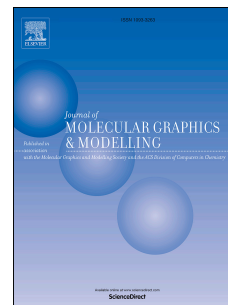


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Loading-device effects on the protein-unfolding mechanisms using molecular-dynamic simulations

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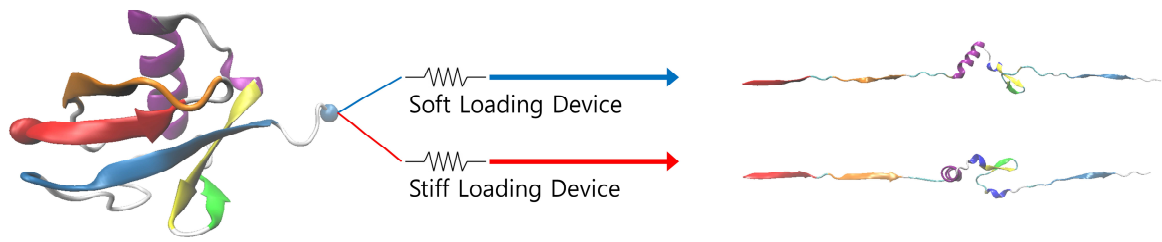
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