

Target Diversity: Investigating Multi-receptor interactions of antipsychotic drugs through molecular docking

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Psychotic Disorders

- Mental disorders affect thinking, emotions, and behavior
- Mental disorders can be treated by the administration of antipsychotic drugs
- Chlorpromazine, discovered in 1951, was the first effective means for treating delusions and hallucinations (often referred to as positive symptoms) of schizophrenia and other psychotic disorders.
- Major issue: Extrapyrarnidal Side Effects (EPS)

Types of Antipsychotic Drugs

TYPICAL ANTIPSYCHOTIC DRUG

- Also known as first generation antipsychotic drug
- It was introduced between 1954 and 1975.
- High EPS

ATYPICAL ANTIPSYCHOTIC DRUG

- Second generation drugs
- Low EPS
- Better efficacy

ATYPICAL ANTIPSYCHOTIC DRUGS:

Act on

- Serotonin (5HT 2A)
- Dopamine (D2)

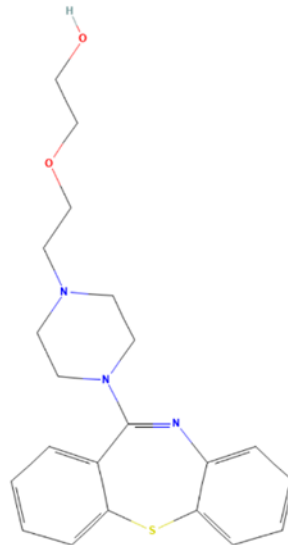
Used as:

- Monotherapy
- Combination therapy

Selected Drugs

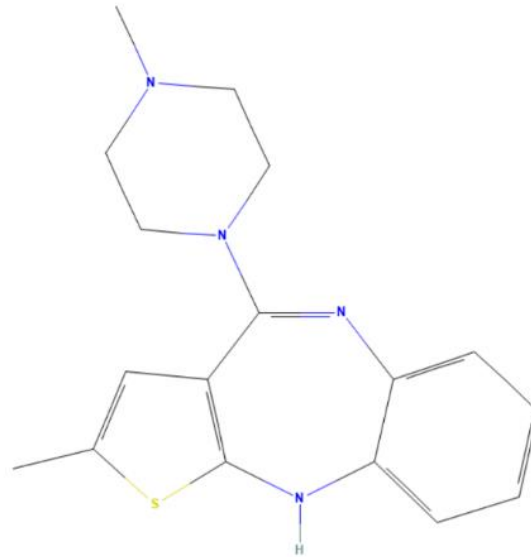
Quetiapine:

A dibenzothiazepine derivative called quetiapine (QTP) is approved for the treatment of bipolar depression



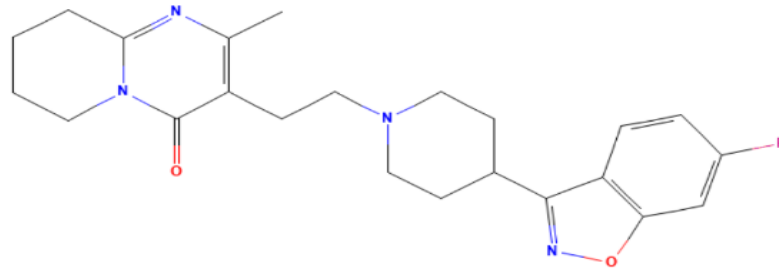
Olanzapine:

Olanzapine (OLZ), a thienobenzodiazepine derivative that exhibits structural similarities with clozapine, is useful in the treatment of acute manic episodes and schizophrenia as well as in preventing the recurrence of bipolar disorders.



Risperidone:

An authorized antipsychotic medication for the treatment of schizophrenia and the acute manic phase of bipolar disorder is risperidone (RSP). It works well for treating positive and negative symptoms of schizophrenia, has a rapid onset of action, and is beneficial in both short- and long-term therapy.



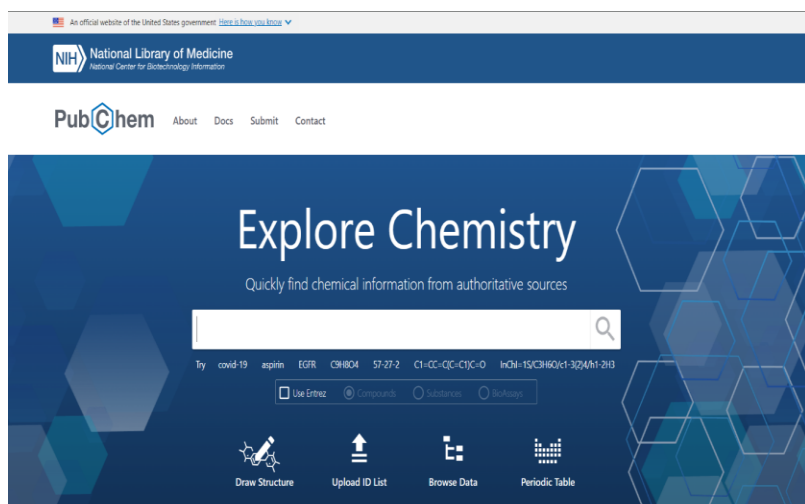
Methodology

TOOLS:

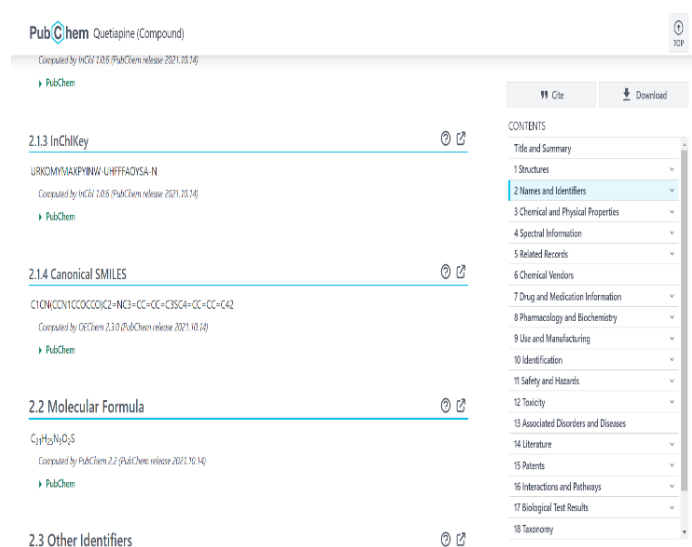
- PubChem
- Swiss ADME
- UniProt
- Maestro Schrödinger

Pubchem

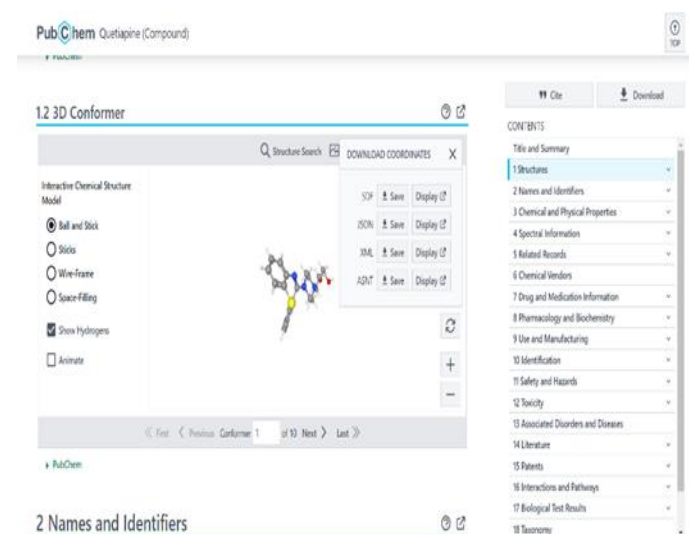
The Canonical SMILE and 3D structure (in SDF format) of the drugs (Quetiapine, Olanzapine, Risperidone) is retrieved from PubChem.



The image shows the PubChem homepage. At the top, it says "An official website of the United States government" and "NIH National Library of Medicine National Center for Biotechnology Information". Below this is the PubChem logo and navigation links: About, Docs, Submit, Contact. The main section is titled "Explore Chemistry" with the subtitle "Quickly find chemical information from authoritative sources". There is a search bar with a magnifying glass icon. Below the search bar, there are several tabs: "Try", "covid-19", "aspirin", "EGFR", "CHB04", "57-27-2", "C1-CC-C(C)-C1C=O", "InChI-15/CH160/c1-3(2)/h1-2h3". There are also buttons for "Use Entrez", "Compounds", "Substances", and "Biologics". At the bottom, there are four icons: "Draw Structure", "Upload ID List", "Browse Data", and "Periodic Table".



The image shows the PubChem page for Quetiapine (Compound). The page title is "PubChem Quetiapine (Compound)". Below the title, it says "Computed by NCBI 2.0.5 (PubChem release 2021.10.14)". There are two buttons: "Cite" and "Download". The page is divided into two main sections: "2.1.3 InChIKey" and "2.1.4 Canonical SMILES". The "2.1.3 InChIKey" section shows the InChIKey "URKDMYMAKPWW-UHFFFAOYSA-N" and the text "Computed by NCBI 2.0.5 (PubChem release 2021.10.14)". The "2.1.4 Canonical SMILES" section shows the Canonical SMILES "C1CN(CCN1CCOCCO)C2=NC3=CC=CC=C3C4=CC=CC=C42" and the text "Computed by OEChem 2.3.0 (PubChem release 2021.10.14)". There is also a "2.2 Molecular Formula" section showing "C₂₇H₃₀N₂O₂S" and the text "Computed by PubChem 2.2 (PubChem release 2021.10.14)". On the right side, there is a "CONTENTS" menu with a list of sections: "Title and Summary", "1 Structures", "2 Names and Identifiers", "3 Chemical and Physical Properties", "4 Spectral Information", "5 Related Records", "6 Chemical Vendors", "7 Drug and Medication Information", "8 Pharmacology and Biochemistry", "9 Use and Manufacturing", "10 Identification", "11 Safety and Hazards", "12 Toxicity", "13 Associated Disorders and Diseases", "14 Literature", "15 Patents", "16 Interactions and Pathways", "17 Biological Test Results", and "18 Taxonomy".



