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Assessment of metal concentrations in the SOD1^{G93A} mouse model of amyotrophic lateral sclerosis and its potential role in muscular denervation, with particular focus on muscle tissue

T. Gabriel Enge University of Wollongong, tge571@uowmail.edu.au

Heath Ecroyd University of Wollongong, heathe@uow.edu.au

Dianne F. Jolley University of Wollongong, djolley@uow.edu.au

Justin J. Yerbury University of Wollongong, jyerbury@uow.edu.au

Bernadett Kalmar University College London

See next page for additional authors

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Abstract

Background: Amyotrophic lateral sclerosis (ALS) is among the most common of the motor neuron diseases, and arguably the most devastating. During the course of this fatal neurodegenerative disorder, motor neurons undergo progressive degeneration. The currently best-understood animal models of ALS are based on the over-expression of mutant isoforms of Cu/Zn superoxide dismutase 1 (SOD1); these indicate that there is a perturbation in metal homeostasis with disease progression. Copper metabolism in particular is affected in the central nervous system (CNS) and muscle tissue.

Methods: This present study assessed previously published and newly gathered concentrations of transition metals (Cu, Zn, Fe and Se) in CNS (brain and spinal cord) and non-CNS (liver, intestine, heart and muscle) tissues from transgenic mice over-expressing the G93A mutant SOD1 isoform (SOD1 ^{G93A}), transgenic mice over-expressing wildtype SOD1 (SOD1^{WT}) and non-transgenic controls.

Results: Cu accumulates in non-CNS tissues at pre-symptomatic stages in SOD1^{G93A} tissues. This accumulation represents a potentially pathological feature that cannot solely be explained by the over-expression of mSOD1. As a result of the lack of Cu uptake into the CNS there may be a deficiency of Cu for the over-expressed mutant SOD1 in these tissues. Elevated Cu concentrations in muscle tissue also preceded the onset of symptoms and were found to be pathological and not be the result of SOD1 over-expression.

Conclusions: It is hypothesized that the observed Cu accumulations may represent a pathologic feature of ALS, which may actively contribute to axonal retraction leading to muscular denervation, and possibly significantly contributing to disease pathology. Therefore, it is proposed that the *toxic-gain-of-function* and *dying-back* hypotheses to explain the molecular drivers of ALS may not be separate, individual processes; rather our data suggests that they are parallel processes.

Disciplines

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T. Gabriel Enge^{1*}, Heath Ecroyd², Dianne F. Jolley³, Justin J. Yerbury², Bernadett Kalmar⁴, Anthony Dosseto¹

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¹Wollongong Isotope Geochronology Laboratory and School of Earth and
 Environmental Sciences, University of Wollongong, Australia

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²Illawarra Health and Medical Research Institute and School of Biological Sciences,
 University of Wollongong, Australia

14

³Center for Medical and Molecular Bioscience and School of Chemistry, University
 of Wollongong, Australia

- ⁴Sobell Department of Motor Neuroscience and Movement Disorders, Institute of
 Neurology, University College London, UK
- 20
- 21
- 22 Corresponding Author:

23 T. Gabriel Enge

- 24 Email: tge571@uowmail.edu.au
- 25 Wollongong Isotope Geochronology Laboratory
- 26 School of Earth and Environmental Sciences
- 27 University of Wollongong
- 28 Wollongong, NSW, 2522
- 29 Australia
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33 Abstract

34 **Background:** Amyotrophic lateral sclerosis (ALS) is among the most common of the 35 motor neuron diseases, and arguably the most devastating. During the course of this 36 fatal neurodegenerative disorder, motor neurons undergo progressive degeneration. 37 The currently best-understood animal models of ALS are based on the over-38 expression of mutant isoforms of Cu/Zn superoxide dismutase 1 (SOD1); these 39 indicate that there is a perturbation in metal homeostasis with disease progression. 40 Copper metabolism in particular is affected in the central nervous system (CNS) and 41 muscle tissue.

42 **Methods:** This present study assessed previously published and newly gathered 43 concentrations of transition metals (Cu, Zn, Fe and Se) in CNS (brain and spinal cord) 44 and non-CNS (liver, intestine, heart and muscle) tissues from transgenic mice over-45 expressing the G93A mutant SOD1 isoform (SOD1^{G93A}), transgenic mice over-46 expressing wildtype SOD1 (SOD1^{WT}) and non-transgenic controls.

47 Results: Cu accumulates in non-CNS tissues at pre-symptomatic stages in SOD1^{G93A} 48 tissues. This accumulation represents a potentially pathological feature that cannot 49 solely be explained by the over-expression of mSOD1. As a result of the lack of Cu 50 uptake into the CNS there may be a deficiency of Cu for the over-expressed mutant 51 SOD1 in these tissues. Elevated Cu concentrations in muscle tissue also preceded the 52 onset of symptoms and were found to be pathological and not be the result of SOD1 53 over-expression.

54 Conclusions: It is hypothesized that the observed Cu accumulations may represent a 55 pathologic feature of ALS, which may actively contribute to axonal retraction leading 56 to muscular denervation, and possibly significantly contributing to disease pathology. 57 Therefore, it is proposed that the *toxic-gain-of-function* and *dying-back* hypotheses to

explain the molecular drivers of ALS may not be separate, individual processes;rather our data suggests that they are parallel processes.

