## The Non-Fibrillating N-terminal of α-Synuclein Binds and Co-Fibrillates with Heparin

4	Line K. Skaanning <sup>1</sup> , Angelo Santoro <sup>23</sup> , Thomas Skamris <sup>1</sup> , Jacob Hertz Martinsen <sup>12</sup> , Anna Maria
5	D'Ursi <sup>3</sup> , Saskia Bucciarelli <sup>1</sup> , Bente Vestergaard <sup>1</sup> , Katrine Bugge <sup>2</sup> , Annette E. Langkilde <sup>1,4</sup> , Birthe B.
6	Kragelund <sup>2,s</sup>
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Figure S1 – Purification of His-SUMO-aSN<sub>1-61</sub>. A) SDS-PAGE of fractions from the initial steps in 24 25 purification of aSN<sub>161</sub>. Stained with InstantBlue<sup>™</sup>. S: Supernatant cell lysis. FT: flow-through from 26 HisTrap step. F: Fractions from HisTrap. D: Dialysate of cleaved with His-ULP1. R: Pooled fraction 27 3-13 from reverse HisTrap D2: dialysate of aSN<sub>1-61</sub> into 50 mM Tris, 10 mM NaCl, pH 8.5. IE: Fractions from ion exchange. B) Chromatogram showing the stepwise elution profile of His-SUMO-28 29 aSN<sub>161</sub> on the HisTrap column. Blue trace corresponds to absorption at 280 nm and red trace 30 corresponds to absorption at 215 nm. C) SDS-PAGE of the fractions from ion exchange of 31 purification of aSN<sub>1.61</sub>. The gel was stained with Coomassie. D) Elution profile of aSN<sub>1.61</sub> from IE 32 column. Blue trace corresponds to absorption at 280 nm and red trace corresponds to absorption at 33 215 nm. The base line for absorbance at 280 nm and 215 nm was set to overlap in the beginning of 34 both chromatograms.



**Figure S2 - SAXS data of aSN**<sub>L61</sub> **at different concentration**. A) Scattering intensities as a function of the momentum transfer. Scattering curves were initially scaled by concentration and final adjustments made by scaling to the highest concentration sample. 223  $\mu$ M (light green), 287  $\mu$ M (lime), 366  $\mu$ M (orange), 446  $\mu$ M (magenta), 510  $\mu$ M (purple) and 573  $\mu$ M (bluu) and B) corresponding Kratky plot, all samples shown (colors as in panel A). C) Guinier range (experimental data in blue) and fit (black) for aSN<sub>L61</sub> at 573  $\mu$ M.

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Figure S3 – Test of fibrillation properties for aSN<sub>1-61</sub>. A) ThT profile of aSN<sub>1-61</sub> during incubation in
conditions known to induce fibrillation of aSN. B) Far-UV CD spectrum before incubation (solid)
and after incubation (dashed).

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**Figure S4 – ThT control measurement**. Fluorescence emission at  $480\pm5$  nm (ThT assays) of heparin at concentrations corresponding to those used in the experiments with incubation of aSN<sub>1.61</sub> and heparin (Figures 2 and 3). The ratios above each panel indicate the corresponding experiments with aSN<sub>1.61</sub>. The specific heparin concentrations are: A) 666  $\mu$ M heparin; B) 333  $\mu$ M heparin; C) 66.7  $\mu$ M heparin; D) 29.3  $\mu$ M heparin; E) 29.3  $\mu$ M heparin; F) 14.7  $\mu$ M heparin; G) 3.7  $\mu$ M heparin; H) 1.8  $\mu$ M heparin; I) 1.5  $\mu$ M heparin; J) 0.7  $\mu$ M heparin. The concentration for heparin in D and E is the same, as the aSN<sub>1.61</sub>:heparin experiments were performed with different aSN<sub>1.61</sub> concentrations.





**Figure S5** – Normalized residual concentration of (A)  $aSN_{1-61}$  and (B) aSN in the supernatant after incubation (black) and before incubation (grey) with different molar ratios of (A)  $aSN_{1-61}$ :heparin and (B) aSN:heparin.



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Figure S6 - SAXS data of heparin in different buffers with different salt concentrations. All data are shown on absolute scale and normalized by concentration. A) heparin in 50 mM NaP 0 mM NaCl. 70 Concentration of heparin 53.3 µM (red), 66.7 µM (purple), 133 µM (magenta), 333 µM (black). B) 71 heparin in 50 mM NaP 150 mM NaCl. Concentration of heparin 53.3 µM (light green), 66.7 µM 72 (green), 133 µM (dark green), 333 µM (grey). C) Heparin in 50 mM NaP 500 mM NaCl. 73 Concentration of heparin 53.3 µM (light blue), 66.7 µM (light purple), 133 µM (dark purple), 333 74 µM (dark blue). D-F) Zoom of low-s region of A-C, respectively. G) heparin in 50 mM Tris, 0 mM

- 75 NaCl. Concentration of heparin 53.3 μM (red), 66.7 μM (purple), 133 μM (magenta), 333 μM L
- 76 (black). H) heparin in 50 mM Tris 150 mM NaCl. Concentration of heparin 53.3 μM (light green),
- 77 66.7 μM (green), 133 μM (dark green), 333 μM (grey). I) Heparin in 50 mM Tris 500 mM NaCl.
- 78 Concentration of heparin 53.3 μM (light blue), 66.7 μM (light purple), 133 μM (dark purple), 333
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Figure S7 – ThT and CD analyses of factors potentially affecting aSN<sub>140</sub> fibrillation. A) ThT assay of heparin alone incubated with 1 M NaCl. B) ThT assay of CS alone at 3 mg/mL (light grey) and 0.9 mg/mL (dark grey). C) ThT assay of β-casein at 333  $\mu$ M. D) ThT assay of aSN<sub>140</sub> incubated in 160 mM ammonium sulfate. E) Far-UV CD spectrum of heparin (green) in 1 M NaCl before (solid) and after (dotted) fibrillation. F) Far-UV CD spectrum of CS at 3 mg/mL (light grey) and 0.9 mg/mL (dark grey) after fibrillation.

