

# Identification of property outliers among ALS-associated SOD1 mutations

## COMMON EFFECT ON SURFACE HYDROGEN BONDS

### SUPPLEMENTARY MATERIAL

**Table S1.** Urea and thermal denaturation data used to plot  $\Delta\Delta G^{\text{norm}}$  versus disease progression ( $ST$ ) (Fig. 5). The stability data was normalized prior to comparison.  $ST$  is the mean survival time after first diagnosis and  $n$  is the number of patients in the data set.

Genotype	$\Delta\Delta G_{\text{D,N}}$ (kcal/mol) (1)	$\Delta\Delta G^{\text{norm}}$	$\Delta T_m$ (°K) (2)	$\Delta T_m$ (°K) (3)	$\Delta T_m$ (°K) (4)	$\Delta T_m$ (°K) (5)	$\Delta T_m$ (°K) (6)	$ST$ (years)	$n$
A4V	-4.31	0.85	-12.01	-12.4	-8.9	-	-	1.2	205
V7E	-	0.52	-6.58	-	-	-	-	5.7	3
G37R	-	0.55	-7.13	-	-	-	-	17.0	27
L38V	-3.24	0.71	-9.57	-	-	-	-	2.4	22
G41D	-3.47	0.80	-	-	-	-	-	14.1	15
G41S	-4.47	0.96	-	-	-	-	-	1.0	16
H43R	-4.05	0.91	-	-	-	-	-	1.8	12
H46R	0.48	0.03	3.64	-	-	-	-	17.6	49
H48Q	-	0.45	-5.18	-	-	-	-	1.2	4
D76V	-0.06	0.15	-	-	-	-	-	18.8	4
D76Y	-0.09	0.16	-	-	-	-	-	9.0	2
L84V	-2.65	0.69	-10.73	-	-	-	-	3.2	10
G85R	-0.92	0.35	-3.63	-	-	-	-3.8	6.0	11
N86D	-0.94	0.32	-	-	-	-	-	-	0
N86K	-1.41	0.41	-	-	-	-	-	1.7	7
N86S	-0.45	0.28	-2.97	-	-	-	-	6.8	4
D90A	-0.65	0.26	-	-	-	-	-	8.0	15
D90V	-1.85	0.49	-	-	-	-	-	2.7	3
G93A	-2.97	0.70	-9.5	-10.5	-10.5	-	-	3.1	16
G93D	-	0.84	-	-	-	-12.8	-	8.8	7
G93R	-	0.60	-8.23	-	-3.3	-10.1	-10.1	5.3	4
G93S	-	0.61	-	-	-	-8.3	-	8.0	11
G93V	-	1.00	-	-	-	-15.9	-	6.0	12
E100G	-2.22	0.52	-12.56	-	-7.8	-	-7.7	4.7	50
E100K	-	0.30	-2.18	-	-	-	-	10.0	1
D101G	-1.39	0.40	-	-	-	-	-	1.9	3

D101N	0.75	0.10	-0.09	-	-	-	-	2.3	17
I104F	-1.24	0.37	-	-	-	-	-	21.3	3
S105L	-2.60	0.63	-	-	-	-	-	3.5	7
L106V	-3.62	0.82	-	-	-	-	-	1.9	6
I113T	-2.48	0.68	-9.79	-	-	-	-12.2	4.3	38
G114A	-3.27	0.76	-	-	-	-	-	2.7	2
D124V	-	0.01	3.42	-	-	-	-	-	0
D125H	-	0.21	-0.53	-	-	-	-	1.8	2
S134N	-	0.23	-0.9	-	-	-	-	1.2	3
N139D	0.36	0.07	-	-	-	-	-	-	-
N139K	-0.24	0.24	-2.12	-	-	-	-	-	-
L144F	-1.89	0.46	-4.29	-	-	-	-	11.8	15
L144S	-1.07	0.36	-3.65	-	-	-	-	12.3	2
V148G	-4.56	1.00	-	-	-	-	-	2.1	11

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**Table S2.** The reference set of 75 confirmed ALS associated SOD1 mutations with fast and slow disease progression as listed in (1). *n* is the number of reported patients for each mutation. The mutants measured in this study are in bold letters.