60

61 **Keywords**: ALS, Copper, Spinal Cord, Brain, Muscle, Distal Motor Neuropathy;

62

63 Introduction

64 Amyotrophic lateral sclerosis (ALS) is a neurodegenerative disease, which leads to 65 selective motor neuron (MN) death, and is ultimately fatal (Redler and Dokholyan, 66 2012; Rowland and Shneider, 2001). The degeneration results in progressive muscular 67 paralysis, affecting mobility, speech, and respiration (Hardiman et al., 2011); most 68 patients die within 5 years from diagnosis (Robberecht and Philips, 2013). Most ALS 69 cases are described as sporadic, but around 5 - 10% are familial (Rowland and 70 Shneider, 2001) and associated with a variety of Mendelian-inherited mutations 71 (Robberecht and Philips, 2013).

72

73 The first gene identified to be mutated in familial ALS encodes Cu/Zn superoxide 74 dismutase 1 (SOD1), accounting for ~20% of familial ALS cases (Rosen et al., 1993). 75 The specific mechanisms leading to the selective degeneration of MNs remains 76 unclear, with a variety having been proposed to play a role, such as oxidative stress, 77 glutamate-mediated excitotoxicity, protein aggregation and transition metal-induced 78 toxicity (Cozzolino et al., 2008). Toxicity induced by mutant SOD1 (mSOD1) is 79 likely to be the result of a toxic-gain-of-function (Bruijn et al., 1998; Gurney et al., 80 1994), driven by the increased destabilization of the protein, which causes it to 81 misfold and aggregate. To-date the most studied mouse model of ALS is the transgenic mouse that overexpresses the SOD1-G93A mutation (SOD1^{G93A}). In this 82

model, human SOD1 harboring the G93A mutation is overexpressed (20-24 fold
higher expression than endogenous murine SOD1) (Gurney et al., 1994). This mouse
model recapitulates many features of human ALS, including axonal and
mitochondrial dysfunction, progressive neuromuscular dysfunction, protein
aggregation and MN loss (Bruijn et al., 1997; Gurney et al., 1994; Ripps et al., 1995).
Results from experimental data suggest that the binding of Cu and Zn by mSOD1 may
be defective (Carri et al., 1994; Eum and Kang, 1999; Hayward et al., 2002).

90

91 While transition metals (e.g. Cu and Zn) are crucial for the function of many metallo-92 proteins (e.g. SOD1), they can be toxic at high concentrations (Gaetke and Chow, 93 2003; Jomova and Valko, 2011; Valko et al., 2005), and their uptake, distribution, 94 storage and excretion are therefore tightly controlled (Hare et al., 2013; Tapiero et al., 95 2003). The role of metals in the pathogenesis of certain neurodegenerative diseases 96 (i.e. Cu, Fe and Al in Alzheimer's disease (Greenough et al., 2013; House et al., 2012; 97 Kaden et al., 2011; Shore et al., 1984)) is becoming evident; however, the particular 98 role of Cu and Zn homeostasis in the pathology of ALS remains unclear (Roberts et 99 al., 2014).

100

Findings in the G93A mouse model of ALS indicate that there is a pathologic perturbation in Cu metabolism. While the accumulation and/or depletion of metals in different tissues reported in the literature are inconsistent, overall there appears to be accumulation in tissues of diseased mice. Tissues associated with the autonomic nervous system (ANS) have been shown to accumulate Cu and Zn (Enge et al., 2017; Hilton et al., 2016), with muscle tissue demonstrating pre-symptomatic increases in concentrations (Enge et al., 2017). Results from central nervous system (CNS) tissues 108 indicate accumulation of Cu in the spinal cord (Li et al., 2006; Tokuda et al., 2015, 109 2014, 2013, 2008) at single time points, as well as over time (Tokuda et al., 2009, 110 2007). Brain tissue showed a similarly varied behavior with studies reporting both 111 elevated concentrations (Hilton et al., 2016), or no change to healthy controls (Enge et 112 al., 2017; Lelie et al., 2011; Li et al., 2006; Tokuda et al., 2009, 2007). These changes 113 in Cu and Zn concentrations have been postulated to play an important role in the 114 toxic-gain-of-function associated with mSOD1 (Borchelt et al., 1994; Hilton et al., 115 2015). Two major strategies to alleviate symptoms of ALS that involve metals have 116 been proposed. These are based on either reducing the general availability of Cu 117 through the application of Cu chelators (i.e. Trietine or *D*-penicillamine), which has been shown to prolong survival of SOD1^{G93A} mice (Hottinger et al., 1997; Nagano et 118 119 al., 2003); or the application of a synthetic Cu-containing therapeutic agent (Cu^{II}-120 atsm) to transport Cu across the blood-brain-barrier, which prolongs survival in SOD1^{G37R} and SOD1^{G93A} mice (Hilton et al., 2017; McAllum et al., 2013; Roberts et 121 122 al., 2014; Soon et al., 2011; Williams et al., 2016).