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- 93 Table S1 Primary analysis of aSN<sub>141</sub> SAXS data as shown in Figure S2A. The apparent values
- 94 of  $R_s$  and  $I_o$  as obtained from Guinier analysis in PRIMUS <sup>+</sup> together with estimated molecular weights
- 95 based on the Bayesian inference approach <sup>2</sup>.

Sample conc	$\mathbf{R}_{s}$	$\mathbf{I}_{o}$	MW estimate (credibility interval)
(µM)	(Å)	relative scale	(kDa)
573	24.0±0.3	1360±5	11.3 (10.0-12.4)
510	24.1±0.4	1358±5	11.2 (10.0-12.4)
446	24.2±0.8	1367±5	9.5 (9.1-12.4)
366	24.8±1.6	1390±6	11.2 (10.0-12.4)
287	24.8±0.6	1409±7	15.5 (14.5-16.5)
223	25.1±1.7	1407±10	11.2 (10.0-12.4)

97 **Table S2 - Primary analysis of SAXS data for heparin (Figure S7).** Heparin in 50 mM NaP with 98 0, 150 or 500 mM NaCl (top) and heparin in 50 mM Tris with 0, 150, or 500 mM NaCl (bottom). The 99 apparent values of  $R_i$  and  $I_o$  (i.e. with no correction for intermolecular interactions) are shown together 100 with the apparent estimated molecular weight based on absolute scaling to illustrate the systematic 101 influence of the interparticle interaction effects.

Heparin	50 mM NaP 0 mM NaCl			50 mM NaP 150 mM NaCl			50 mM NaP 500 mM NaCl		
(µM)	$R_{s}$ (nm)	I <sub>o</sub>	MW	$R_{s}(nm)$	I <sub>o</sub>	MW	R <sub>s</sub> (nm)	I.	MW
			(kDa)			(kDa)			(kDa)
333	3.91±0.43	5549±16	5.5	4.17±1.62	7815±32	7.8	5.19±0.48	7386±19	7.4
133	4.53±0.6	6885±31	6.9	4.88±0-15	9038±39	9	4.72±0.09	7406±27	7.4
66.7	4.73±0.49	7588±47	7.6	5.08±0.23	9597±54	9.6	4.77±0.7	7296±54	7.3
53.3	4.78±0.78	7734±57	7.7	5.08±0.33	9578±89	9.6	4.67±0.32	6795±45	6.8
Heparin	50 mM Tr	is 0 mM NaO	C1	50 mM Tris 150 mM NaCl		50 mM Tris 500 mM NaCl			
(µM)	$R_{s}$ (nm)	I <sub>o</sub>	MW	R <sub>s</sub> (nm)	I <sub>o</sub>	MW	R <sub>s</sub> (nm)	I.	MW
			(kDa)			(kDa)			(kDa)
333	3.37±0.97	7255±14	7.2	4.5±1.8	10263±27	10.3	4.48±0.78	10431±33	10.4
133	4.74±0.35	11042±35	11	4.8±0.25	907±40	9	4.91±0.86	12311±48	12.3
66.7	5.14±0.67	12885±51	12.9	5.07±0.36	5602±44	5.6	5.30±0.47	9498±52	9.5
53.3	5.25±0.34	12562±56	12.6	5.29±0.47	7051±59	7	5.21±0.51	13950±70	13.9
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- **Table S3** Overview of fragments of aSN, and their fibrillation/aggregation potential reported
- 106 from the literature.

Fragment	Observation	Reference
1-60	No fibrillation Fibrillation in trifluoroethanol	3,4
1-65	Aggregation	5
1-70	Aggregation	3
1-87	Fibrillation	6
1-95	Aggregation	5
1-100	Fibrillation Fibrillation in trifluoroethanol	3,4
1-108	Fibrillation	7
1-110	Fibrillation	8 9
1-120	Fibrillation	6,8-10
1-124	Fibrillation	7
1-130	Fibrillation	9
11-140	Fibrillation	11
15-95	Aggregation	5
15-65	Aggregation	5
16-35	Fibrillation	5
21-140	Fibrillation	11
31-140	Fibrillation	11
32-140	Fibrillation	10
41-140	Fibrillation	11
51-140	Fibrillation	11
58-140	Aggregation	10
60-95	Fibrillation	12
NAC 1-18 (61-78)	Aggregation	13

NAC8-15 (68-75)	No fibrillation	13
NAC8-16 (68-76)	No fibrillation	13
NAC8-18 (68-78)	Fibrillation	13
NAC9-16 (69-76)	No fibrillation	13
71-140	No fibrillation	11
NAC12-18 (72-78)	No fibrillation	13
NAC 19-35 (79-95)	No fibrillation	13
93-115	Fibrillation	5
116-136	Fibrillation	5
Δ71-82	No fibrillation	14
Δ71-74	Fibrillation	3
Δ66-74	Fibrillation	3
Δ1-74	Fibrillation	3
Δ71-82	Fibrillation	п
Δ73-83	Fibrillation	и
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