

WEBLEM 2**BIOVIA Draw software & Open Babel Tool****Aim:**

Introduction to Chemical Structure (1D, 2D and 3D), Drawing using BIOVIA DRAW Software and File conversion using Open Babel Tool

Introduction:**Chemical Structure:**

A drug's chemical composition impacts its physicochemical characteristics, which in turn influence its ADME/Tox characteristics, which in turn influence its pharmacological action. By changing the shape of drug molecules, medicinal chemists can control the pharmacological activity of a given substance. A drug's ring systems and functional groups are crucial parts. The percentage of heavy atoms in a medicine is determined by the ratio of non-hydrogen to non-hydrocarbon atoms. The three variables have a lot of potential for determining whether organic compounds have drug-like characteristics. To the best of our knowledge, however, no studies have been done to systematically examine the simultaneous impacts of the quantity of aromatic and non-aromatic rings, the quantity of specific functional groups, and the fraction of heavy atoms on the drug-like qualities of an organic molecule.

1. BIOVIA Draw Software:**History:**

Accelrys was formed in 2001 as a wholly owned subsidiary of Pharmacia, Inc. from the fusion of five companies: Molecular Simulations Inc., Synopsys Scientific Systems, Oxford Molecular, the Genetics Computer Group (GCG), and Synomics Ltd. MSI, itself a result of the combination of Biodesign, Cambridge Molecular Design, Polygen and, later, Biocad and Biosym Technologies.

Introduction:

BIOVIA is a software company headquartered in the United States, with representation in Europe and Asia. It provides software for chemical, materials and bioscience research for the pharmaceutical, biotechnology, consumer packaged goods, aerospace, energy and chemical industries. Previously named Accelrys, it is a wholly owned subsidiary of Dassault Systèmes after an April 2014 acquisition and has been renamed BIOVIA.

BIOVIA Draw has the same look-and-feel as ISIS/ Draw, but brings additional speed and efficiency to chemical structure drawing:

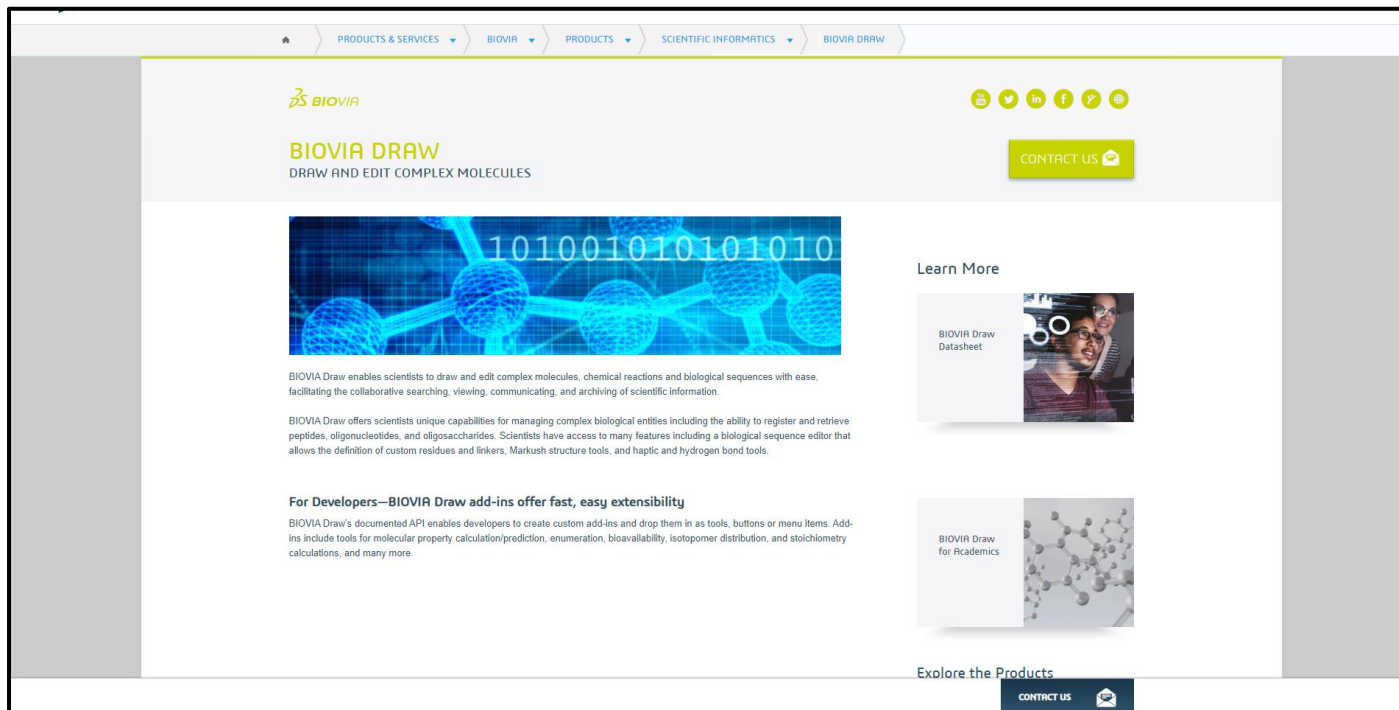
- Continuously draw bonds, pull out rings and add atoms using all-purpose drawing tool
- Drag-and-drop commonly-used structures and chemical abbreviations onto the toolbar for reuse
- Right-click for atom, bond, fragment properties and query options
- Hover over atoms and edit them without right-click
- Quickly retrace steps using Multiple Undo/Redo
- Easily create structures with R groups for queries or enumerations
- Annotate reaction schemes with text, colour and a variety of arrow styles
- Easily create publication quality structures for inclusion in Microsoft Office Documents and presentations.