123

124 In SOD1, Cu and Zn ions are crucial for its function: Zn provides structural stability, 125 while Cu is essential for the protein's catalytic activity. In SOD1 mouse models of 126 ALS, mutant SOD1 accumulates in a Cu-deficient form (Lelie et al., 2011; Roberts et 127 al., 2014; Williams et al., 2016). This partially metalated SOD1 is proposed to lead to 128 aggregation and toxicity (Bruijn et al., 1998; Gurney et al., 1994; Roberts et al., 129 2014). Treatment of mice with the synthetic Cu-containing therapeutic agent Cu^{II}(atsm) converts the partially metalated SOD1 into holo SOD1 through the 130 131 delivery of Cu (Roberts et al., 2014), improving the phenotype significantly (Lelie et

132	al., 2011;	Roberts	et al.,	2014;	Williams	et	al.,	2016).	This	points	towards	a
133	misbalanc	e of bioav	ailabili	ty and d	emand for	Cu	in th	e CNS	(Hilto	n et al.,	2016).	

134

135 The potential role of metals as well as muscle tissue (Gonzalez de Aguilar et al., 136 2008; Luo et al., 2013; Xiao et al., 2015) in ALS pathology has recently gathered 137 further attention. In addition to the involvement of metals in binding to SOD1, recent 138 research has suggested that toxic accumulation of metals in muscle tissue may also 139 play a role in ALS pathology (Enge et al., 2017). With pathological hallmarks that 140 include the destruction of the neuromuscular junction (NMJ) and axonal retraction 141 preceding MN death (Zhou et al., 2015), ALS could be described as a distal motor 142 neuropathy (Enge et al., 2017; Fischer et al., 2004; Frey et al., 2000; Kiernan et al., 143 2011).

144

In this study concentrations of transition metals in transgenic mutant SOD1 mice (G93A) and non-transgenic controls were assessed. Through the longitudinal comparison of diseased and healthy CNS and non-CNS tissues, this research sought to expand current knowledge on the role and presence of transition metals in the pathology of ALS. <u>Metal accumulations in muscle tissue were assessed for their</u> possible contribution to axonal retraction.

151

152 METHODS

153 Samples

154 This study used two types of transgenic over-expression models, SOD1^{G93A} and 155 SOD1^{WT}; only female mice were used in this work. The SOD1^{G93A} mouse model of 156 ALS expressed the G93A mutant form of human SOD1 (B6SJL-

157 Tg(SOD1*G93A)1Gur/J)(Gurney et al., 1994) (Jackson Laboratory, ME, USA), 158 backcrossed onto a black 6 background. Healthy controls for these mice were nontransgenic littermates (NTG^{UOW}). These mice were bred at Australian Bioresources 159 160 (Mossvale, AU) in accordance with the approved University of Wollongong ethics 161 clearance (AE14/28). At ages 30, 60, 90 and 120 (±2) days, capturing the 162 characteristic disease progression (Olsen et al., 2001), mice were euthanized by CO₂ 163 asphysiation and blood collected via puncturing of the left ventricle. The mice were 164 not perfused and following blood collection immediately dissected and tissue samples 165 (brain, spinal cord, liver, intestine, heart and muscle) were snap frozen in liquid 166 nitrogen. Muscle tissue was samples of *Quadriceps femoris* from both legs and 167 intestine samples were 2 cm sections immediately following the stomach.

168

The second type of mouse model used was SOD1^{WT} mice, which overexpress the non 169 170 mutated form of human SOD1 (B6SJL-Tg(SOD1)2Gur/J) (Gurney et al., 1994) (Jackson laboratories). Healthy controls were non-transgenic littermates (NTG^{UCL}). 171 172 Mice were bred at University College London - Biological Services (London, UK) in 173 accordance with the Guide for the Care and Use of Laboratory Animals as adopted by 174 the US National Instituted of Health and under license from the UK Government 175 (Animals and Scientific procedures) Act 1986 (Amended Regulations 2012), 176 following ethical approval from the University College London Institute of 177 Neurology. The samples taken from this mouse model were brain, spinal cord and 178 muscle tissue (Quadriceps femoris) samples. These samples were collected at 60 and 179 90 d to assess if the previously observed increase in Cu concentration in diseased 180 tissue (Enge et al., 2017) is the result of the over expression of the mSOD1 or presents 181 a pathologic feature, independent of mSOD1 over-expression.

182

183 Samples of DORM-2 dogfish muscle, certified reference material of the National
184 Research Council Canada (CRM, NRCC), were used to test the completeness and
185 accuracy of the sample digestion protocol (Enge et al., 2016).

186

187 Analytical techniques

188 Sample digestion

189 Mouse tissues and DORM-2 aliquots were weighed and pre-digested in MARSXpress 190 20 mL PFA vessels in a 2.5:1 mixture of 15 M Ultrapur® HNO₃ (Merck) and 191 Ultrapur[®] 30% H₂O₂ (Merck) for 30 min. Pre-digested samples were completely 192 digested using MARS6 (CEM Corporation, North Carolina, USA) microwave 193 systems. Temperature was ramped to 210 °C over 15 min and held constant for 150 194 min to ensure all organic carbon was driven off as gaseous CO₂. For quality control 195 purposes, one blank (acid only) and two DORM-2 aliquots were added to each 196 digestion batch. Recovery of elements (Cu, Fe, Zn and Se) from the DORM-2 197 certified reference material was used to validate that the digestion of biological 198 samples was complete (Table 1). Selenium concentrations were only determined in 199 liver, intestine and heart tissues.

