Supplemental figures

Supplementary Figure 1 *A*, Comparison of all known synuclein sequences. *B*, Comparison of human, mouse and α -, β - and γ -synuclein sequences. Program used: ClustalW2 (http://www.ebi.ac.uk/Tools/clustalw2/index.html).

Supplemental Figure 2: Mapping the CK1 phosphorylation sites: WT, S129A and S129E were phosphorylated for 24 h with CK1 and the different phosphorylated species (I,II,III) were separated by semi-preparative RP-HPLC and analyzed by MALDI-MS. A, The Coomassie blue stained SDS gels of the mono-, di-, and unphosphorylated species of each protein. The S129A α-syn was subjected to further trypsin digestion and tandem mass spectrometry. B, 4280-4380 m/z inset covering the pseudomolecular region of C-terminal peptide [103-140] of S129A analyzed by MALDI-MS in positive ion mode. The peak corresponding to the monophosphorylated peptide at m/z 4363.4 could not be detected. C, MALDI-MS spectra in negative ion mode showing the m/z region 1450-1850 of the tryptic digest of unphosphorylated (blue), monophosphorylated (red), and diphosphorylated (green) peaks obtained from an in vitro phosphorylation of S129A with CK1. The peak at m/z 1478.0 corresponds to the sequence TVEGAGSIAAATGFVKK ([81-96]), which contains three potential phosphorylation sites. The difference in mass of 80 or 160 Da is consistent with the addition of one and two phosphate groups, respectively. D, CID spectrum of the doubly modified peptide [81-96]. Fragmentation pattern of the [M + 2H]2+ 81-96 peptide bearing two phosphates on S87 and T92 residues. (Top right) Sequence coverage based on detected v and b fragment ions. Ions carrying the phosphate group are shown with one or two stars (* and **) on the sequence.

Supplemental Figure 3. CK1-mediated phosphorylation inhibits the fibrillization of S129A and S129E α-syn. *A*, ThT fluorescence measurements of CK1-phosphorylated S129A and *B*, S129E α-syn (100 μM) (white) and their unphosphorylated control (black). S129A and S129E α-syn were phosphorylated for 24 h with CK1 and then aggregated for the indicated length of time. Negatively stained TEM images of phosphorylated S129A and S129E and their unphosphorylated control after 12 h of incubation at 37°C under agitating conditions (scales bar 0.2 μm). *C*, ThT fluorescence measurements of CK1-phosphorylated S129E α-syn (20 μM). S129E was phosphorylated for 24 h with CK1 and then the different phosphorylated species were separated by RP-HPLC. S129E and S129E/S87-P samples were aggregated for the indicated length of time. Negatively stained TEM images of S129E and S129E/S87-P after 24 h of incubation at 37°C under agitating conditions (scales bar 0.2 μm).

Supplemental Figure 4. *A*, Comparison of two-dimensional 1H -15N HSQC spectra of unphosphorylated WT (blue) and phosphorylated WT (red) α-syn. A dashed rectangle marks glutamine and asparagine side chain resonances. *B*, Comparison of 3J(HN,Hα) scalar couplings observed in nonphosphorylated WT (black), nonphosphorylated S129D (blue) and phosphorylated S129D (grey) α-syn at 15 °C. *C*, Comparison of 15N R1ρ spin relaxation rates

in nonphosphorylated WT (black) and phosphorylated S129D (grey). The domain organization of α -syn is shown on the top: basic N-terminal domain (red), hydrophobic NAC region (yellow), acidic C-terminal domain (blue) and the six repeats (green).

Supplemental Figure 5. *In vitro* phosphorylation of α -syn monomers and fibrils as a function of time by CK1. Samples were separated on a 12% SDS gel and probed with anti- α -syn (211, 1:500), anti-S129-P (1:5000) or anti-S87-P (1:100) α -syn antibodies.

