

NEW TRENDS AND STRATEGIES IN THE CHEMISTRY OF ADVANCED MATERIALS WITH RELEVANCE IN BIOLOGICAL SYSTEMS, TECHNIQUE AND ENVIRONMENTAL PROTECTION

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LYS A:102

ZINC38321654

gscore = -9.917

MM-GBSA(ΔG_{bind})=-64.88

NATURAL PRODUCTS AS ANTI-HIV AGENTS: AN IN SILICO INSIGHT

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Introduction

In the present work, in silico approaches were applied to find natural products (NPs) with similar bioactivity to non-nucleoside reverse transcriptase inhibitors (NNRTIs) but with different chemotypes, as promising candidates to treat human immunodeficiency virus type 1 (HIV-1) infection. The FDA-approved drug, etravirine, was used as reference molecules in a virtual screening experiment. The ZINC15 database of 224205 NPs was screened and filtered using 3D-similarity search, ADMET, and molecular docking simulations. Three NPs having higher docked scores and superior ADMET profile than etravirine were selected for further investigations. For these NPs, hydrogen bonds and hydrophobic interactions with key binding site residues (Lys101, Tyr181, Tyr188, Trp229, and Tyr318), along with free binding energies, argue that ligands can bind to HIV-1 reverse transcriptase. In conclusion, these three compounds were proposed as potential anti-HIV inhibitors. Moreover, our proposed workflow might be helpful to design novel potential NNRTIs from natural sources.

Results and Discussions

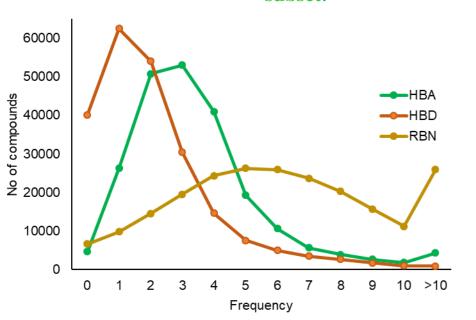
224205 **QUERY Natural Products** Etravirine ZINC15 database ADMET 3D-similarity search **Drug likeness 18 Natural Products** Glide Docking and Prime MM-GBSA **Results Analysis** Free energy of binding, RMSD, Interactions,

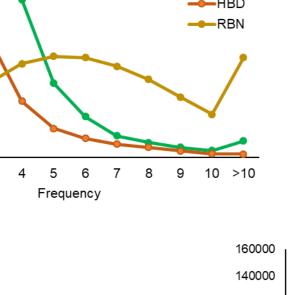
Percent inhibition prediction of selected compounds

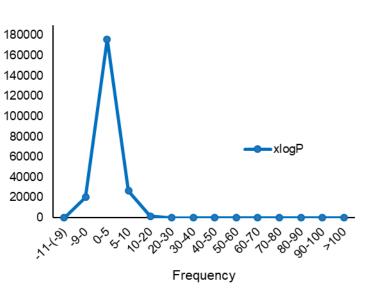
Workflow

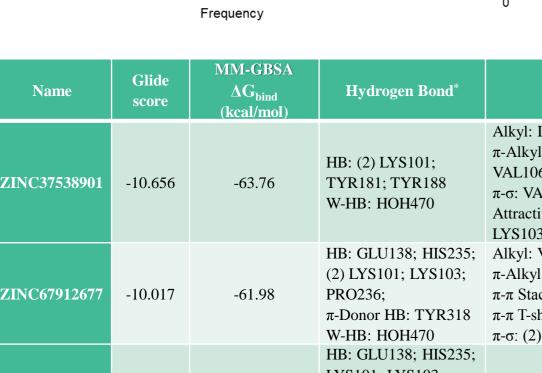
- > To expand the chemical space of NPs, the 3D similarity search was employed.
- ➤ The FDA-approved drugs, **ETRAVIRINE** was designated as query in the current
- investigation. ETRAVIRINE, a Non-nucleoside reverse transcriptase inhibitor (NNRTI), has been designed to be active against HIV1-RT.
 - > The Food and Drug Administration (FDA) approved its use for patients with established resistance to other drugs, in January 2008.