200

Table 1 – Recoveries of select metals from DORM-2 (CRM, NRCC), in mg kg⁻¹,
during processing of samples

Element	Certified Value	2SD	Measured Value	2SD	Recovery (%)	Ν
Cu	2.34	0.16	2.4	0.4	104	21
Fe	142	10	133	31	94	21
Zn	25.6	2.3	26	10	109	21
Se	1.40	0.09	1.4	0.2	101	8

205 Copper, Fe, Zn and Se concentrations were determined using a Thermo Scientific 206 iCAP-O quadrupole-inductively coupled plasma-mass spectrometer (O-ICP-MS) at 207 the Wollongong Isotope Geochronology Laboratory, University of Wollongong 208 (WIGL, UOW). The concentrations were quantified using a multi-element standard 209 external calibration curve; long-term instrument drift was corrected using a 50 ppb Ga 210 solution as an internal standard. Accuracy of the measurements was assessed through 211 the analysis of the DORM-2 CRM, which yielded recoveries of 104% for Cu, 94% for 212 Fe, 109% for Zn and 101% for Se (Table 1). Total procedure blanks were assessed as 213 <4 ng Cu, <9 ng Fe, <18 ng Zn, and <13 ng Se, and deemed negligible (<0.13% for 214 Cu, Zn and Fe; <2% for Se) compared to the total amount processed (see 215 supplementary tables).

216

217 Statistical Methods

218 Statistical analysis was conducted using the statistical program R v3.4.2 (R Core 219 Team, 2016). Prior to analysis, outliers were removed using the median average 220 deviation (3*mad). Any measurements outside 3*mad were considered to be measurement artifacts or the result of contamination. Transgenic SOD^{WT} muscle 221 222 samples were analysed qualitatively, due to the low sample number. Data were 223 analyzed using linear regression models with disease state and time used as 224 independent variables and Cu, Fe, Zn and Se concentrations, as dependent variables. 225 Likelihood ratio tests were used to determine which independent variables were 226 significant (Barr et al., 2013). The assumptions of the linear regression model were 227 checked (linearity, independence, normality and equality of variance), including 228 normality tests for residuals. A significance level of $\alpha = 0.05$ was used.

229

230 **RESULTS**

231 Peripheral tissues

232 Tissues controlled by the ANS, i.e. the liver, intestine and heart, demonstrate a significant difference in Cu concentration (p < 0.0005) between the SOD1^{G93A} and 233 234 healthy controls (Figure 1A-C), which is evident starting at 30 d, with concentrations 235 in heart and liver increasing over time in the diseased tissues, while they remained 236 relatively constant in the intestine. Zinc concentrations show a similar behaviour to Cu: they vary significantly between SOD1^{G93A} and healthy controls in heart and liver 237 238 tissue (p < 0.0005), and intestine (p = 0.03) (Figure 1D-F) starting at 30 d. In heart 239 and liver, the Zn concentrations increase over time, while concentrations in intestine 240 remain largely constant. In contrast to Cu and Zn, Fe concentrations only vary 241 significantly between the diseased and healthy liver (p = 0.05) and the intestine (p = 0.05)242 0.006) tissue (Figure 2 A-C), and over time in liver (p < 0.0005) and intestine (p =243 0.001). Concerning Se concentrations, only intestine was found to show a significant 244 difference between healthy and diseased tissue (p = 0.04), while Se concentrations in heart (p = 0.002), liver and intestine (p = 0.0005) changed over time (Figure 2D-F). 245

246

247 Muscle tissue (Quadriceps)

As previously shown (Enge et al., 2017), Cu (p < 0.0005), Zn (p < 0.0005) and Fe (p = 0.004) are significantly different between SOD1^{G93A} and NTG^{UOW} mice (Figure 3). Muscle tissues from mice that overexpress the SOD1^{WT} were collected at 60 d and 90 d and tissues from non-transgenic littermates from the same colony were collected at 90 d of age (Figure 4). The concentrations of Cu, Zn and Fe in the SOD^{WT} samples are in agreement with results from muscle tissue collected from non-transgenic mice from a different colony (NTG^{UOW}) (Enge et al., 2017). The metal concentrations in SOD1^{WT} tissues match well these found in NTG^{UOW} non-transgenic controls (Figure 3).

257

258 Central nervous system tissues

It was previously shown that there is no significant difference between brain and spinal cord tissue from SOD1^{G93A} and NTG^{UOW} mice in Cu, Zn and Fe concentrations (Enge et al., 2017). Samples from mice over-expressing SOD1^{WT} and non-transgenic littermates from the same colony (NTG^{UCL}) were collected at 60 and 90 d of age, and are in general agreement with previously presented results of SOD1^{G93A} and NTG^{UOW} mice from a different colony (Supplementary Figure 1).

