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Author	Sinisa Bjelic	
Title (English)	Computational study of <i>P. falciparum</i> plasmepsin II inhibitor binding	
Title (Swedish)		
Abstract	Prediction of free energy of binding for two inhibitors with four isomers each was performed against malaria parasite <i>Plasmodium falciparum</i> protease plasmepsin II by molecular dynamics simulations and linear interaction energy method. The binding energies were compared further for isomer dependent activity.	
Keywords	Molecular dynamics, linear interaction energy method, malaria	
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