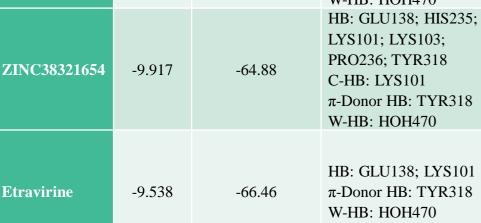
> Distribution of drug-like properties for the ZINC15 NPs









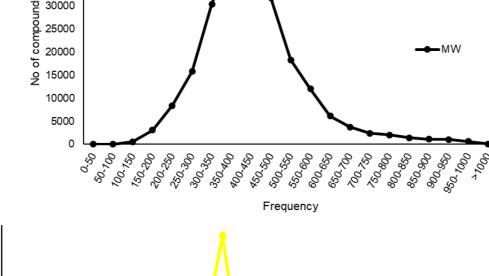


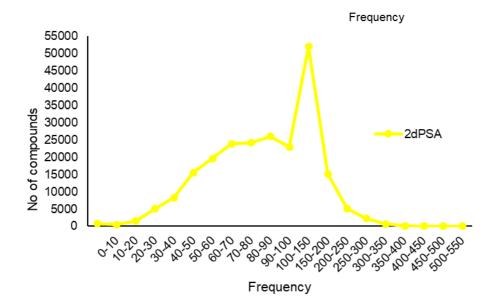
 π -π Stacked: TYR188 π -π T-shaped: (2) TRP229 π-σ: LEU100; VAL179; TRP229

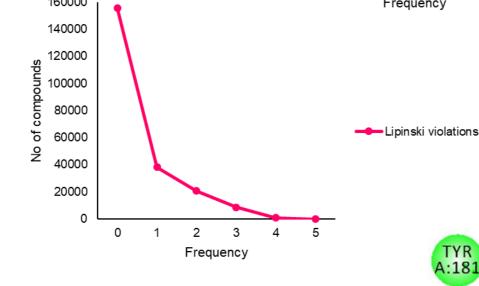
Known therapeutic benefits of the proposed natural products

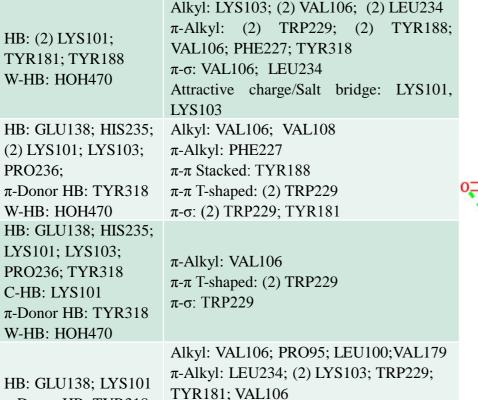
40000 35000

25000 20000 ≥ 15000 10000

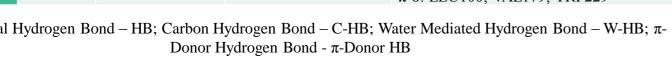






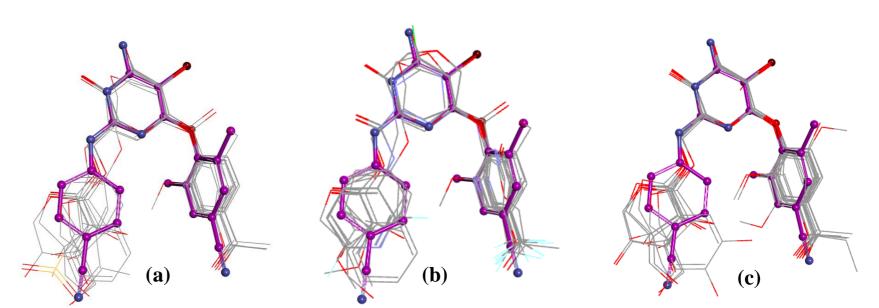


Hydrophobic interactions



- ✓ **ZINC37538901** is a natural derivative of β -D-glucopyranoside type known to develop anticancer, anti-inflammatory, antiseptic and many other activities.
- ✓ Canthoside D (ZINC038321654), a phenolic compound isolated from aerial parts of Salsola tetragona specie, is known to possess anticancer, antimicrobial, anti-inflammatory, antioxidant, antidepressant, and antihypertensive activities.
- ✓ Geoside (ZINC67912677), one out of more than 30 steviol glycosides, is a natural sweetness compound extracted from Stevia rebaudiana leaves. The steviol glycosides are used in food industry, especially as sweeteners in fruit juices. They also exhibit antinflammatory, antibacterial, antiviral, antitumor, antihyperglycemic, antioxidant activities, etc. In short, stevia has zero calories, many benefits. Like most natural compounds, they are safe for human health and could be consumed without restriction by diabetics.