265

266 **DISCUSSION**

267 Accumulation of Cu and Zn in tissues controlled by the ANS

268 Tissues controlled by the ANS play an as yet unknown role in the development of 269 ALS. The accumulation of Cu in the heart and liver could either be pathological and 270 driven by ALS, or the result of the over-expression of SOD1 in the mouse model. As 271 the mouse model of ALS overexpresses (20 - 24 times) human SOD1 harboring the 272 G93A mutation (Gurney et al., 1994), overexpression of the protein could result in it 273 acting as a Cu sink (Tokuda and Furukawa, 2016). However, if this were the case a uniform offset with regards to the Cu concentration in tissues from SOD1^{G93A} mice 274 275 compared to the non-transgenic mice would be expected (similar to that observed for 276 the intestine where concentrations did not change significantly over time). A lack of 277 increases of SOD1 concentration with time in spinal cord and brain of various mouse 278 models of ALS was observed and attributed to reflecting high-level relative steady-279 states (Jonsson et al., 2006). Though work by Turner et al. (2003) has shown age280 dependent accumulation of hSOD1 in the lumbar spinal cord, the sciatic nerve, and 281 the gastrocnemius muscle. In our work (Enge et al., 2017) it was acknowledged that differences in Cu and Zn concentration between the SOD1^{G93A} and non-transgenic 282 mice in muscle tissue could be due to over-expression of human SOD1. Here, 283 284 observations of a lack of elevated concentrations in Fe and Se (Figure 2) point 285 towards a great contribution of over-expressed SOD1 to the Cu accumulation. This 286 accumulation was also reported in previous work (Tokuda et al., 2013): in spinal cord tissue of SOD1^{WT} mice (over-expression of wild-type SOD1) and SOD1^{G93A} mice Cu 287 concentrations were elevated (up to 3.2-fold) compared to non-transgenic controls. 288 289 Furthermore, Cu concentrations in spinal cord were significantly elevated in SOD1^{G93A} samples compared to SOD1^{WT} samples (Tokuda, 2017). This indicates that 290 at least part, if not all of the accumulation of Cu and Zn in these tissues could be 291 292 attributable to SOD1 over-expression. However, the distinct accumulation of Cu in heart and liver over time (Figure 1) compared to controls may represent disease 293 pathology: previously it was found that in SOD1^{G93A} spinal cord tissue, Cu was bound 294 295 to copper-binding proteins other than SOD1 and this increased over time (Tokuda et 296 al., 2013). Accumulation of Cu in cupro-proteins besides SOD1 may therefore 297 contribute to the observed signal. The observation of a subtle age-dependent increase 298 of hSOD1 in the gastrocnemius muscle (Turner et al., 2003) could though indicate 299 that a large proportion if not all of the observed metal accumulation is a result of 300 SOD1 accumulation with age. The lack of SOD1 concentration measurements here 301 presents a limitation to the conclusions that can be drawn. Without tissue-matched measurements from mice that overexpress SOD1^{WT}, or determination of SOD1 302 concentrations, estimation of how much of the observed increase in Cu concentration 303

is attributable to the over-expression of SOD1 in peripheral system tissues remainsuncertain.

306

307 Zinc concentrations were found to be significantly elevated in all three tissues (liver, 308 heart and intestine) and their concentrations in heart and liver increase over time 309 (Figure 1D-F). Similar to Cu, increased concentration of Zn in heart and liver of SOD1^{G93A} mice are likely to be related to the over-expression of SOD1, as other 310 311 elements (Fe and Se) were not increased (Figure 2). Another factor may also contribute: work using spinal cord tissue from SOD1^{G93A} mice showed an 312 313 accumulation of Zn not only in the zinc-binding site of SOD1, but also suggested an 314 increase of Zn binding in other Zn binding proteins (Tokuda et al., 2013). The 315 increase in Zn concentration was also statistically significant in the intestine; the 316 difference is lower and relatively constant over time.

317

While the accumulation of Cu and Zn in these tissues precede disease pathology and
symptoms, coinciding with initial protein aggregation at 30 d (Gould et al., 2006)
(Table 2), the specific binding of the metal(s) to SOD1 has to be further scrutinized.

322 Table 2 – Typical ALS symptoms and features in transgenic SOD1 mice

Feature/symptom	Average time of onset (d)	Reference
Initial protein aggregation	30	(Gould et al., 2006)
Selective neuromuscular junction degeneration	47	(Fischer et al., 2004)
Maximum running speed reduced	52	(Veldink et al., 2003)
Reduction of blood flow through the spinal cord	60	(Zhong et al., 2008)
Axonal loss prominent	80	(Fischer et al., 2004)
Significant motor neuron loss	100	(Fischer et al., 2004;

		Seki et al., 2007)
Onset of paralysis	100	(Chiu et al., 1995)

323

324 Lack of up-regulation of Cu transport into CNS

325 Previously no increase in Cu and Zn concentrations was found in the CNS (Enge et 326 al., 2017); in this work we show these do increase in non-CNS tissues (Figure 1). This 327 could be the result of the non-uniform expression of mSOD1 in various tissues. Results in humans (Human Protein Atlas, 2017) as well as in the SOD1^{G37R} (Hilton et 328 al., 2016) and SOD1^{G93A} (Gajowiak et al., 2016) mouse models have shown that 329 SOD1 protein expression varies between tissues; SOD1^{G37R} CNS tissues expressed 330 331 much greater amounts of SOD1 compared to peripheral nervous system tissues (Hilton et al., 2016). In tissues from SOD1^{G93A} mice, higher amounts of SOD1 were 332 333 reported in spinal cord compared to liver and muscle (Gastrocnemius) (Gajowiak et 334 al., 2016). Thus, if Cu and Zn concentrations reflected the amount of SOD1 present, 335 it would be expected that CNS tissues would have higher concentrations of Cu and Zn 336 compared to other tissues. The origin for the discrepancy in Cu and Zn concentrations 337 between CNS and non-CNS tissues is therefore more likely to be the response of the 338 blood-CNS barriers. While an increased Cu requirement in the CNS, due to the SOD1 339 over-expression, may not be met (Hilton et al., 2016), it is unclear why Zn 340 concentrations are not elevated.