The image shows the PubChem page for Quetiapine (Compound) with the "3D Conformer" section selected. The page title is "PubChem Quetiapine (Compound)". Below the title, it says "Computed by NCBI 2.0.5 (PubChem release 2021.10.14)". There are two buttons: "Cite" and "Download". The "3D Conformer" section shows a 3D ball-and-stick model of the Quetiapine molecule. There are buttons for "Download Coordinates" and "Display" for different formats: "SDF", "JSON", "MOL", and "ASCI". The "2 Names and Identifiers" section is also visible on the right side of the page.

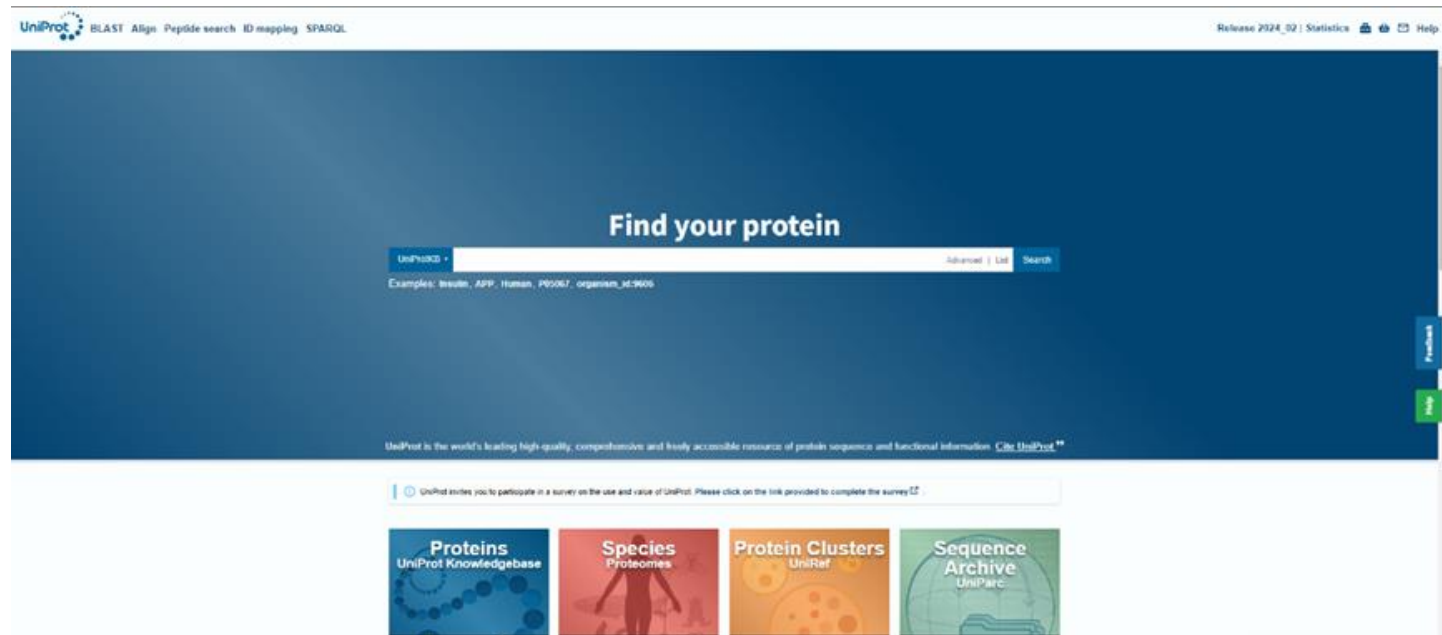
SwissADME:

- The Canonical SMILE of Quetiapine, Olanzapine, and Risperidone is imported in SwissADME to get the ADME properties of the respective drugs.

The screenshot displays the SwissADME web application interface. At the top, a navigation bar includes links for SwissDrugDesign, SwissDock, SwissParam, SwissSidechain, SwissBioisostere, SwissTargetPrediction, **SwissADME**, SwissSimilarity, and About us. The main header features the SIB logo (Swiss Institute of Bioinformatics) and the title "SwissADME". Below the header, a text block explains the website's purpose: "This website allows you to compute physicochemical descriptors as well as to predict ADME parameters, pharmacokinetic properties, druglike nature and medicinal chemistry friendliness of one or multiple small molecules to support drug discovery." It also provides references for the underlying methodologies: iLOGP, BOILED-Egg, and BOILED-Score. A survey prompt asks users to provide feedback. The central area is divided into two main sections. On the left, a "Marvin JS by ChemAxon" logo is displayed within a chemical editor interface. On the right, a text input field is labeled "Enter a list of SMILES here:" and contains the Canonical SMILES for Quetiapine: C1CN(CCN1CCOCCO)C2=NC3=CC=CC=C3SC4=CC=CC=C42CC1=CC2=C(S1)NC3=CC=CC=C3N=C2N4CCN(CC4)C. Below the input field, there are buttons for "Fill with an example", "Clear", and "Run!". At the bottom, a "Show BOILED-Egg" button is visible on the left, and a "Retrieve data:" section with a CSV icon and "POWERED BY ChemAxon" logo is on the right.

