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In the case of SALS previous studies have revealed only two motifs consisting of a few 10 amino acids, called Wiscott-Aldrich syndrome homology 2 (WH2) domains, that are intrinsically disordered protein regions (IDR) of low structural complexity. Considering their role, they possess actin-binding properties. Depending on the number and sequence of domains, proteins containing WH2 show multifunctional properties.

In our previous research we completed the functional analysis of the SALS WH2 domains (SALS-WH2) [2]. Based on our results both of the SALS WH2 domains interact with the actin, and through their activities shift the monomer:filament ratio towards monomeric actin. We further aimed to characterize the structural and conformational dynamic properties of SALS-WH2 by using in silico and experimental approaches. Our bioinformatics analysis suggests that the SALS-WH2 domains have IDR elements. Our predictionbased results were experimentally verified by fluorescence spectroscopy and thermal analysis.

Keywords: IDP; IDR; WH2; SALS; actin

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Structural and Conformational Dynamics of a Disordered **Protein Motif +**

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SALS (sarcomere length short), a WH2-domain-containing protein was identified in Drosophila as an important regulator of the assembly of sarcomeric actin structures (Bai et al. [1]). It contributes to the establishment of sarcomere length and organization by promoting the lengthening of actin filaments at the pointed end. The absence of SALS is already lethal in the embryonic age. This may be due to the shortening of the length of sarcomeric actin filaments, and/or the disruption of their organization.

SALS is a relatively large protein, consisting of 935 amino acids. According to our bioinformatics analysis, it is an intrinsically disordered protein (IDP). IDPs are biologically active proteins, that, however, do not have a well-defined three-dimensional structure. They possess specific physicochemical properties, different from those of ordered proteins (e.g., hydrophilic/charged:hydrophobic amino acid ratio, thermal stability, electrophoretic mobility).

Abstract

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Conflicts of Interest:

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