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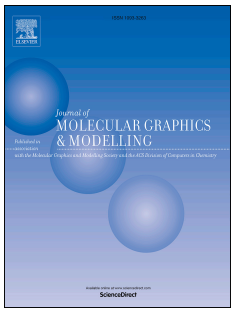

Computational approach for generating robust models for discovering novel molecules as Cyclin Dependent Kinase 4 inhibitors

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