UniProt:

The Electron Microscopic structure of the receptor 5-HT1A (7E2X), 5-HT2B (7SRQ), 5HT6 (8JLZ), D1 (7JVQ), D2 (7JVR), D3 (8IRT), α 2A (7EJ8), α 2B (6K41), GABA subunit alpha-1 (7QNE) , H1 (7DFL), H2 (7UL3) and x-ray structure of 5-HT1B (4IAR), 5-HT2A (7WC8), GABA TYPE SUB-2 (6WIV) are retrieved from UniProt as 3D structures in pdf format.



Maestro Schrodinger

- Initially, ligand preparation was done using the Ligprep tool.
- LigPrep is designed to rapidly generate high-quality small-molecule ligand structures for structure-based virtual screening
- It functions by explaining stereoisomers, ring conformation, and tautomeric and ionization states.
- The protein (receptor) retrieved from PDB was imported and protein preparation was done by protein preparation wizard tool.
- A first-in-class, structure-based drug development effort faces several early challenges, one of which is locating druggable pockets.

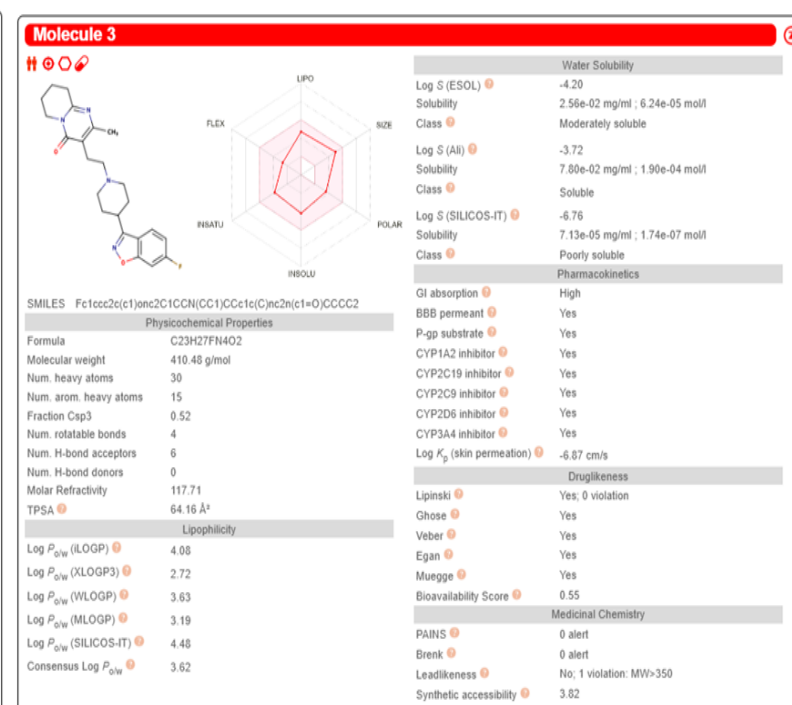
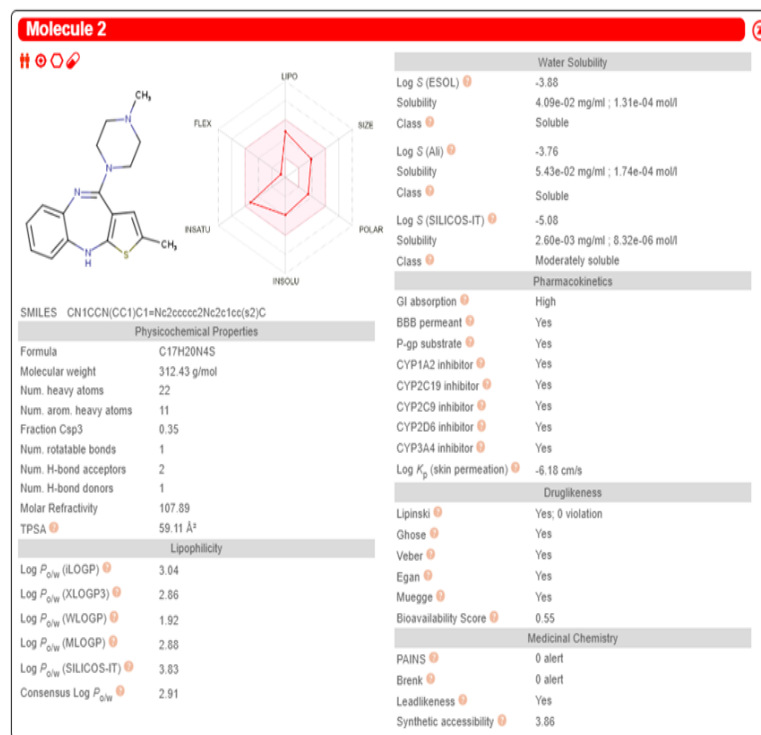
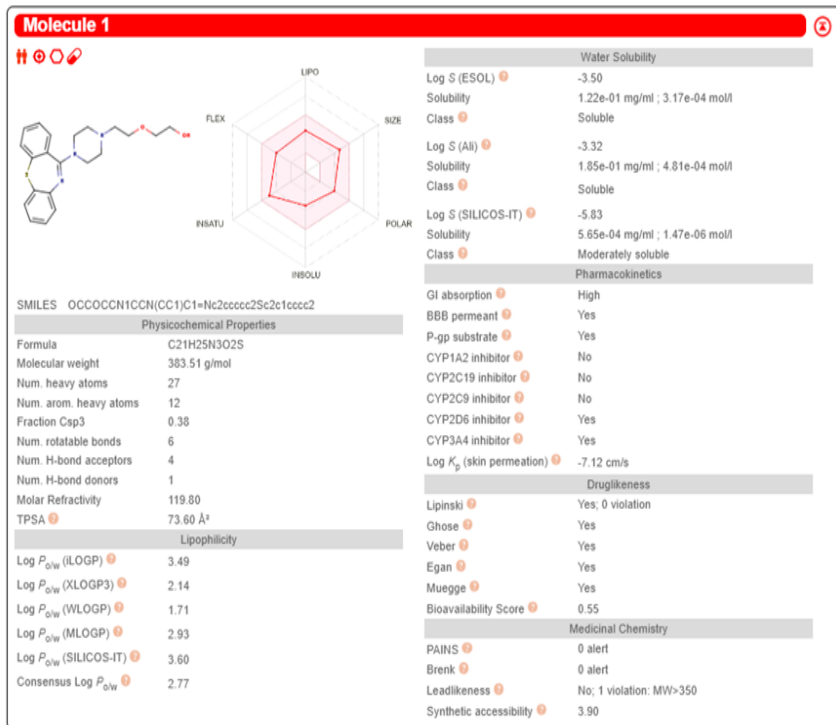
- The binding sites and drugabilities of protein-protein interfaces and allosteric binding sites can be identified with the aid of the SiteMap technique.
- Grid generation must be performed before running a virtual screen with glide.
- To define the grid surrounding the ligand and retain all the functional residues within it, the receptor grid generation technique applied.
- Finally, the protein-ligand docking was done and the docking was done