341

Our results show an accumulation of Cu and other metals in the heart and liver tissue compared to the CNS, as well as compared to non-transgenic controls (Enge et al., 2017; Hilton et al., 2016). Blood of ALS patients was shown to not accumulate Cu and Zn, compared to controls (Garzillo et al., 2014; Kapaki et al., 1997; Nagata et al., 1985; Pamphlett et al., 2001; Roos et al., 2013). This agrees with our previous

findings in the SOD1^{G93A} mouse (Enge et al., 2017). Blood carries a large amount of 347 348 the total body Cu, which can be regarded as a theoretically infinite pool for the 349 individual compartments. The observed increases in concentration in tissues such as heart and liver in the SOD1^{G93A} mice, despite no change in blood (Enge et al., 2017), 350 351 suggests that acquisition of Cu from the liver into the blood does not vary. Overall the 352 increased demand for Cu in the organism due to higher mSOD1 levels in these tissues 353 is satisfied. Copper concentrations in the heart, liver and intestine may therefore serve as an additional proxy for disease in the SOD1^{G93A} mouse model of ALS. 354

355

356 The lack of an increase in Cu concentration in the CNS may reflect a disconnect 357 between the rate of Cu uptake and Cu requirement (Hilton et al., 2016) as a response 358 to the over-expression of mSOD1. Recent results have shown that the amount of Cu in both brain and spinal cord tissue of the SOD1G37R mouse model is 359 360 disproportionately small compared to the amount of SOD1 protein expressed (Hilton 361 et al., 2016). This indicates that as well as a generally slower turnover, Cu transport 362 into the CNS is not up-regulated to satisfy the increased Cu requirement due to 363 mSOD1 over-expression. This effectively 'starves' the over-expressed mSOD1 364 (Figure 4) (Hilton et al., 2016). As a result, the protein could be destabilized, making 365 it aggregation-prone (Lelie et al., 2011). The result could be MN death through direct 366 toxicity of the partially metalated SOD1 (Gil-Bea et al., 2017; Roberts et al., 2014).

367

Pathological accumulation of Cu in muscle tissue may contribute to ALS development
Muscle tissue (Gonzalez de Aguilar et al., 2008; Luo et al., 2013; Xiao et al., 2015),
and other (neighbouring) cell types (Boillée et al., 2006) may play a significant role in
the development of ALS pathology. Distinct and rapid muscle atrophy caused by MN

372 death is a pathologic feature of ALS. Skeletal muscle comprises ~40% of whole body 373 lean mass and, combined with bone, makes up ~50% of total body Cu in humans 374 (Evans, 1973). We previously showed consistently elevated metal concentrations in SOD1^{G93A} muscle tissue compared to samples from non-transgenic animals (Enge et 375 376 al., 2017), in accordance with previous findings (Hilton et al., 2016). These increases 377 of Cu and Zn concentrations precede the onset of disease symptoms (Figure 4A-B), 378 while Fe trails them (Figure 4C) (Enge et al., 2017). We hypothesized that this 379 observed accumulation could be the result of the over-expression of mSOD1 in the SOD1^{G93A} mouse model and that further testing, involving tissues from mice that 380 over-express SOD1^{WT} was necessary to determine whether a pathologic accumulation 381 382 of Cu occurred in muscle tissue.

383

384 Here, Cu, Zn and Fe concentrations measured in samples from mice over-expressing 385 wild type SOD1 were similar to those found in tissues from non-transgenic controls 386 (Figure 4A-C). This is in agreement with results from a previous study comparing Cu 387 in spinal cord tissues from mice over-expressing wild-type SOD1 and non-transgenic 388 controls (Tokuda et al., 2009). Since the accumulation of these metals in muscle tissue 389 is not related to SOD1 over-expression, we contend that it represents a pathological 390 feature of ALS that warrants further investigation, particularly as the accumulation of 391 Cu is pre-symptomatic (Figure 4A). While the method applied here does not enable us 392 to determine the origin of the accumulation, several hypotheses regarding its involvement in ALS pathology are presented below: 393

394

First, as previously proposed (Enge et al., 2017), Cu accumulation may result in toxicity at the NMJ and the onset of ALS as a distal motor neuropathy that proceeds

