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Title:	Fluorescence Depolarization Kinetics Captures Short-Range Backbone Dihedral Rotations and Long-Range Correlated Dynamics of an Intrinsically Disordered Protein
Authors:	Das, Debapriya (/jspui/browse?type=author&value=Das%2C+Debapriya) Arora, Lishaa (/jspui/browse?type=author&value=Arora%2C+Lishaa) Mukhopadhyay, Samrat (/jspui/browse?type=author&value=Mukhopadhyay%2C+Samrat)
Keywords:	Polarization Magnetic properties
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Abstract:	Intrinsically disordered proteins (IDPs) do not autonomously fold into well-defined three-dimensional structures and are best described as a heterogeneous ensemble of rapidly interconverting conformers. It is challenging to elucidate their complex dynamic signatures using a single technique. In this study, we employed sensitive fluorescence depolarization kinetics by following picosecond time-resolved fluorescence anisotropy decays to directly capture the essential dynamical features of intrinsically disordered α -synuclein (α -syn) site-specifically labeled with thiol-active fluorophores. By utilizing a long-lifetime (≥ 10 ns) anisotropic label, we were able to discern three distinct rotational components of α -syn. The subnanosecond component represents the local wobbling-in-cone motion of the fluorophore, whereas the slower (~ 1.4 ns) component corresponds to the short-range backbone dynamics governed by collective torsional fluctuations in the Ramachandran Φ - Ψ dihedral space. This backbone dihedral rotational time scale is sensitive to the local chain stiffness and slows down in the presence of an adjacent proline residue. We also observed a small-amplitude ($\leq 10\%$) slower rotational correlation time (6–10 ns) that represents the long-range correlated dynamics involving a much longer segment of the polypeptide chain. These intrinsic dynamic signatures of IDPs will provide critical mechanistic underpinnings in a mosaic of biophysical phenomena involving internal friction, allosteric interactions, and phase separation.
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