

Time of export: 16.05.2024. 13:50:12

Repository: dabar.srce.hr

Number of records on this URL: 27

Records exported: 27

Title	URL	Authors	Host item title
In Silico Design of Alkhumra Virus NS3 Protease Inhibitors		Novak, Jurica	
Computational and experimental therapeutic efficacy analysis of andrographolide phospholipid complex self-assembled nanoparticles against Neuro2a cells		Novak, Jurica	
Gaussian field-based 3D-QSAR and molecular simulation studies to design potent pyrimidine-sulfonamide hybrids as selective BRAFV600E inhibitors		Novak, Jurica	
MD simulacije novosintetiziranih inhibitora u aktivnom mjestu BRAF proteina		Novak, Jurica	
Virtual screening, structure based pharmacophore mapping, and molecular simulation studies of pyrido[2,3-d]pyrimidines as selective thymidylate synthase inhibitors		Novak, Jurica	
Mukoadhezija farmaceutskih otopina koje sadrže natrijev hijaluronat - utjecaj koncentracije i molekulske mase natrijeva hijaluronata		Strbad, Klara	
Sinteza 131-[N-(2-Aminoetil)maleimid] derivata klorina e6		Rončević, Marina	
Imunogenost lijekova koji sadrže monoklonska protutijela		Lekić, Eleonora	
Sinteza BODIPY spojeva koji ciljaju unutarstanične organele		Vidoša, Lorena	
Sinteza, ciklizacija i karakterizacija katalitičkog peptida CG11		Šimunić, Lorena	
Analysis of potential neuroinflammatory pathology in the spinal cords of mice with optineurin insufficiency		Kolarić, Marta	

Analysis of DNA methyltransferase DNMT1 structure and function and development of noncovalent inhibitors using computational methods		Morović, Viktor	
Structural Modifications Introduced by NS2B Cofactor Binding to the NS3 Protease of the Kyasanur Forest Disease Virus		Kandagalla, Shivananda; Kumbar, Bhimanagoud; Novak, Jurica	
CYP3A4		Novak, Jurica	
Can Resveratrol Influence the Activity of 11 β - Hydroxysteroid Dehydrogenase Type 1? A Combined In Silico and In Vivo Study		Novak, Jurica; Tseilikman, Vadim E.; Tseilikman, Olga B.; Lazuko, Svetlana S.; Belyeva, Lyudmila E.; Rahmani, Azam; Fedotova, Julia	
The design of compounds with desirable properties – The anti-HIV case study		Novak, Jurica; Pathak, Prateek; Grishina, Maria A.; Potemkin, Vladimir A.	
Computational and experimental therapeutic efficacy analysis of andrographolide phospholipid complex self-assembled nanoparticles against Neuro2a cells		Mourya, Atul; Pingle, Purva; Babu, Chanti Katta; Veerabomma, Harithasree; Sainaga Jyothi, Vaskuri G.S; Novak, Jurica; Pathak, Prateek; Grishina, Maria; Verma, Amita; Kumar, Rahul; Singh, Pankaj Kumar; Khatri, Dharmendra Kumar; Singh, Shashi Bala; Madan, Jitender	
Structural modifications introduced by NS2B cofactor binding to the NS3 protease of the Kyasanur forest disease virus		Novak, Jurica	
Gaussian field-based 3D-QSAR and molecular simulation studies to design potent pyrimidine-sulfonamide hybrids as selective BRAFV600E inhibitors		Singh, A. K.; Novak, Jurica; Kumar, A.; Singh, H.; Thareja, S.; Pathak, Prateek; Grishina, M.; Verma, V.; Yadav, J. P.; Khalilullah, H.; Pathania, V.; Nandanwar, H.; Jaremko, M.; Emwas, A.-H.; Kumar, P.	
Računalna studija mehanizmom temeljenih ireverzibilnih inhibitora monoamin oksidaze B		Vrban, Lucija	
Analysis of ADAR protein during productive HSV-1 Infection		Echeta, Justina Oluchi	
Lijekovi protiv herpes simpleks virusa 1 i 2		Krišto, Ruža	
Withasomniferol C, a new potential SARS-CoV-2 main protease inhibitor from the <i>Withania somnifera</i> plant proposed by in silico approaches		Kandagalla, Shivananada; Rimac, Hrvoje; Krishnamoorthy, Gurushankar; Novak, Jurica; Grishina, Maria; Potemkin, Vladimir	
Molekulsko dinamičke simulacije retinala u otopini		Novak, Jurica	
Kiralne stacionarne faze		Novak, Jurica	
Exploring potential inhibitors against Kyasanur forest disease by utilizing molecular dynamics simulations and ensemble docking		Kandagalla, Shivananda; Novak, Jurica; Shekarappa, Sharath Belenahalli; Grishina, Maria A; Potemkin, Vladimir A; Kumbar, Bhimanagoud	

Dinamika modelnih bioloških sustava u osnovnom i pobuđenim elektronskim stanjima		Novak, Jurica	
--	--	---------------	--