Applications:

Simulate, visualize and analyse chemical and biological systems and to communicate the results to other scientists.

Installation Steps:

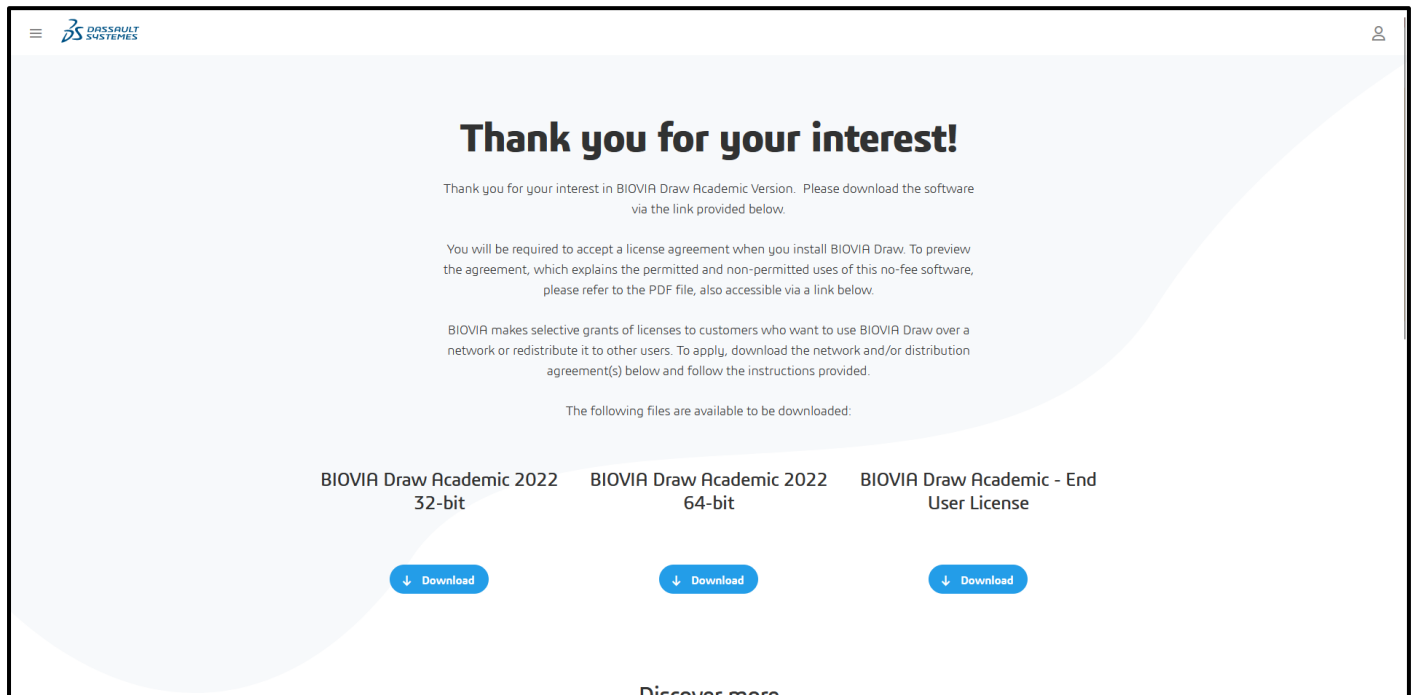
1. Open Homepage of BIOVIA DRAW software website and then click on BIOVIA Draw for Academics (URL: <https://www.3ds.com/products-services/biovia/products/scientific-informatics/biovia-draw/>).