397 via a 'dying-back' mechanism (Enge et al., 2017; Fischer et al., 2004; Frey et al., 398 2000; Kiernan et al., 2011). In this case distal axonal degeneration precedes neuronal 399 degeneration, whereby axons and NMJs are affected early and the MN withdraws due 400 to perturbations arising from muscle tissue (Zhou et al., 2015). This contrasts the 401 toxic-gain-of-function and 'dying-forward' hypotheses in which denervation results 402 from the CNS. To understand the potential role of muscular Cu as a trigger for axonal 403 denervation, its location within the tissue must be considered: Cu may reside in the 404 cytosol or extracellular space. In the cytosol, it could lead to increased reactive 405 oxygen species (ROS)-related stress and ultimately cellular damage and apoptosis 406 (Linder, 1991). However, Cu is unlikely to be present as a free ion, as this would 407 trigger an anti-oxidant response through SOD1 and other protective mechanisms, in 408 order to reduce its impact. Mechanisms to protect against Cu toxicity include the 409 intrinsic stress response of the heat shock proteins (e.g. heat shock protein 70) (Urani 410 et al., 2001), which can be triggered by sub-lethal concentrations of Cu. Additionally, 411 the metal-responsive transcription factor 1 (MTF-1) (Balamurugan and Schaffner, 412 2006) acts under both high and low Cu concentrations to control the expression of 413 metallothioneins and other components (Cu importer/transporter 2) that control Cu 414 homeostasis. Copper in the extracellular space can bind to a variety of proteins, 415 including ceruloplasmin, extracellular SOD1, extracellular metallothionein and 416 albumin (Linder and Hazegh-Azam, 1996). The work presented here is limited to 417 organ-level resolution and therefore does not provide any insight into how Cu 418 accumulates in tissues, including whether it is located inside or outside cells, or both.

419

420 Second, the accumulation of Cu in muscle tissue may be the consequence of 421 hypermetabolism. Increases in energy expenditure are intrinsically linked to ALS

progression, preceding axonal retraction and denervation (Dupuis et al., 2004; Dupuis
and Loeffler, 2009; Ferri and Coccurello, 2017). Even though previous work has
shown that mitochondrial defects precede the onset of MN loss (Basun et al., 1991;
Jaarsma et al., 2000; Kong and Xu, 1998), specific mechanisms and causal links
between hypermetabolism and mitochondrial dysfunction remain largely unknown.

427

428

429 Mitochondrial respiration produces electrons, which may escape the electron transport 430 chain to induce the formation of reactive oxygen species (ROS) (Adam-Vizi and 431 Chinopoulos, 2006; Liemburg-Apers et al., 2015; Murphy, 2009). Mitochondrial 432 dysfunction, including muscle tissue, has been identified as one of the key features of 433 ALS (Leclerc et al., 2001); aggregated, swollen, vacuolated and fragmented 434 mitochondria, as a possible result of mSOD1 interaction (Pickles et al., 2016), are 435 able to explain other pathological features such as oxidative stress, glutamate 436 excitotoxicity and apoptosis (Abel et al., 2012; Faes and Callewaert, 2011; Martin, 437 2011; Smith et al., 2017). It was suggested that dys-regulation of Cu homeostasis may 438 play a substantial role in the development of the mitochondrial defects associated with 439 ALS (Son et al., 2007).

440

The observed Cu accumulation in muscle tissue may be the result of more than one mechanism, whereby the initial increase could result from an unidentified process that also increases ROS production. Increases in ROS could stimulate glucose uptake, which in return could stimulate ROS production leading to a positive feedback loop (Liemburg-Apers et al., 2015). Simultaneously, Cu transport into mitochondria could be up regulated to meet the demand of cellular respiration, resulting in the observed

447 hypermetabolism and overall accumulation of Cu in the tissue.

448

449 An updated view on the role of Cu in muscle tissue in ALS

450 The specific role of accumulated Cu in muscle tissue associated with ALS pathology 451 remains unclear. The data presented here shows an increase in metal concentrations in muscle tissues of the SOD1^{G93A} mouse model over time. This is interpreted as a 452 453 pathological process (Figure 5C), which may result in both a *toxic-gain-of-function* 454 (partially metalated SOD1 aggregates are toxic and lead to axonal retraction), as well 455 as a *dying-back* mechanism (e.g. accumulation of insoluble mSOD1 aggregates and 456 glutamate toxicity result in ALS presenting as a distal axonal retraction that precedes 457 death of MNs [see Ref (Dadon-Nachum et al., 2011) for a comprehensive review]) of 458 ALS onset and progression. The unifying feature in both these mechanisms is Cu, be 459 it the above-described lack of up-regulation in the CNS or the apparent accumulation 460 in ANS tissues. We therefore propose a new theoretical model here, in which both the 461 toxic-gain-of-function and the dying-back hypotheses could be occurring in parallel, 462 to explain the mechanisms of ALS pathology (Figure 5C). The relative proportion to 463 which each process contributes to the end result of muscular denervation remains 464 unclear.

465

The model proposed here combines findings of general Cu accumulation in peripheral tissues, specifically muscle tissue, and lack of up-regulation of Cu in the CNS. It is proposed that there are separate processes (Figure 5A, B) associated with Cu (either as a trigger or consequence), whose specific distribution on an organ level we have tested (Figure 5C). This specific distribution may contribute to the development of 471 ALS in these mice (see above). Furthermore, the model explains why Cu chelators 472 (Hottinger et al., 1997; Nagano et al., 2003), as well as synthetic Cu-containing 473 therapeutic agent (Roberts et al., 2014) likely act to prolong the survival of SOD1 474 mouse models. In addition to the transformation of labile intermediate forms of SOD1 475 into apo-SOD1, (Roberts et al., 2014) Cu chelators may also remove some generally 476 available Cu to prevent its accumulation in tissues. Simultaneously the synthetic Cucontaining therapeutic agent Cu^{II}(atsm), which is able to readily cross the BBB, was 477 478 found to deliver Cu to mSOD1 so it is in a (stable) holo-SOD1 state (Hilton et al., 479 2017; Soon et al., 2011; Williams et al., 2016). Prolonged survival (>20%) in G93A mice treated with Cu^{II}(atsm) has been reported, however this lifespan is greatly 480 481 extended in G93A mice co-expressing the copper chaperone for superoxide dismutase 482 (CCS) (G93AxCCS) and treated transdermally from prenatal stages (Williams et al., 483 2016). Even though this presents a very important milestone in finding a feasible 484 treatment option for ALS, it is not a cure as all mice developed typical end stage 485 motor neuron disease. Furthermore, G93AxCCS mice removed from treatment after 21 days had the same life span as SOD1^{G93A} mice (Williams et al., 2016). This 486 487 indicates that at least part of the toxic-gain-of-function of SOD1 resides in mSOD1 488 being metal deficient, while pathological Cu accumulation in muscle tissue may also 489 contribute.

