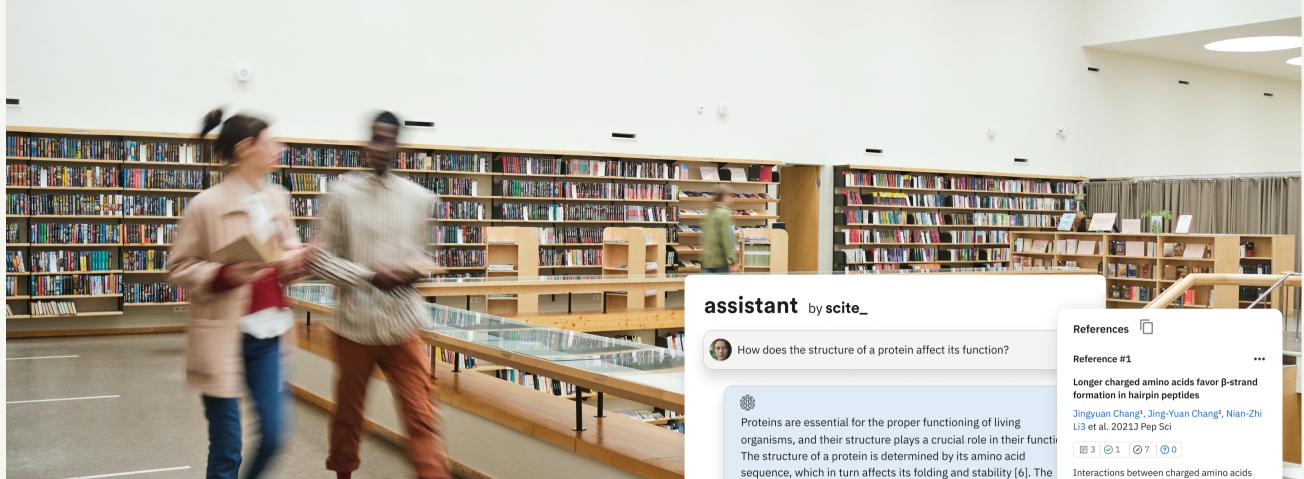
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Interactions between charged amino acids significantly influence the structure and function of proteins. The encoded charged amino acids Asp, Glu, Arg, and Lys have different number of hydrophobic methylenes linking the backbone to the charged functionality...

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Reference #2

Understanding protein non-folding

Vladimir N. Uversky¹, A. Keith Dunker² 2010 Biochimica et Biophysica Acta (BBA) - Proteins and Proteomics

This review describes the family of intrinsically disordered proteins, members of which fail to form rigid 3-D structures under physiological conditions, either along their entire lengths or only in localized regions...

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interactions between charged amino acids significantly influenc

hydrophobic methylenes linking the backbone to the charged

functionality of amino acids affects protein structure stability [1

Protein secondary structures, such as β -sheet structures, are th

structures [1]. Intrinsically disordered proteins (IDPs) are a fam of proteins that fail to form rigid 3-D structures under physiolog

conditions, either along their entire lengths or only in localized

regions [2]. IDPs exist as dynamic ensembles within which atom

positions and backbone Ramachandran angles exhibit extreme

temporal fluctuations without specific equilibrium values [2].

fundamental three-dimensional building blocks of protein

the structure and function of proteins [1]. The number of



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