2. Fille the registration details and click on submit

The screenshot shows the "BIOVIA Draw for Academics" registration page. The left side features a large image of a molecular structure and the text "BIOVIA Draw for Academics" and "Free access for academics and non-commercial users". Below this, it says "Draw and edit complex molecules" and "Students, teachers and researchers in an academic settings can—at no charge—download BIOVIA Draw for rapidly drawing chemical structures and chemically intelligent queries." A link "Register to download the installer." is provided. The right side has a "Register now" section. It includes a link for non-gmail users, a progress indicator "STEP 2 OF 2", and several input fields: "First Name*", "Last Name*", "Company*", a country dropdown menu (set to "India"), "City*", and a phone number field (set to "+91 81234 56789"). A checkbox for consent is present, and a blue "Submit" button is at the bottom.

3. Download BIOVIA DRAW Academic 2022



The screenshot shows the BIOVIA Draw Academic 2022 download page. At the top left is the Dassault Systèmes logo. The main heading is "Thank you for your interest!". Below it, a paragraph says: "Thank you for your interest in BIOVIA Draw Academic Version. Please download the software via the link provided below." Another paragraph states: "You will be required to accept a license agreement when you install BIOVIA Draw. To preview the agreement, which explains the permitted and non-permitted uses of this no-fee software, please refer to the PDF file, also accessible via a link below." A third paragraph mentions: "BIOVIA makes selective grants of licenses to customers who want to use BIOVIA Draw over a network or redistribute it to other users. To apply, download the network and/or distribution agreement(s) below and follow the instructions provided." Below this, it says: "The following files are available to be downloaded:". There are three download links, each with a "Download" button: "BIOVIA Draw Academic 2022 32-bit", "BIOVIA Draw Academic 2022 64-bit", and "BIOVIA Draw Academic - End User License". At the bottom right, there is a "Discover more" link.

Thank you for your interest!

Thank you for your interest in BIOVIA Draw Academic Version. Please download the software via the link provided below.

You will be required to accept a license agreement when you install BIOVIA Draw. To preview the agreement, which explains the permitted and non-permitted uses of this no-fee software, please refer to the PDF file, also accessible via a link below.

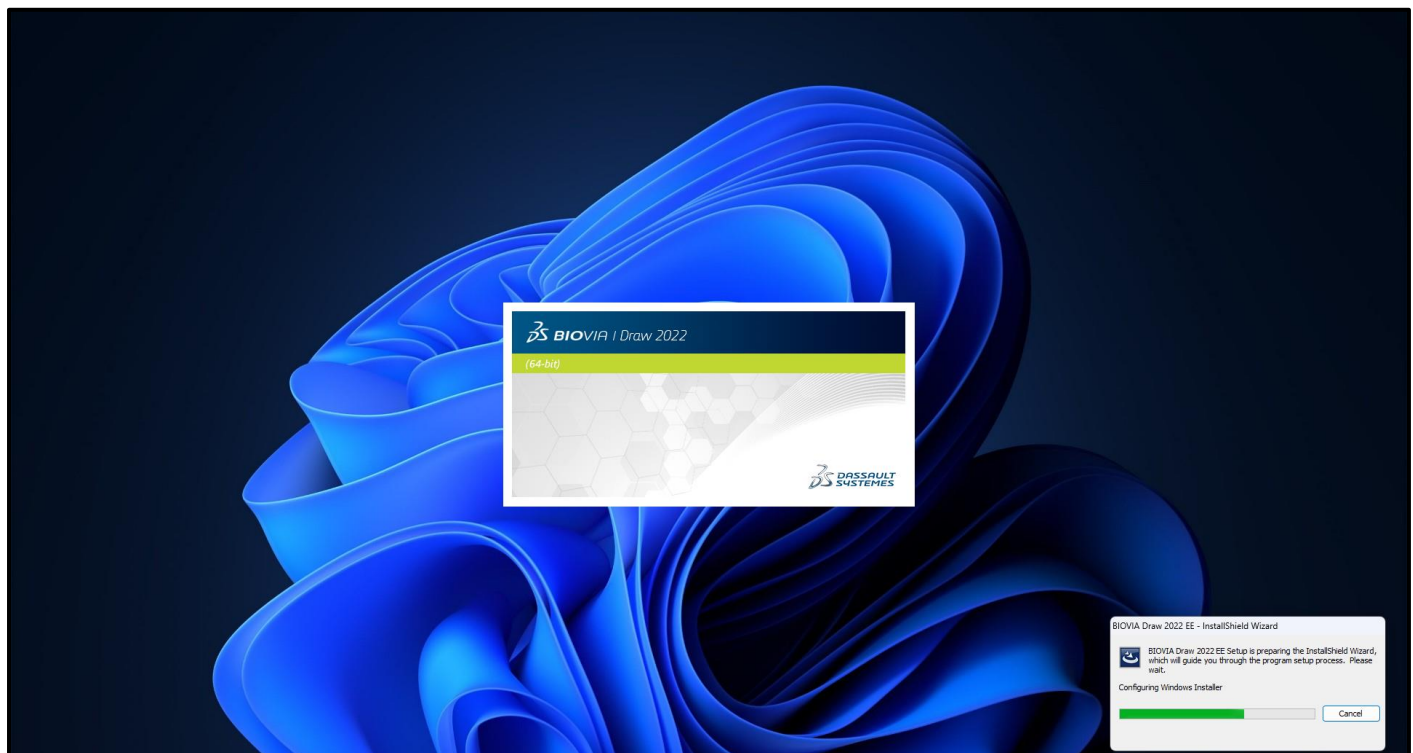
BIOVIA makes selective grants of licenses to customers who want to use BIOVIA Draw over a network or redistribute it to other users. To apply, download the network and/or distribution agreement(s) below and follow the instructions provided.