Α

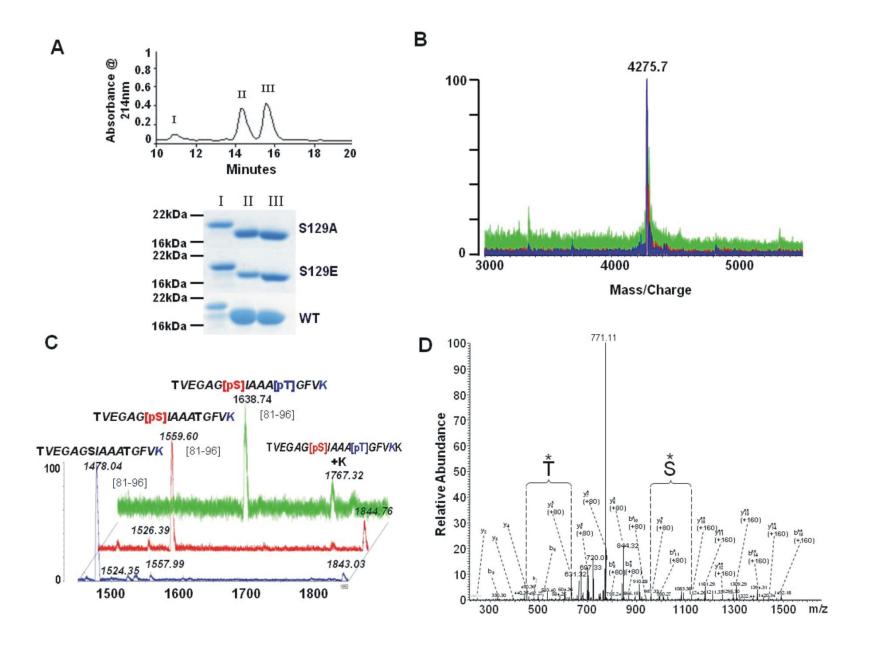
Suppl. Figure 1

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Canary	MDVFMKGLISKAKEGVVAAAENTKQGVAEAAGNTKEGVLYVGSHTKEGVVHGVTTVAENTMEDVSNVGGAVVTGVTAVAQNTVEGAGNLAAATGLVKKDQL 100
Chicken	MDVFMKGLINKAKEGVVAAAENTROGNAEAAGNTKEGVLYVGSHTKEGVVHGVTTIVAENTMEDVSNIVGGAVVTGVTAVAQNTVEGAGNITAAATGLVKKDOLI 100
Mouse	MDVFMKGLISKAKEGVVAAAEHTKQGVAEAAGHTKEGVLYVGSHTKEGVVHGVTTVAEHTMEDVTNVGGAVVTGVTAVAQHTVEGAGNLAAATGFVKKDQM 100
Rat	MDVFMKGLSKAKEGVVAAAENTKQGYAEAAGNTKEGVLYVGSNTKEGVVHGYTTVAENTNEEVTNVGGAVYTGYTAVAQNTVEGAGNIAAATGFVKKDQN 100
Bovine	MDVFMKGUSKAKEGVVAAAEHTKQGVAEAAGHTKEGVLYVGSHTKEGVVHGVTTVAEHTMEDVTNVGEAVVTGVTAVAQHTVEGAGSIAAATGFGKKDHM 100
Pig	MDVFMKGUSKAKEGVVAAAENTKQGVAEAAGNTKEGVLYVGSNTKEGVVHGVTTVAENTNEBDYTNVGEAVVTGVTAVAQNTVEGAGSIAAATIGFGKKDQL 100
Bonobo	MDVFMKGLSKAKEGVVAAAEMTKQGYAEAAGMTKEGVLYVYGSMTKEGVVHGYATVAEMTWEBYTNVGGAVYTGYTAVAQMTVEGAYSTAAATGFVKKDYL 100
Chimpanzee	MDVFMKGUSKAKEGVVAAAEHTKQGVAEAAGHTKEGVLYVGSHTKEGVVHGVATVAEHTMEQVTNVGGAVVTGVTAVAQHTVEGAGSIAAATGFVKKDQU 100
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Common-woolly-monkey	MDVFMKGLISKAKEGVVAAAENTKOGVAEAAGHTKEGVLYVGSHTKEGVVHGVTTVAEHTNEGVTSVGGAVVTGVTAVAOHTVEGAGNIAAATIGFVKKDHS 100
Black-handed-spider-monkey	MDVFMKGLSKAKEGVVAAAENTKOGVAEAAGHTKEGVLYVGSKTKEGVVHGVTTVAEHTNEGVTSVGGAVVTGVTAVAOHTVEGAGNTAAATIGFVKKDHS 100
	MOVFMKGLSKAKEGVVAAAENTKOGVAEAAGNTKEGVLYVOSNTKEGVVHGVTTVAENTNEGVTNVGGAVVTGVTAVAONTVEGADNIAAATGFVRKDHL 100
