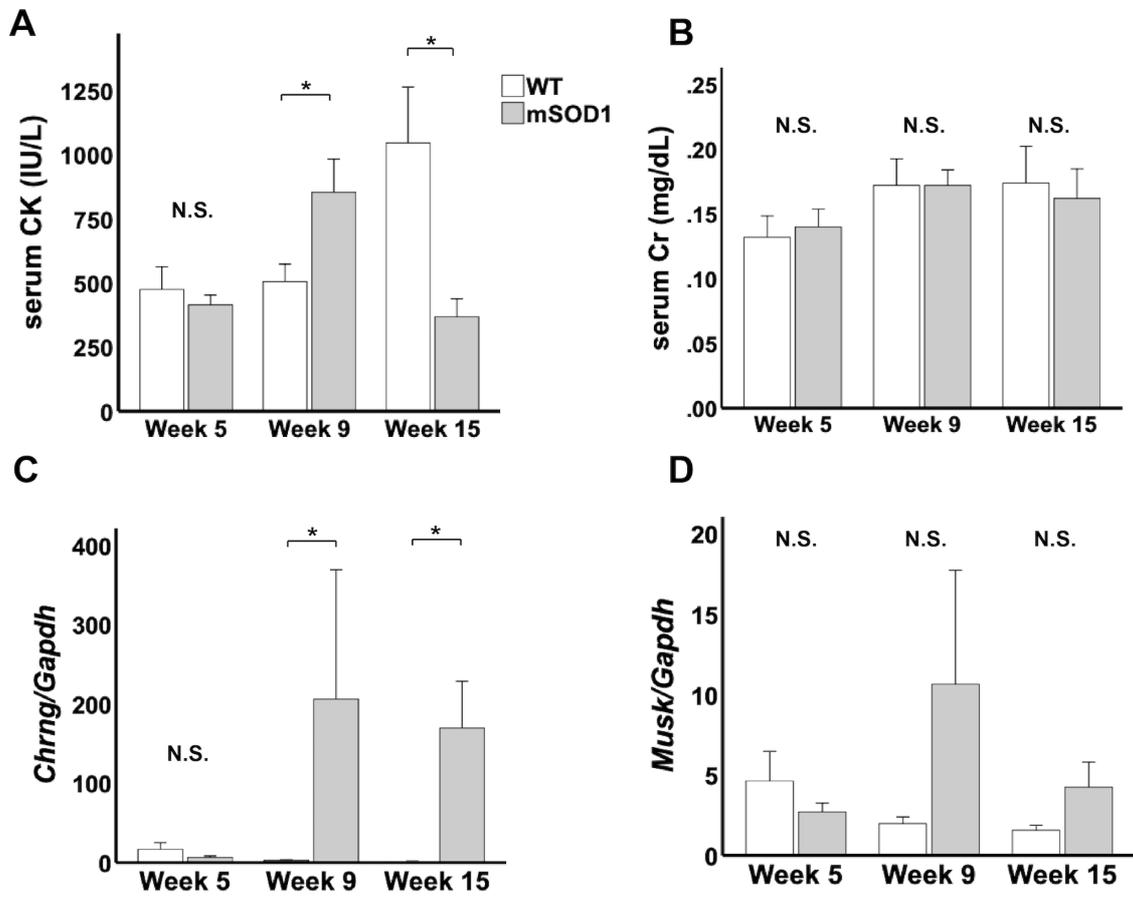
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529 Figure 5



530