The following files are available to be downloaded:

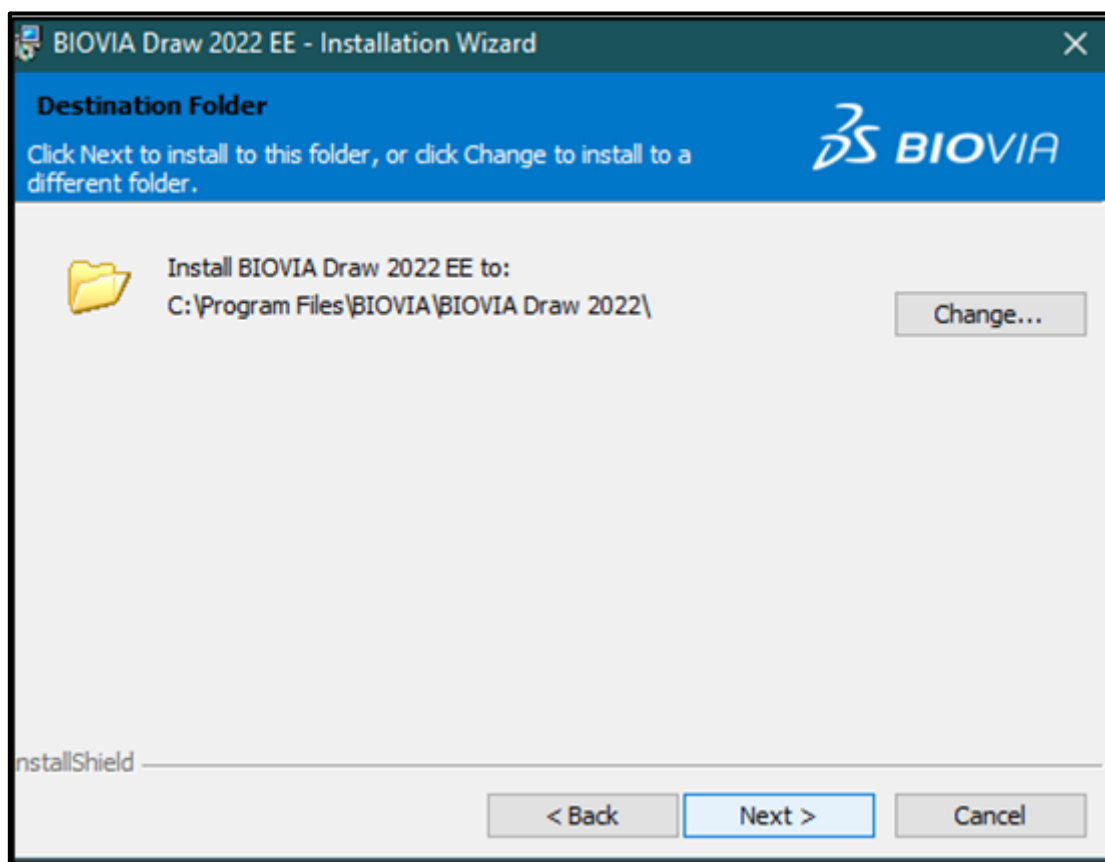
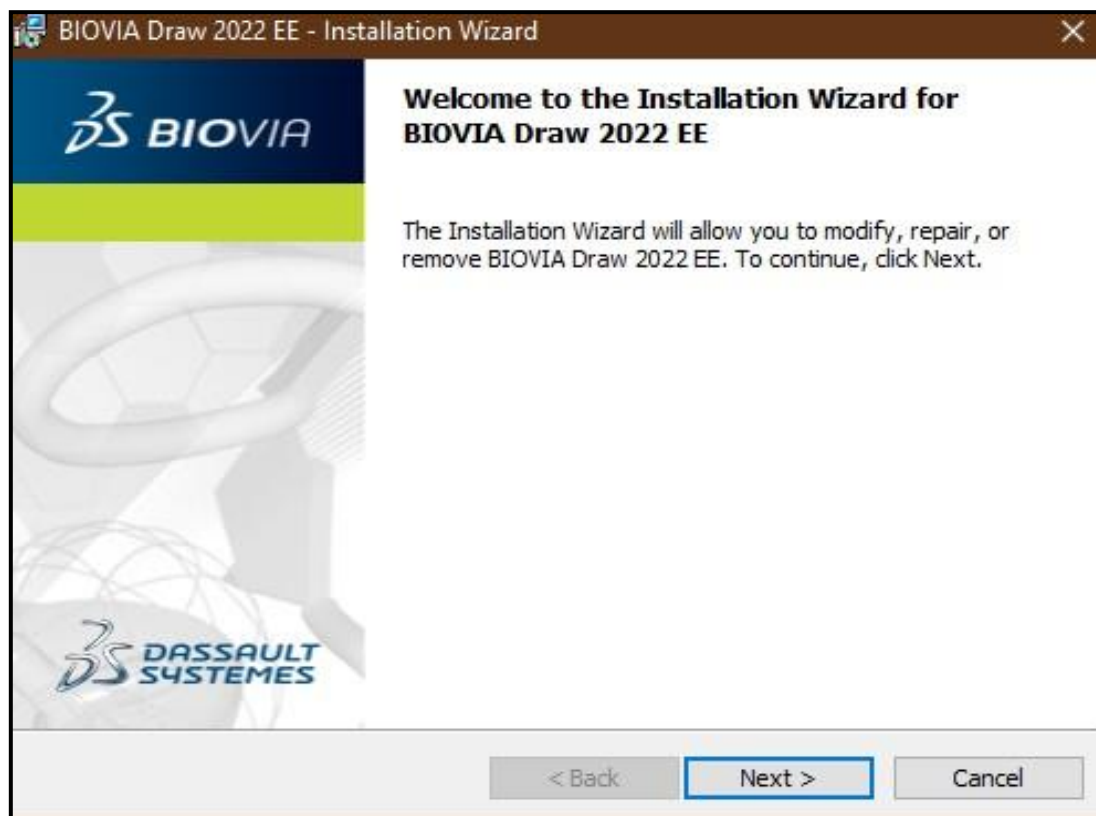
File Name	Action
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BIOVIA Draw Academic 2022 64-bit	Download
BIOVIA Draw Academic - End User License	Download

