

# POSTER PRESENTATION



POSTER NO	AUTHOR NAME	ABSTRACT TITLE
P1	Imran Verirsen	Development Of Novel Fluorescent Tyrosinase Enzyme Sensor And Tyrosinase Activable Photodynamic Therapy Agent
P2	Hasan Tahsin Şen	Creation Of Homology Models Of Some Hiv-1 Integrase Mutations And Investigation Of Their Relationship With Cabotegravir By In Silico Methods
P3	Şeyda Aydoğdu	The Degradation Reaction Mechanism of Ampicilin with Hydroxyl Radical
P4	Gulce Davutlar	Molecular Docking Study of Some Nitro-Substituted Benzamide Compounds as Cyclin-Dependent Kinase Inhibitors and The Anticancer Activity in TNBC cells
P5	Nebahat Sahin	In Silico Determination of Potential Candidates as AXL Kinase Inhibitors
P6	Kamer Nisa BAZ	{In-silico} Study on the Effect of Pro 120 Ser Mutation on NPC-2 Protein
P7	Lalehan Özalp	In silico discovery of potential azole containing mPGES 1 inhibitors by virtual screening, pharmacophore modeling and molecular dynamics simulations
P8	Verda Begüm Tezcan	Tacrine-Hydrazone Compound as Multi-Target Agent for The Treatment of Alzheimer's Disease and Type II Diabetes Mellitus: Design, Synthesis and Bioactivity Studies
P9	Metin Yazar	Pathogenicity prediction of cancer-related variants with protein sequence, structure, dynamics-based features
P10	Aslıhan Özcan	Computational approach for inhibition of ATPases acting antagonistically in Type IV Pilus of Neisseria meningitidis by FDA-approved natural drugs
P11	Melek Gul	Novel Antipyrine Derivatives As Antioxidant Reagents: Synthesis, In-Silico Pharmacokinetics, Molecular Docking Studies, An Implication In Identifying The Nitric Oxide Synthase Inhibition
P12	Derya Yidiz	The combinatorial drug repurposing strategy enhances cytotoxicity in hepatocellular carcinoma
P13	Hazal SARI	Transcriptional regulation of YY1 with a novel G-quadruplex stabilizer
P14	Zeliha Nur YILMAZ	Quantum Chemical, Molecular Docking, Molecular Dynamics and ADMET Studies of Potent Anticancer Diimine-Dioxime Agents
P15	Tolga Corbaci	Discovery of HDAC6 selective inhibitors by deep machine learning methods
P16	Fatma Betul Ertem	Structural studies of IDH1
P17	Muhammad Ammar Zahid	Prediction of CYP450–related pharmacokinetic drug-drug interactions using Graph Convolutional Neural Network
P18	Esmâ Nur YAZ	R448Q Mutation In The Transcription Factor CTCF: Novel Insight Into Structure - Function Relationship From In Silico Analysis
P19	Maide Önder	Towards Selective Stabilization of G-Quadruplex DNA/RNA Structures with Structurally Optimized Ligand
P20	Clara Xazal Buran	Investigation Of The Structural Differences Between Wild-type And Mutant Forms Of MutsA Heterodimer With Molecular Dynamic Simulations
P21	Pınar Erkekoğlu	New Hexahydroquinoline Derivatives And Their Cytotoxic Properties On HePG2 Cells
P22	Harun Nalçakan	Molecular Docking and in Silico ADMET Analyses of Novel Fedratinib Derivatives as Potent JAK2 Inhibitors
P23	Mustafa Filik	AlphaFold Based Pathogenicity Prediction of Missense Variants
P24	İlgaz Taştekil	Investigation of OXA enzyme inhibitors using computational tools
P25	Busra Arvas	Design and Synthesis of New Coumarin-Bistriazole Hybrids as Potential Drug Candidates
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P27	Begüm Nisa Kasaplı	The Development of Antagonists for Knockdown of Axl Overexpression
P28	Gözde Karakadioğlu	Development new generation of imatinib using structural biology techniques at ambient temperature
P29	Gizem Nur Ayan	Synthesis, Purification and Application of Isotopic Derivatives of Various Pharmaceutical Active Ingredients in Analytical Determination Methods
P30	Şura YILMAZ	Synthesis of Bisindolylmaleimide Based Potential Antitumor Compounds
P31	Betul Uzulmez	Investigation of Breast Cancer Cell Responses to the Fullerenol-Dexamethasone Dual Therapy
P32	Mehmet Ali Yucel	Deep Learning Based Drug Repurposing Study for JAK2 Inhibitors
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P34	Rabia Kutlu	Trace Levels in Blood and Urine of Colchicine Active Substance Used in the Treatment of Goutand Behçet's Disease, Also Used in the Treatment of COVID-19, with the HPLC system determinationt
P35	İrem Kulu	Synthesis, pharmacokinetic and biological evaluation of 5-aminoquinoline-triazole hybrid derivatives
P36	Selin Sezer	Molecular Docking Studies Reveal Toxin Binding Differences That Enable Selectivity For Sodium Channels
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P42	Ahmet Çağan	Investigation of Tomentosin β-Amino Alcohol Derivatives Interactions with Topoisomerase II Enzyme Using Molecular Docking Method
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P65	Safa Haddad	Construction of new hERG blocker models based on heteroatom numbers from ultra large ligand libraries
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P67	Anwar Abuelrub	Investigation of the role of CLDNs and TRPs proteins and their associated proteins in peritumoral brain edema (PTBE) by evaluating gliovascular unit (GVU) in disrupted blood brain barrier in glioma IDH 1 mutant vs IDH 1 wild type.
P68	İlayda Tolu	IN SILICO SCREENING OF THE APPROVED DRUGS, PEPTIDOMIMETICS AND DESIGNING OF NEW PEPTIDES AGAINST AXL-GAS6 TARGET