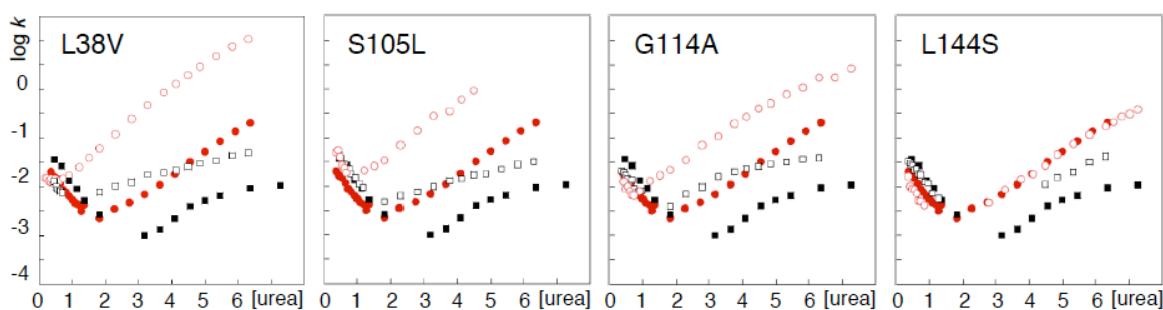
Mutant	Fast <5 years	<i>n</i>	Slow >5years	<i>n</i>
A4S	>3	0		
A4T	1,5	21		
A4V	1,2	205		
C6F	1	1		
C6G	0,2	2		
V7E <sup>1</sup>			5,7	3
L8Q	1,6	5		
G10V	1,1	5		
G12A	>4	0		
G12R <sup>2</sup>			5,5	2
V14G	1,7	1		
N19S	2,1	3		
F20C	2	7		
E21G <sup>2</sup>			17,2	5
G37R <sup>2</sup>			17	27
L38R <sup>2</sup>	>2	0		
<b>L38V</b>	2,4	22		
G41D <sup>1</sup>			14,1	15
G41S	1	16		
H43R <sup>2</sup>	1,8	12		
F45C			>6	0
H46R <sup>2</sup>			17,6	49
H48Q	1,2	4		
N65S			>14	0
G72C	4,4	1		
G72S	3,3	2		
<b>D76V<sup>2</sup></b>			18,8	4
<b>D76Y<sup>2</sup></b>			11	2
L84F			5,8	18
L84V	3,2	10		
G85R <sup>2</sup>			6	11
<b>N86D<sup>1</sup></b>	>4	0*		
<b>N86K<sup>2</sup></b>	1,7	7		
<b>N86S</b>			6,8	4
A89V	>3,8	0		
D90A <sup>2</sup>			8	15
<b>D90V<sup>2</sup></b>	2,7	3		
G93A	3,1	16		
G93C			12,1	27
G93D <sup>1</sup>			8,8	7
G93R <sup>2</sup>			5,3	4
G93S			8	11
G93V			6	2
E100G <sup>2</sup>	4,7	50		
E100K <sup>2</sup>			10	1
<b>D101G<sup>2</sup></b>	1,9	3		
D101H <sup>2</sup>	1,2	2		
<b>D101N<sup>2</sup></b>	2,3	17		
D101Y <sup>2</sup>	0,9	1		

I104F			21,3	3
<b>S105L</b>	3,5	7		
L106V	1,9	6		
G108V	4	2		
I112M	3	9		
I112T	0,9	2		
I113F			>6	0
I113T	4,3	38		
<b>G114A</b>	2,7	2		
R115G <sup>1</sup>	2,5	2		
D124V <sup>2</sup>	>2	0		
D125H <sup>2</sup>	1,8	2		
L126S			8	5
L126X	3,8	14		
S134N	1,2	3		
N139H	3	6		
A140G			10	1
G141E <sup>1</sup>	3,5	2		
L144F			11,8	15
<b>L144S</b>			12,3	2
A145T	1,6	2		
C146R <sup>2</sup>	2,8	2		
V148G	2,1	11		
V148I	1,7	5		
I149T	2,7	15		
<b>I151T</b>			20	1

<sup>1</sup>Increase of the net repulsive charge (blue in plot). <sup>2</sup>Decrease of the net repulsive charge (red in plot).

\*Patient still alive and displays a slow disease progression (P.M. Andersen, unpublished data).

1. Wang, Q., Johnson, J. L., Agar, N. Y. R., & Agar, J. N. (2008) Protein aggregation and protein instability govern familial amyotrophic lateral sclerosis patient survival *Plos Biology* 6: 1508-1526.



Supplementary Fig. 1. Chevron plots of L38V, S105L, G114A and L144S.  
apoSOD1<sup>pWT</sup><sub>mono</sub> apoSOD1<sup>pWT</sup><sub>mono</sub> (●), apoSOD1<sup>mut</sup><sub>mono</sub> apoSOD1<sup>mut</sup><sub>mono</sub> (○), apoSOD1<sup>pWT</sup><sub>dimer</sub> apoSOD1<sup>pWT</sup><sub>dimer</sub> (■) and apoSOD1<sup>mut</sup><sub>dimer</sub> apoSOD1<sup>mut</sup><sub>dimer</sub> (□).