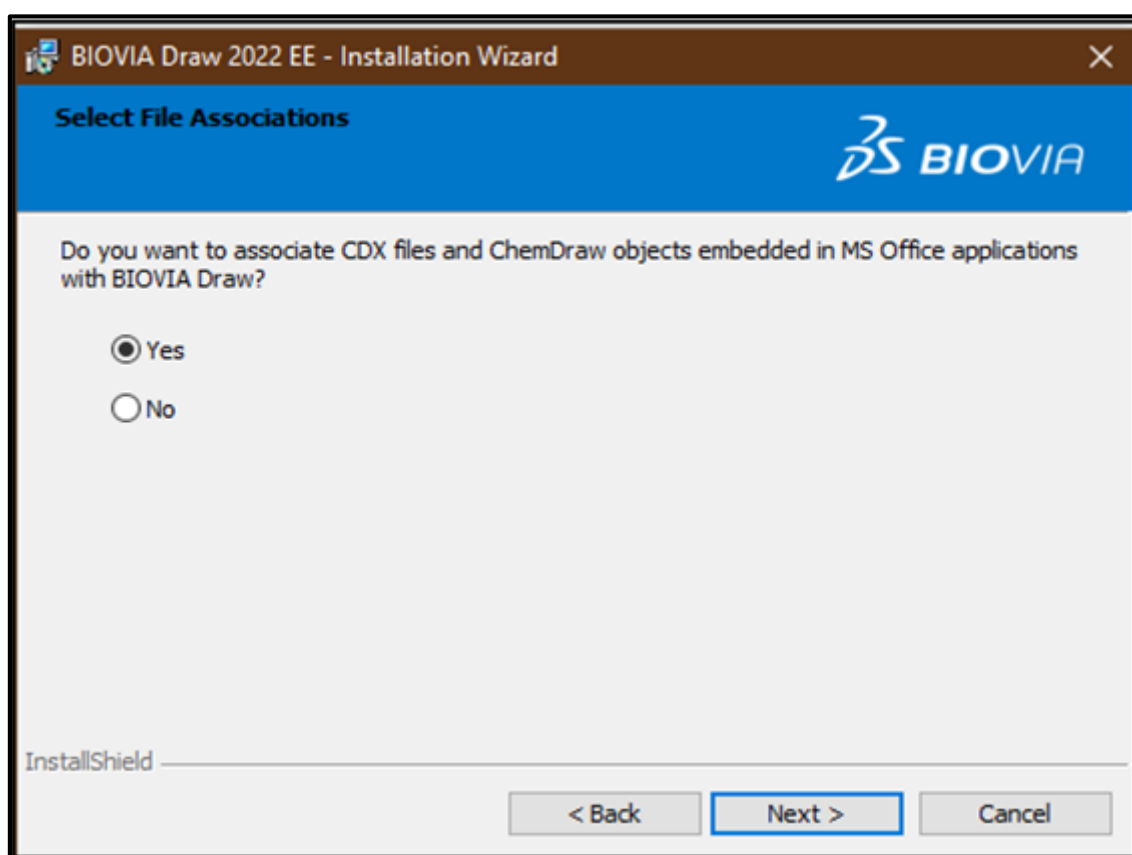
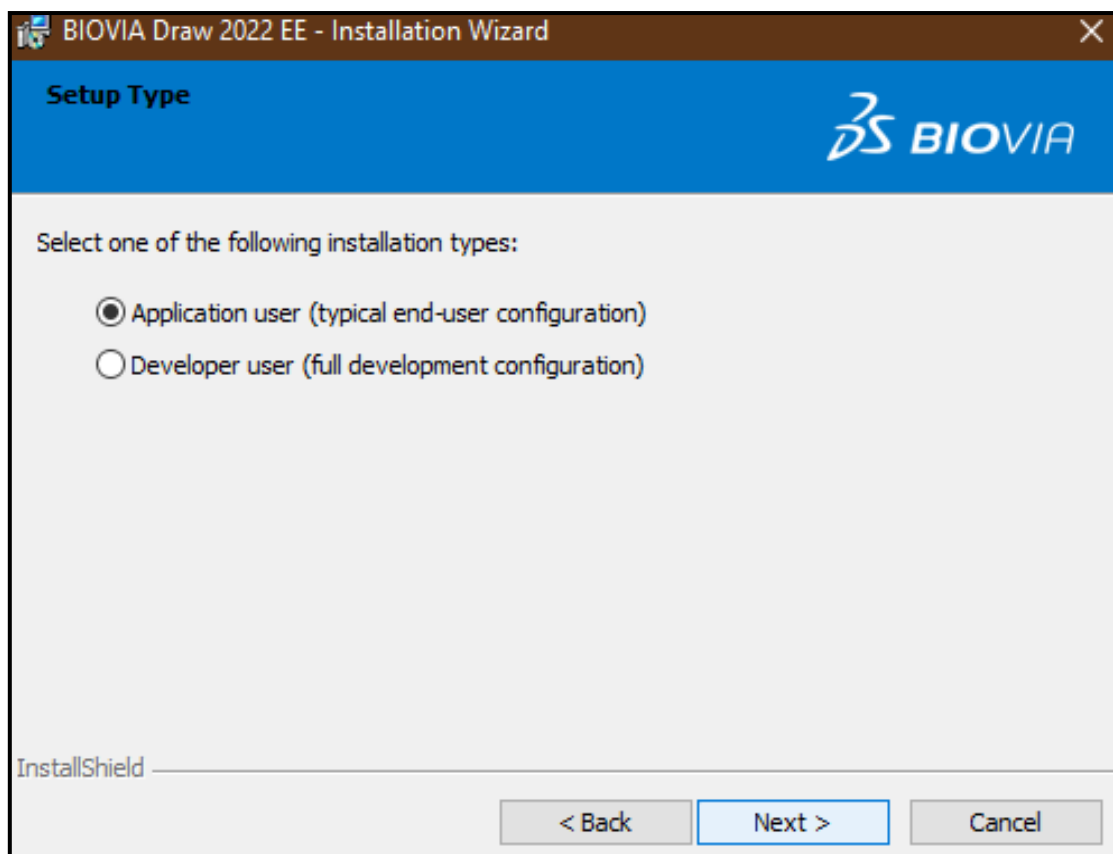
