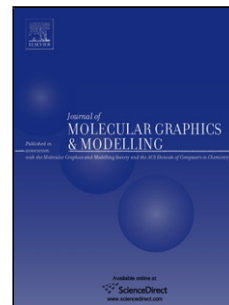


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Homology Modeling of Parasite Histone Deacetylases to Guide the Structure-Based Design of Selective Inhibitors

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