490

Based on the model we have proposed here, a treatment combining a specific Cu chelator (to prevent Cu accumulation in non-CNS tissues), and a synthetic Cucontaining therapeutic agent (to deliver Cu to the CNS) could be most beneficial to treat ALS in the here assessed G93A mouse model. An application to humans with SOD1 familial ALS, and potentially other genotypes, could be beneficial as well;

496 even though studies on human tissue samples did not observe Cu accumulation in 497 neither liver or kidneys (Sillevis Smitt et al., 1992), nor in brain (Gellein et al., 2003) of ALS patients compared to controls. The lack of general accumulation of Cu in 498 499 human ALS patients is non conclusive at this point due to the overall limited number 500 of studies available. Future studies should systematically assess Cu concentrations in 501 human ALS patient organs, including muscle tissues, target tissues of ALS. 502 Furthermore, Cu accumulation in ALS models has to further be examined as it may 503 serve as a biomarker of disease.

504

505 CONCLUSIONS

506 The assessment of metal concentrations in several tissues from control and ALS mice 507 showed accumulation of Cu in tissues controlled by the ANS compared to controls, 508 which was also not observed in CNS tissues. This accumulation was identified to be 509 pre-symptomatic. Muscle tissue showed accumulation of Cu, which was not driven by 510 over-expression of SOD1 (Enge et al., 2017), but rather was associated with disease 511 pathology in SOD1^{G93A} mice and was evident at pre-symptomatic stages. The results 512 further indicate that there is a lack of Cu uptake into the brain despite the higher Cu requirement due to mSOD1 over-expression in the SOD1^{G93A} mouse model of ALS. 513 514 The observed pre-symptomatic changes in metal concentrations in tissues controlled 515 by the ANS provide further evidence for a role of metals, in particular Cu, in the 516 development of ALS pathology.

517

518 The accumulation of Cu in muscle tissue leads us to propose a revised model of the 519 mechanisms underpinning ALS pathology, which unites the *toxic-gain-of-function* 520 and *dying-back* hypotheses. Our model offers another explanation to elucidate the axonal retraction of MNs in ALS, a process potentially underpinned by Cu accumulation. Further investigation into the potential applicability of metal concentrations as biomarkers for ALS is therefore warranted. Future studies should expand to study human tissues, and further establish in disease models the location and binding of Cu in ANS tissues to help further elucidate the role that Cu plays in the aetiology of ALS.

527

528 LIST OF ABBREVIATIONS

529	Ab	Albumin
530	ALS	Amyotrophic lateral sclerosis
531	ANS	Autonomic nervous system
532	ATOX	Copper metal chaperone
533	ATP7A	ATPase copper transporting alpha
534	BBB	Blood-brain-barrier
535	BCB	Blood-cerebrospinal fluid-barrier
536	CCS	Copper chaperone for superoxide dismutase
537	CNS	Central nervous system
538	COX	Cytochrome c oxidase
539	СР	Ceruloplasmin
540	CTR1	Copper transporter 1
541	CTR2	Copper transporter 2
542	Cu ^{II} (atsm)	Diacetyl-bis(4-methylthiosemicarbazonato)copper ^{II}
543	DMT1	Divalent metal transporter 1
544	FUS	Fused in Sarcoma
545	MN	Motor neuron
546	MTF1	Metal regulatory transcription factor 1
547	NMJ	Neuromuscular junction
548	Q-ICP-MS	Quadrupole-inductively coupled plasma-mass spectrometer
549	ROS	Reactive oxygen species
550	SOD1	Cu,Zn superoxide dismutase 1
551 552	TDP-43	TAR DNA-binding protein 43

- 553 **DECLARATIONS**
- 554 Ethical approval: UOW Animal Ethics committee approval: AE14/28; Approved
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- 556 *Consent for publication:* Not applicable

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- 565 HE, DFJ and AD interpreted the data and conceived the manuscript. TGE, HE, DFJ,
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926 Supplementary material:927

947		
928	File Name:	Supplementary Material_Data
929	File Format:	.docx
930	Title:	Raw data
931	Description:	This data set includes data relevant to this paper.
932		
933		
934	File Name:	Supplementary Material_Figure 1
935	File Format:	.docx
936	Title:	Supplementary Figure 1
937	Description:	This Figure presents Cu and Zn concentrations ($\mu g g^{-1}$ of dry tissue) in
938		brain and spinal cord tissue as a function of time (days). This includes
939		previously published results and newly produced results.