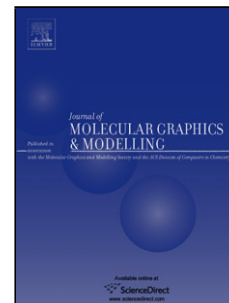


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Sampling conformational space of intrinsically disordered proteins in
explicit solvent: comparison between well-tempered ensemble approach
and solute tempering method

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