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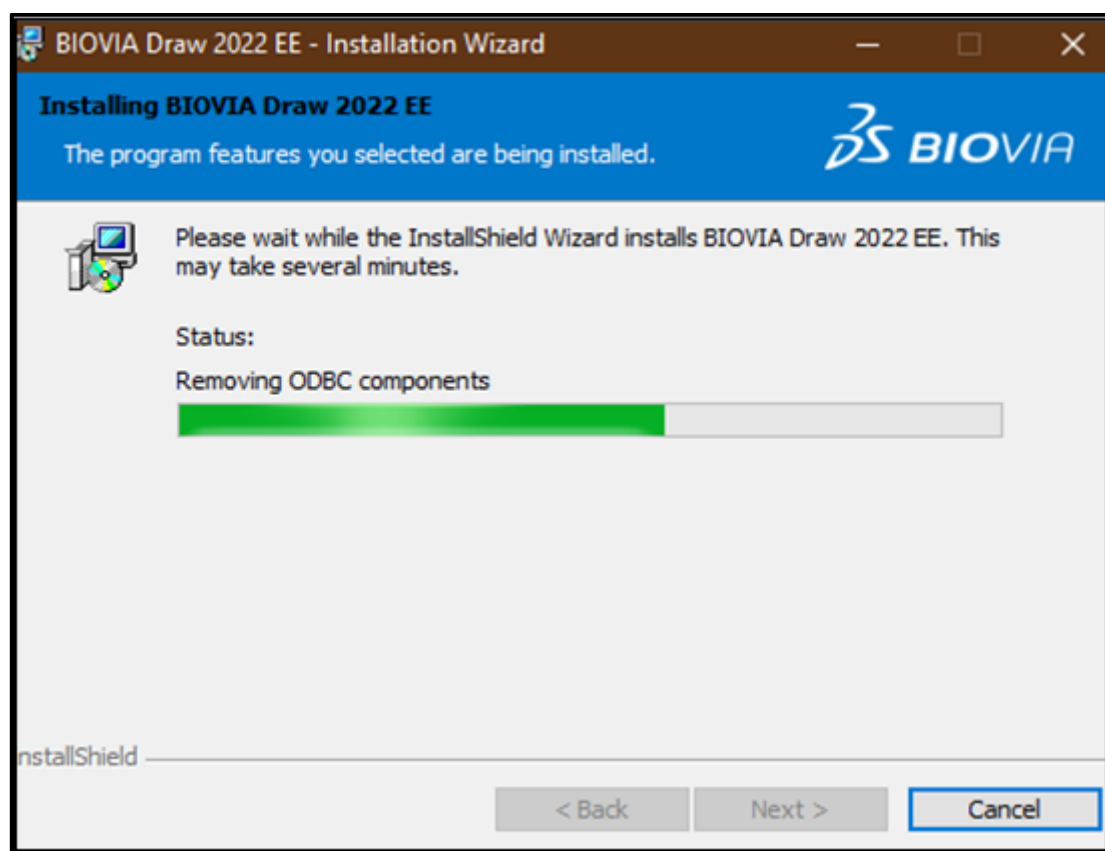