Fugu-rubripes	MDAFMKGFSKAKDGVVAAAEHTKOGVTGAAENTKDGVMFVGTHTKDGVTVVAGHTNSSVSQVGGAMVTGVTAVAGHTVESAGSTAAATGLVKKE-P 95
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Zebra-finch	AKQNEEGFLDEGMUNNIGVAVDPENEA/EMFPEEEYQDYEPEA 143
Canary	AKONEEGFLDEGMUNNIGAAVDPONEA/EMPPEEEYÖDYEPEA 143
Chicken	AKONEEGHLDEGMUNNTDIPVDPENEA/EMPPEEEYODYEPEA 143
Mouse	GK-GEEGYPDEGILEDMPVDPGSEAVEMPSEGYODYEPEA 140
Rat	GK-GEEGYPDEGTUED-MPVDPSSEAVEMFSEEGYODYEPEA 140
Bovine	an later a lie Second con Later Appendix to Later 140
	GK I DE GAS DE GTI I EN LIMPANDANE A MEMBER GOVEN DE DE A 140
	GK-GEEGASDEGTUEDMPVDPONEA/EMFSEEGYQDYEPEA 140
Pig	GK-NEEGAPQEGILEDMPVDPQNEAVEMESEEGYQDYEPEA 140
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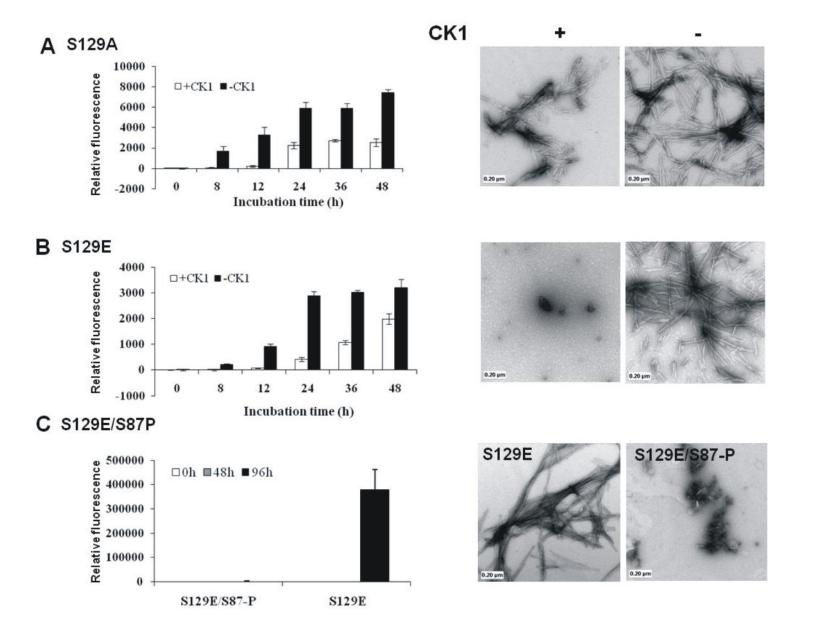
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Rat-α
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Mouse-β
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Rat-β
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Mouse-y
Rat-γ
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Mouse-a
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Human-β
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Mouse-β
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Rat-γ
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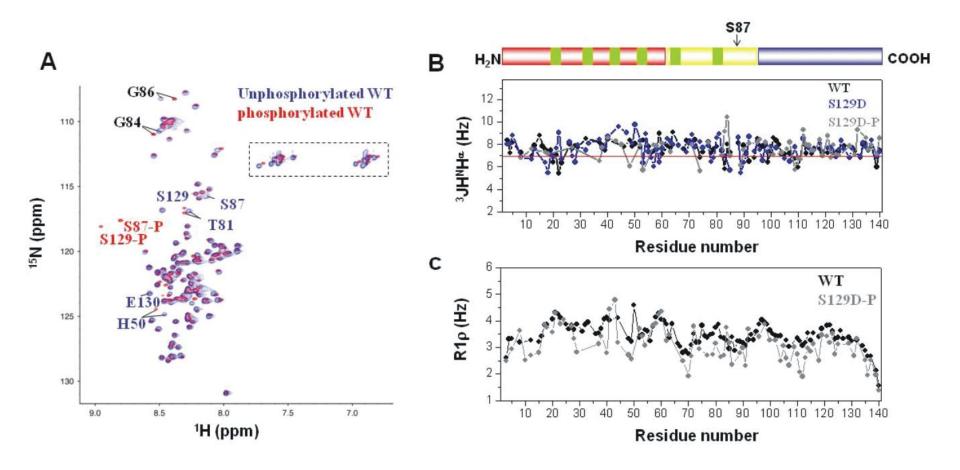
Suppl. Figure 2



Suppl. Figure 3



Suppl. Figure 4



Suppl. Figure 5