Results

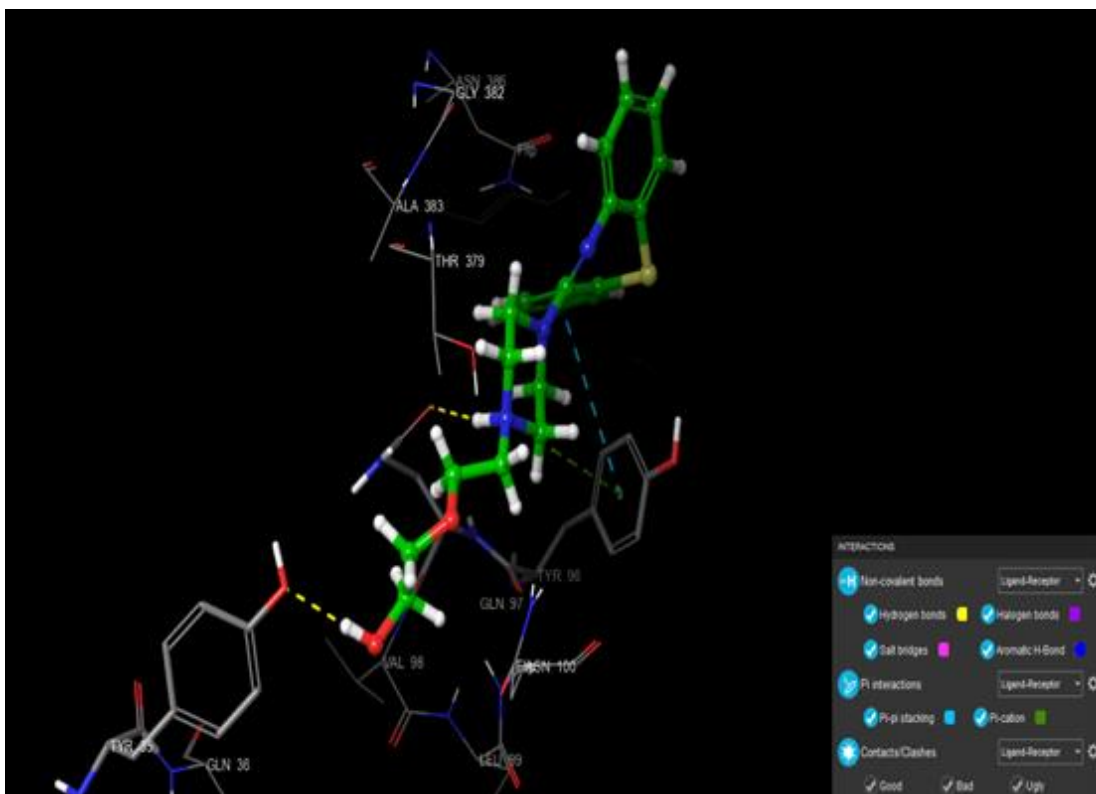
- Drugs and their canonical SMILES

S.No	Name of the Drug	Drug Type and status	Canonical SMILE	Punchem ID
1.	Quetiapine	Atypical antipsychotic drug, Approved by FDA	<chem>C1CN(CCN1CCOCCO)C2=NC3=CC=CC=C3SC4=CC=CC=C42</chem>	5002
2.	Olanzapine	Atypical antipsychotic drug, Approved by FDA	<chem>CC1=CC2=C(S1)NC3=CC=CC=C3N=C2N4CCN(CC4)C</chem>	135398745
3.	Risperidone	Atypical antipsychotic drug, Approved by FDA	<chem>CC1=C(C(=O)N2CCCCC2=N1)CCN3CCC(CC3)C4=NOC5=C4C=CC(=C5)F</chem>	5073