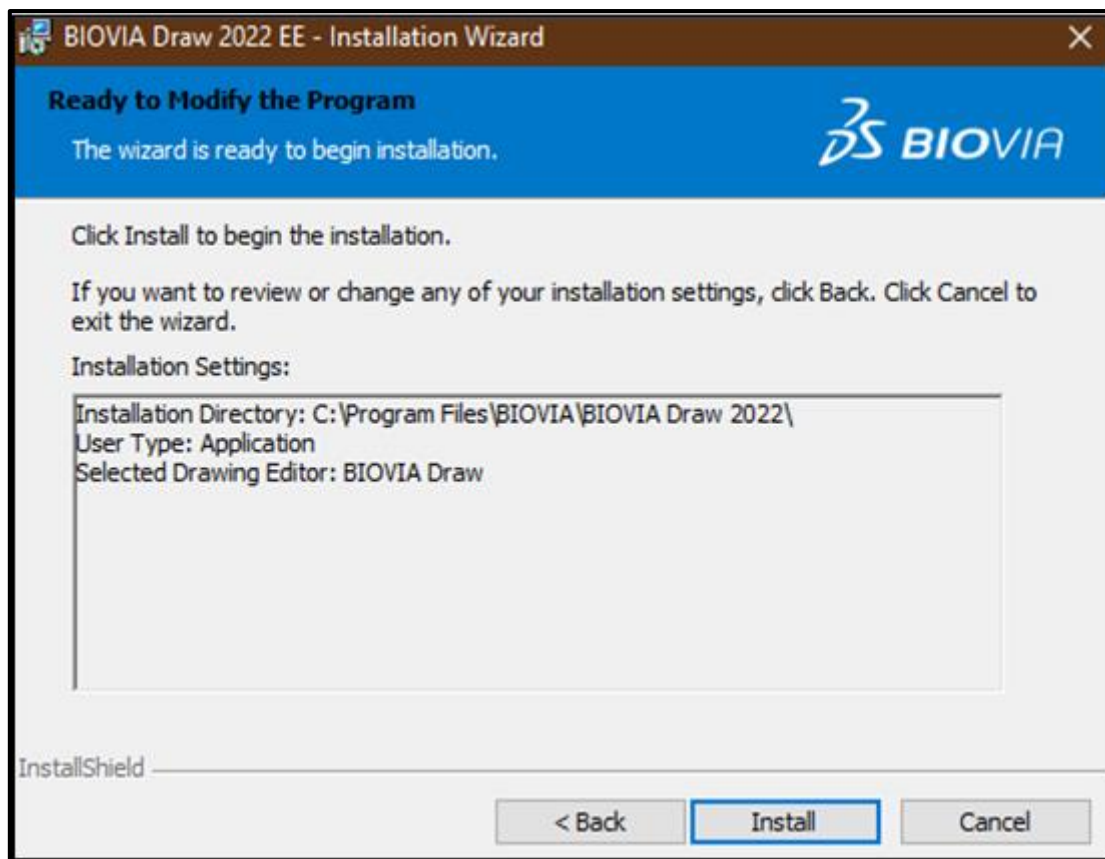
4. After Downloading, open it on your PC or Laptop



