

**Supplementary Figure 1.** Spearman rank test correlation between *HDAC4* and *FGF21* mRNA expression levels (assessed by qPCR) in 9 post-mortem ALS muscle samples.



Supplementary Figure 2. FGF21 mRNA is increased in muscle and liver tissue in the SOD1<sup>G93A</sup> mouse. RNA was extracted from wild-type (WT) and SOD1<sup>G93A</sup> mouse tissue at post-natal day 60 and assessed by qPCR for FGF21 mRNA. All values represent fold-change compared to WT muscle which was set at 1. Data points represent individual mice and bars represent the mean  $\pm$  SD. \**P* < 0.05, \*\*\*\**P* < 0.0001; one-way ANOVA followed by Tukey's multiple comparisons test.



Supplementary Figure 3. Oxidative stressors induce *ATF4* and *PGC-1a* in NSC-34 motor neuron-like cells and C2C12 myoblasts. (A) and (B) *ATF4* and *PGC-1a* mRNA levels in NSC-34 cells were quantified after treatment with 100  $\mu$ M H2O2 or methionine-cysteine (MetCys)-depleted media for 24 h. (C) and (D) *ATF4* and *PGC-1a* mRNA levels were quantified in C2C12 myoblasts after exposure to the same conditions as in (A) and (B). Data points are independent biological samples and bars represent the mean  $\pm$  SD.\**P* = 0.048, \*\*\*\**P* < 0.0001; one-way ANOVA followed by Tukey's multiple comparisons test.



Supplementary Figure 4. Recombinant FGF21 treatment reverses cytotoxicity of NSC-34 motor neuron-like cells expressing SOD1<sup>G93A</sup>. NSC-34 cells expressing either WT SOD1 or SOD1<sup>G93A</sup> were treated with recombinant FGF21 (100 ng/ml). Cell viability was measured 24 hours later as described in the methods. Bars represent the mean  $\pm$  SD of 4 independent biological samples. \*\*P = 0.003; unpaired two-tailed t-test.



Supplementary Figure 5. C2C12 cells expressing FGF21 rescue SOD1<sup>G93A</sup>-mediated toxicity in NSC-34 motor neuron like cells in co-culture. (A) Schematic of co-culture system used. (B) Cell viability of NSC-34 cells expressing either WT-SOD1 or SOD1<sup>G93A</sup> (lower well) was assessed in the presence or absence of FGF21-expressing C2C12 cells (upper well). Data points represent biological replicates and bars are the mean  $\pm$  SD. \**P* = 0.025, \*\**P* = 0.002, \*\*\*\**P* < 0.0001; oneway ANOVA followed by Tukey's multiple comparisons test.

Cell Line	Sex	Age (y)	<b>Clinical Diagnosis</b>	Primary tissue	Mutation
CS0002iCTR	Μ	51	Normal	PBMC	N/A
CS83iCTR	F	21	Normal	Fibroblast	N/A
CS188iCTR	Μ	80	Normal	PBMC	N/A
CS14iCTR	F	35	Normal	Fibroblast	N/A
CS00iCTR	Μ	6	Normal	Fibroblast	N/A
FA0000011	F	49	Normal	Fibroblast	N/A
NN0003920	Μ	64	Normal	Fibroblast	N/A
CS0118iALS-SOD1-I114T	F	73	ALS	Fibroblast	SOD1 I113T
CS28iALS	Μ	47	ALS	Fibroblast	C9ORF72 (HRE ~800)
CS29iALS	Μ	47	ALS	Fibroblast	C9ORF72 (HRE ~800)
CS52iALS	Μ	49	ALS	Fibroblast	C9ORF72 (HRE ~800)
CS30iALS	F	51	ALS	Fibroblast	C9ORF72 (HRE ~70)
NN0004306	F	51	ALS	Fibroblast F10330	C9ORF72 (HRE 2.7kb)
NN0004307	М	57	ALS	Fibroblast F09152	C9ORF72 (HRE 6-8kb)

Supplementary Table 1: Demographic and clinical data for iPSC-derived motor neurons.

ALS = amyotrophic lateral sclerosis; F = female; HRE = hexanucleotide repeat expansion; M = male; ORF = open reading frame; PBMC = peripheral blood mononuclear cells; SOD1 = superoxide dismutase 1; y = years

	Biopsy		Autopsy	
	Normal	ALS	Normal	ALS
Number	24	36	22	23
Mean age (years) <sup>a</sup>	$52 \pm 15$	57 ± 13	$67 \pm 12$	$64 \pm 11$
Age range (years)	24 - 77	27-86	34 - 83	40 - 81
Gender (M:F)	11:13	21:15	18:4	18:5
Duration <sup>b</sup> (m)		$15 \pm 9$		$51 \pm 32$
Diagnosis		Spinal onset (33)		Spinal onset (20)
C		Bulbar onset (3)		Bulbar onset (3)
Muscle sampled				
Biceps brachii	5	2		2
Deltoid	3	11	4	3
Vastus lateralis	15	8	3	5
Tibialis anterior	1	15		Triceps (1)

## Supplementary Table 2. Demographic and clinical data of tissue samples.

<sup>a</sup> Mean age ( $\pm$  SD) at time of sample collection. <sup>b</sup> Mean duration ( $\pm$  SD) from onset of symptoms to sample collection. Duration was unknown for three ALS patients in the biopsy pool.

Supplementary Table 3: Demographic and clinical data of plasma sample
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	Normal	ALS
Number	23	28
Mean age (y) <sup>a</sup>	$61 \pm 9$	$59 \pm 10$
Age range (y)	45 - 84	35-82
Gender (M:F)	12:11	18:10
Duration <sup>b</sup> (m)		$26 \pm 20$
Onset		Spinal onset (22)
		Bulbar onset (6)

 $\label{eq:stars} \begin{array}{l} F = \mbox{female; } M = \mbox{male; } m = \mbox{months; } y = \mbox{years} \\ {}^a \mbox{Mean age} \ (\pm \mbox{SD}) \mbox{ at time of sample collection.} \\ {}^b \mbox{Mean duration} \ (\pm \mbox{SD}) \mbox{ from onset of symptoms to sample collection.} \end{array}$ 

Plasma FGF21 (FC) <sup>a</sup>	< 1.5	≥ 1.5	-
Number	7	9	-
Age (y) <sup>b</sup>	$65 \pm 9$	$57 \pm 9^{d}$	
Age range (y)	58 - 82	$42 - 70^{d}$	
Gender (M:F)	4:3	7:2	
Duration <sup>c</sup> (m)	$18 \pm 10$	$26 \pm 23^{d}$	
Onset	Bulbar (4)	Spinal (9)	
	Spinal (3)	-	

**Supplementary Table 4: ALS study patients** 

 $\begin{array}{l} F = \mbox{female; FC} = \mbox{fold-change; M} = \mbox{male; m} = \mbox{months; y} = \mbox{years} \\ {}^{a}\mbox{Fold-change over normal control group} \\ {}^{b}\mbox{Mean age ($\pm$ SD) at time of sample collection.} \\ {}^{c}\mbox{Mean duration ($\pm$ SD) from onset of symptoms to sample collection.} \\ {}^{d}\mbox{No significant difference between the 2 groups} \end{array}$