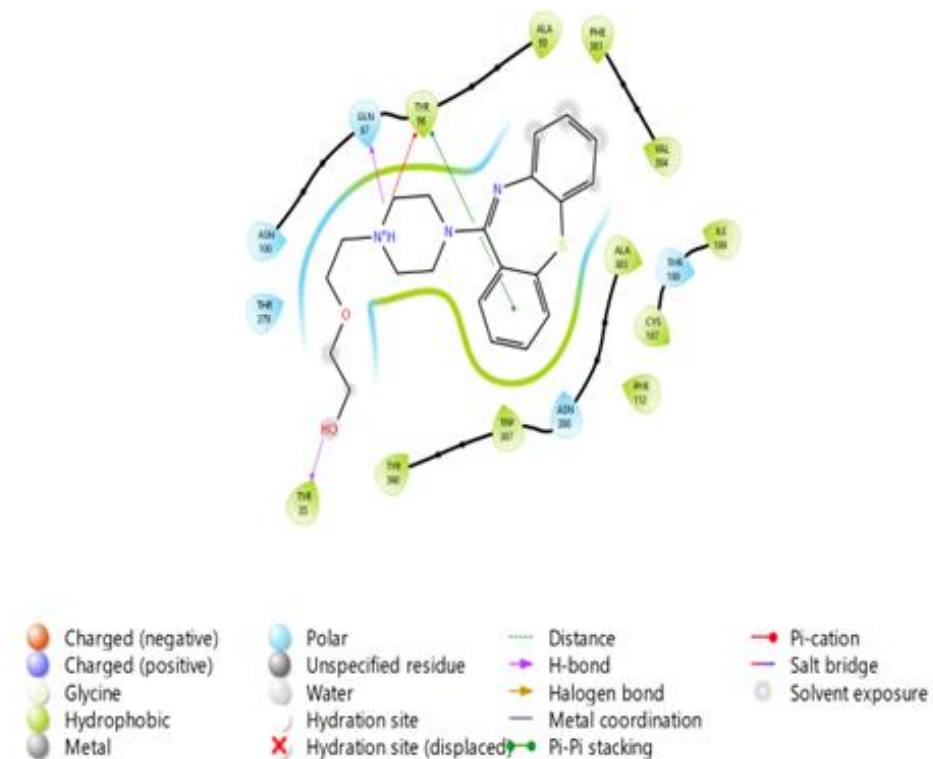
ADME properties of Quetiapine, Olanzapine and Risperidone



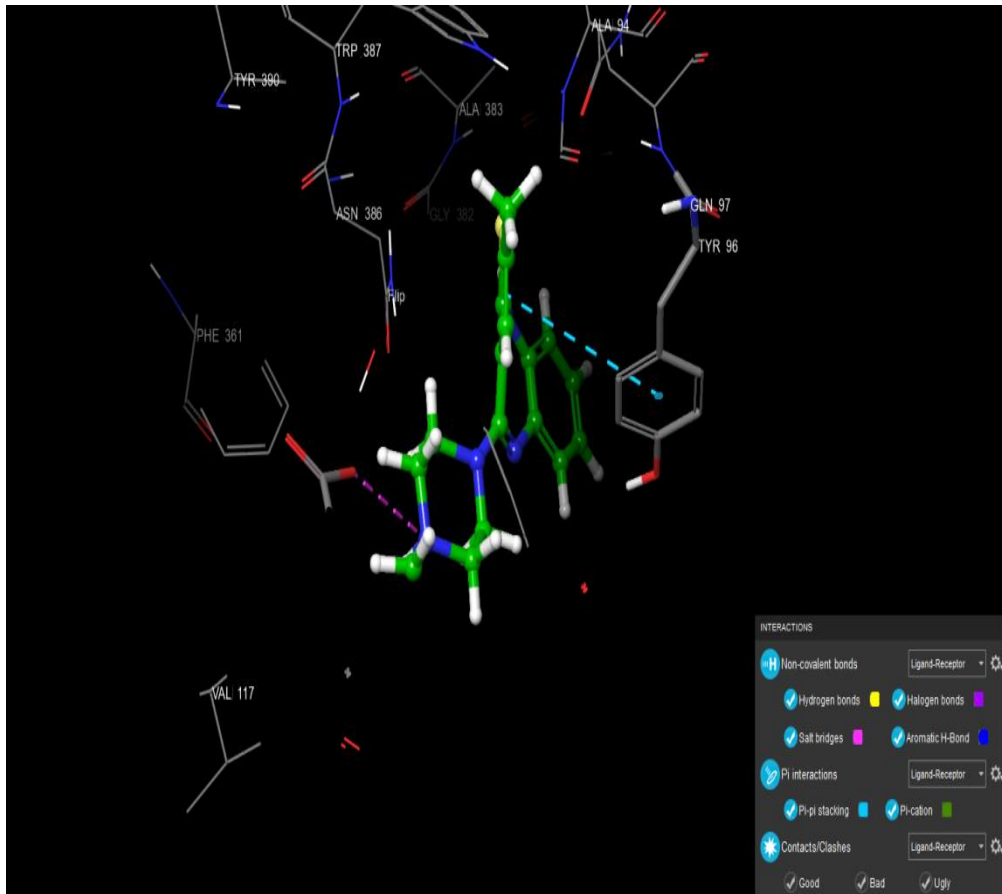
Molecular Docking of 5-HT1A with Quetiapine, Olanzapine, and Risperidone respectively