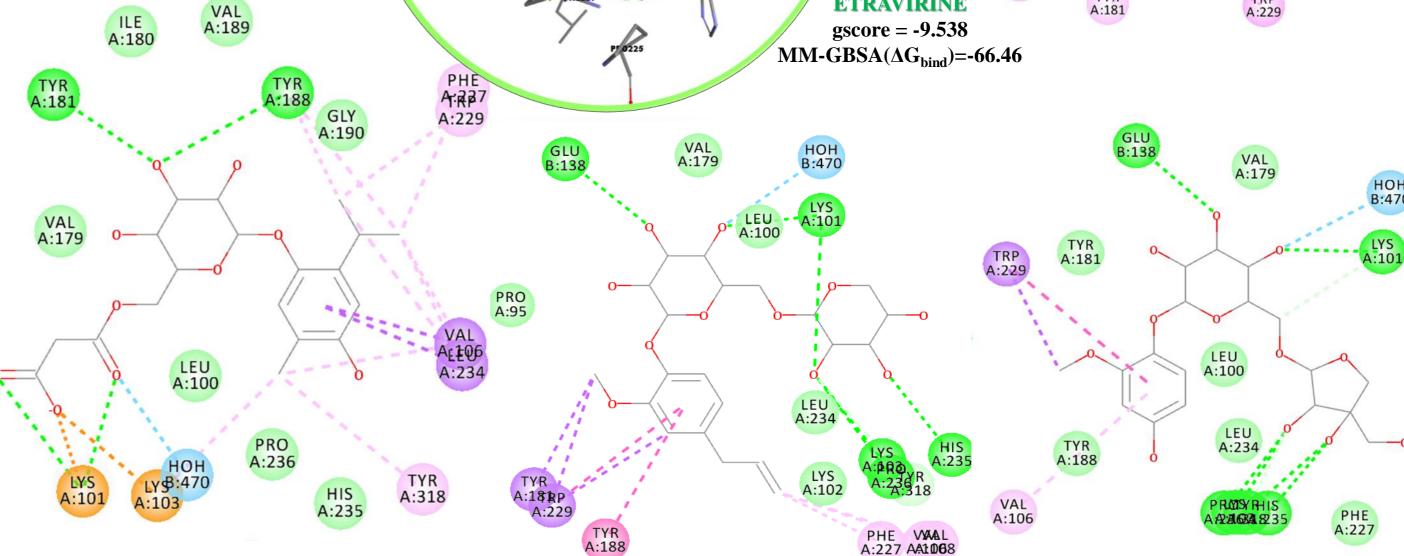
- > The TanimotoCombo (TC), ShapeTanimoto (ShT), and ComboScore (CS) were selected as scoring parameters to rank NPs with respect to etravirine.
- > The top ten molecules aligned by ROCS using the criteria: TC values (a), ShT values (b), CS values (c)



3D structure of Etravirine is depicted in magenta

■ The very low RMSD value of 0.695Å validates the docking procedure.

- The RMSD between the best docking pose of Etravirine and its X-ray structure coordinates, was calculated.
- VAL A:106 A:103 A:234 VAL A:179 TYR A:181 **ETRAVIRINE** A:229 gscore = -9.538MM-GBSA(ΔG_{bind})=-66.46 A:227



*Conventional Hydrogen Bond – HB; Carbon Hydrogen Bond – C-HB; Water Mediated Hydrogen Bond – W-HB; π-

Conclusions

❖ 3D-similarity search, ADMETox, molecular docking, and MM-GBSA simulations were performed within 224205 natural compounds of the ZINC15 database to identify potential new antiviral agents against HIV-1 RT.

ZINC67912677

gscore = -10.017

MM-GBSA(ΔG_{bind})=-61.98

- ❖ The *in-silico* analysis revealed that three (ZINC37538901, ZINC67912677, and ZINC38321654) out of twenty five selected natural products fulfilled all the parameters investigated such as 3D-similarity coefficients, ADMETox parameters, the IC50/percent inhibition of HIV RT protein, docking scores, and free binding energies.
- **❖** Docking outcomes suggested that residues Lys101, Tyr181, Tyr188, Trp229, and Tyr318, involved in essential hydrogen bonding and π-π stacked interaction which stabilized ZINC NP in HIV-1 RT active site, played essential roles for anti-HIV activity.
- **Concerning the antiviral activity evaluated by HIVprotI, all ZINC NPs were predicted to show activity against** HIV-1 RT with an IC50 range of 2-99.86μM and 27.65-52.46% inhibition.

References

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ZINC37538901

gscore = -10.656

MM-GBSA(ΔG_{bind})=-63.76

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