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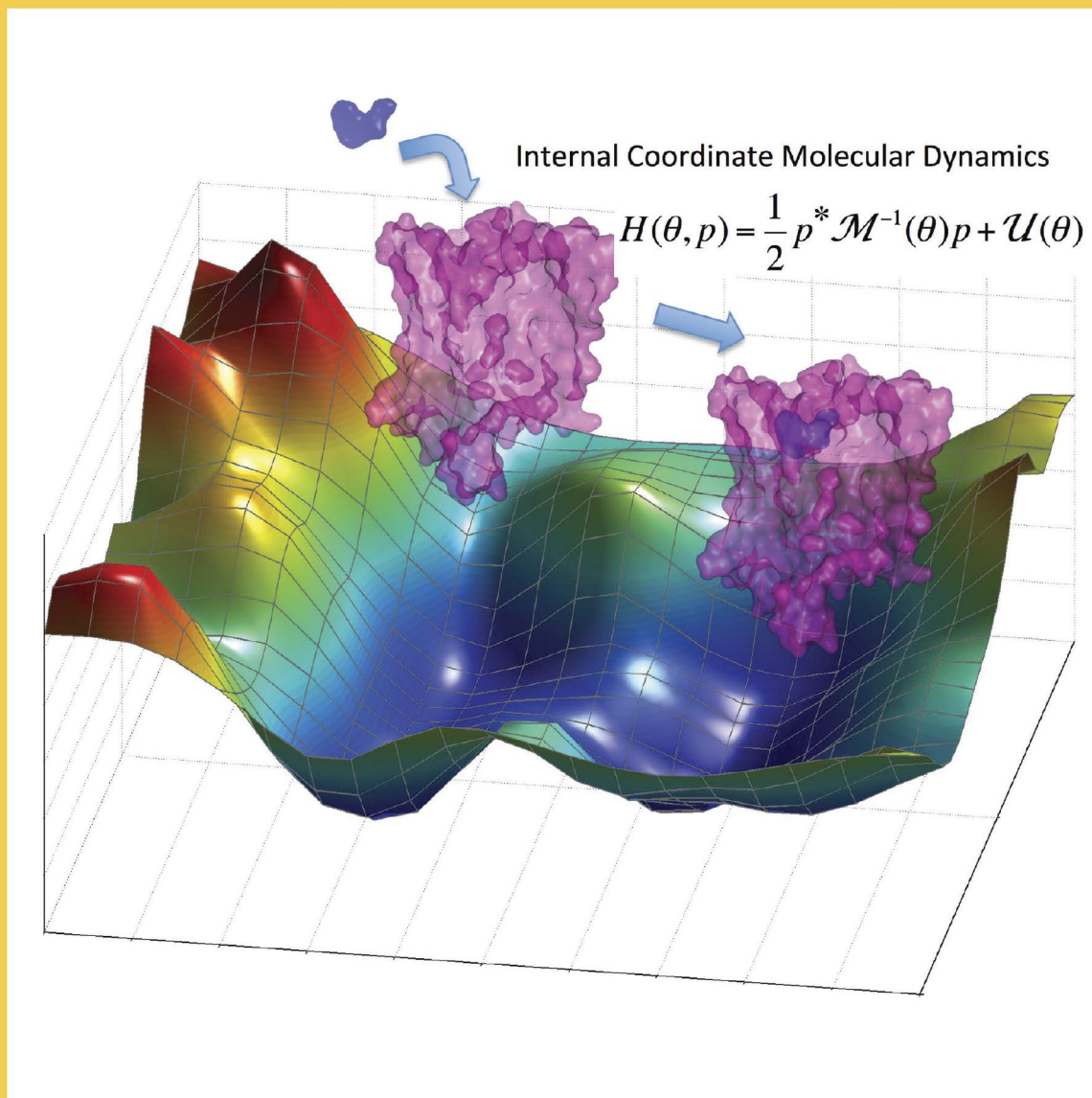
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# THE JOURNAL OF PHYSICAL CHEMISTRY

B

Simulating Large Scale Conformational Changes in Proteins Using GNEIMO, a Multiscale Internal Coordinate Molecular Dynamics Simulation Method (see page 1233)



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