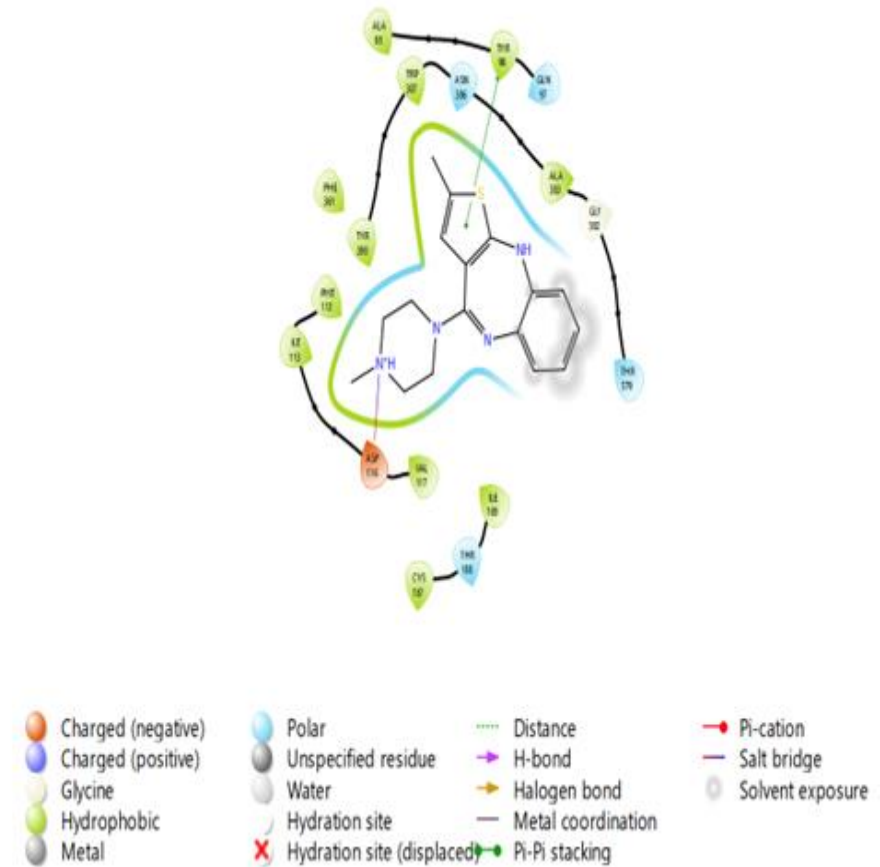
docking of Quetiapine with 5-HT1A



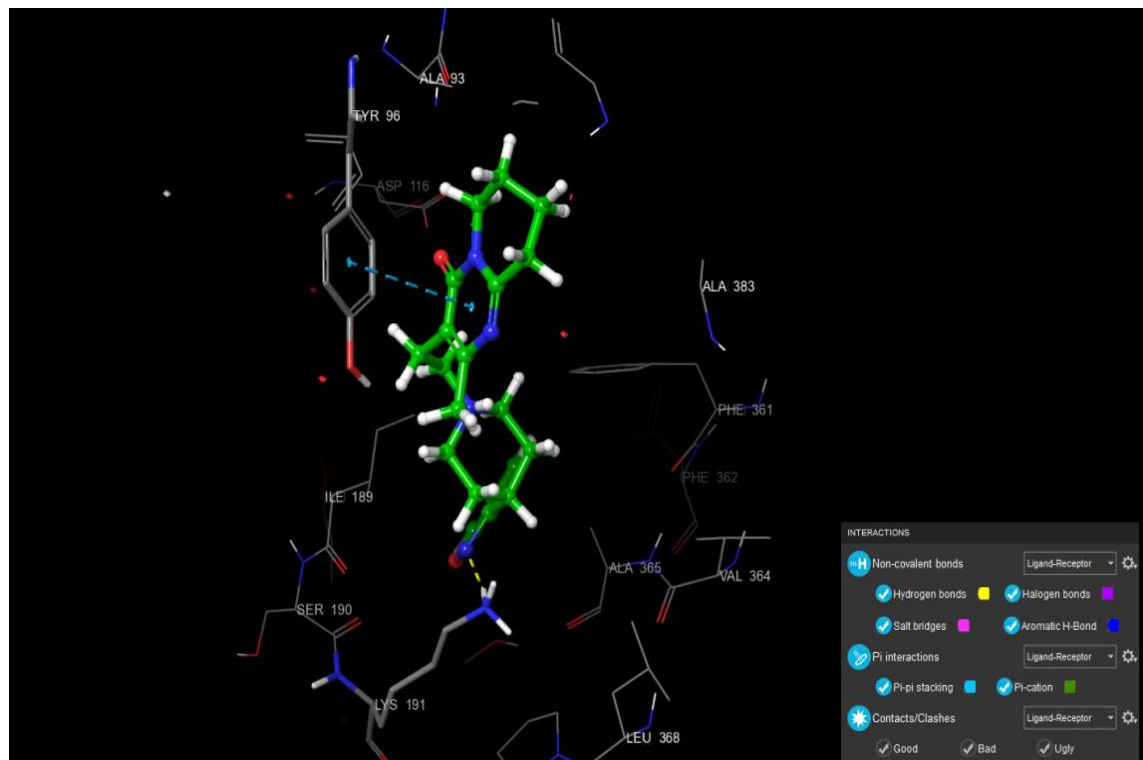
ligand interaction of Quetiapine with 5-HT1A



docking of Olanzapine with 5-HT1A



ligand interactions of Olanzapine with 5-HT1A



docking of Risperidone with 5-HT1A



ligand interaction of

Risperidone with 5-HT1A

Docking score:

- Quetiapine:

1	5-HT1A(7E2X)	-5.231	TYR 96, GLN 97
2	5-HT1B(4IAR)	-5.445	ASP 352
3	5-HT2A(7WC8)	-6.720	ASP 155
4	5-HT2B(7SQR)	-6.988	ASP 135, PHE 341
5	D1(7JVQ)	-5.487	SER 275, SER 147
6	α_{2A} (7EJ8)	-5.004	ASN A: 346, 347, ASP A: 262, ARG A:349
7	α_{2B} (6K41)	-5.183	ARG B: 22, GLN B: 259, LYS B: 15
8	GABA SUB α -1(7QNE)	-6.292	HIS A: 56, GLU E: 152, GLN G 53
9	H1(7DFL)	-6.565	HIS A: 218

Olanzapine:

S.No.	Recptor	Docking score Kcal/mol	Amino acids involved in interactions
1	5-HT1A(7E2X)	-5.525	ASP 116, TYR 96
2	5-HT1B(4IAR)	-7.096	ASP 129, PHE 331, SER 212
3	5-HT2A(7WC8)	-6.462	ASP 155
4	5-HT2B(7SQR)	-8.136	ASN 344, ASP 133
5	5-HT6(8LZ)	-6.706	ASP 106
6	D1(7JVQ)	-5.668	ILE 232
7	D2(7JVR)	-6.341	ASP 114
8	α_{2A} (7EJ8)	-4.809	ASN 346
9	α_{2B} (6K41)	-4.113	ASP B: 205, LYS B: 209
10	GABA SUB α -1(7QNE)	-7.689	ILE E: 47, MET A: 141

Risperidone:

S.No	Receptor	Docking score kcal/mol	Amino acids involved in interactions
1	5-HT1A(7E2X)	-7.043	LYS 191, TYR 96
2	5-HT1B(4IAR)	-7.386	ASP 129
3	5-HT2B(7SQR)	-8.657	ASP 135, PHE 341, TRP 131
4	D2(7JVR)	-6.944	ASP 114, PHE 390, TYR 408
5	α_{2A} (7EJ8)	-4.600	ASP R: 447, HIE R: 446
6	α_{2B} (6K41)	-5.523	ARG B: 19, LYS B: 209
7	GABA Type B sub-2 (6WIV)	-7.793	TYR 667
8	GABA SUB α -1(7QNE)	-8.005	MET E: 49
9	H1(7DFL)	-6.401	ARG B: 150, ASN B: 230, HIS A: 218
10	H2(7UL3)	-6.909	ASP 98, VAL 176, PHE 254, PH 251

Conclusion:

The docking between the drug and the respective receptors is done. The drugs involved in this experiment bind to multiple receptors and there are variations in binding affinity when a drug binds with different receptors. Quetiapine shows high affinity towards the receptors 5-HT_{2A}, 5-HT₂₈, GABRA1, and H1 with a docking score (Kcal/mol) of -6.720, -6.988, -6.292, and -6.565 respectively. Olanzapine shows high affinity towards the receptors 5-HT_{2A}, 5-HT₁₈, 5-HT₂₈, 5-HT₆, D1, and GABRA1 with a docking score of -6.462, -7.096, -8.136, -6.706, -6.341, and -7.689 respectively. Risperidone shows high affinity towards the receptors 5-HT₂₈, 5-HT_{1B}, GABBR2, and GABRA1 with a docking score of -7.386, -8.657, -7.793 and -8.005 respectively. Molecular docking holds great promise for accelerating the drug discovery process and addressing complex biomedical challenges. However, further in vitro, and in vivo study is important for analyzing the drug properties.