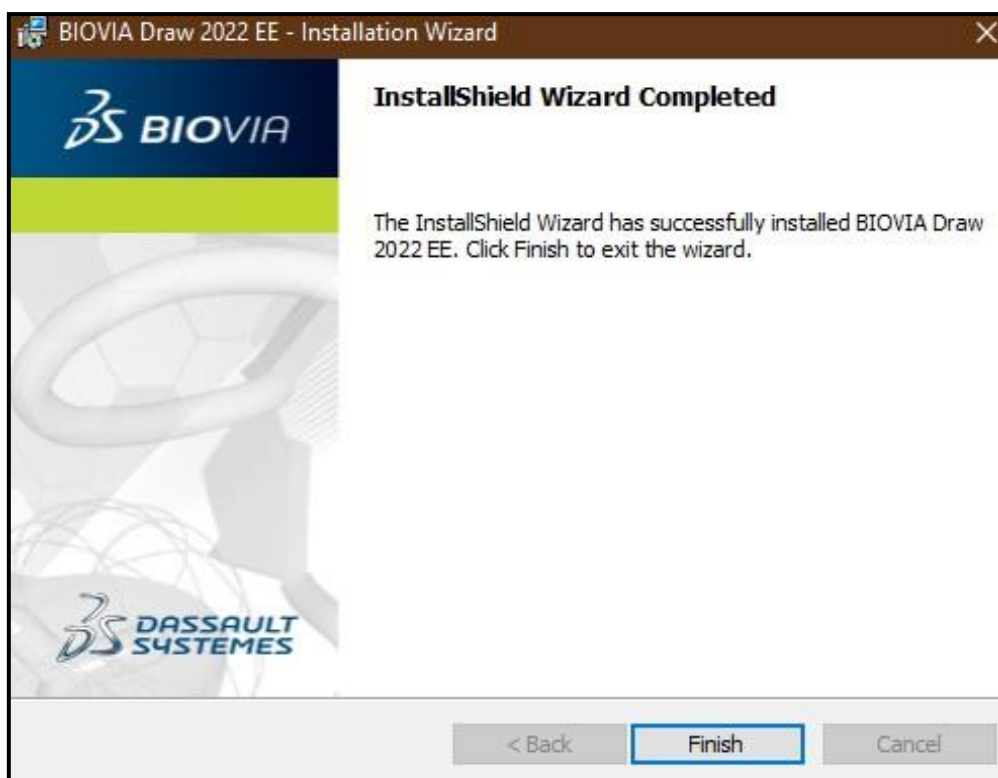
5. Check and select the optional components, and then click on 'Next'



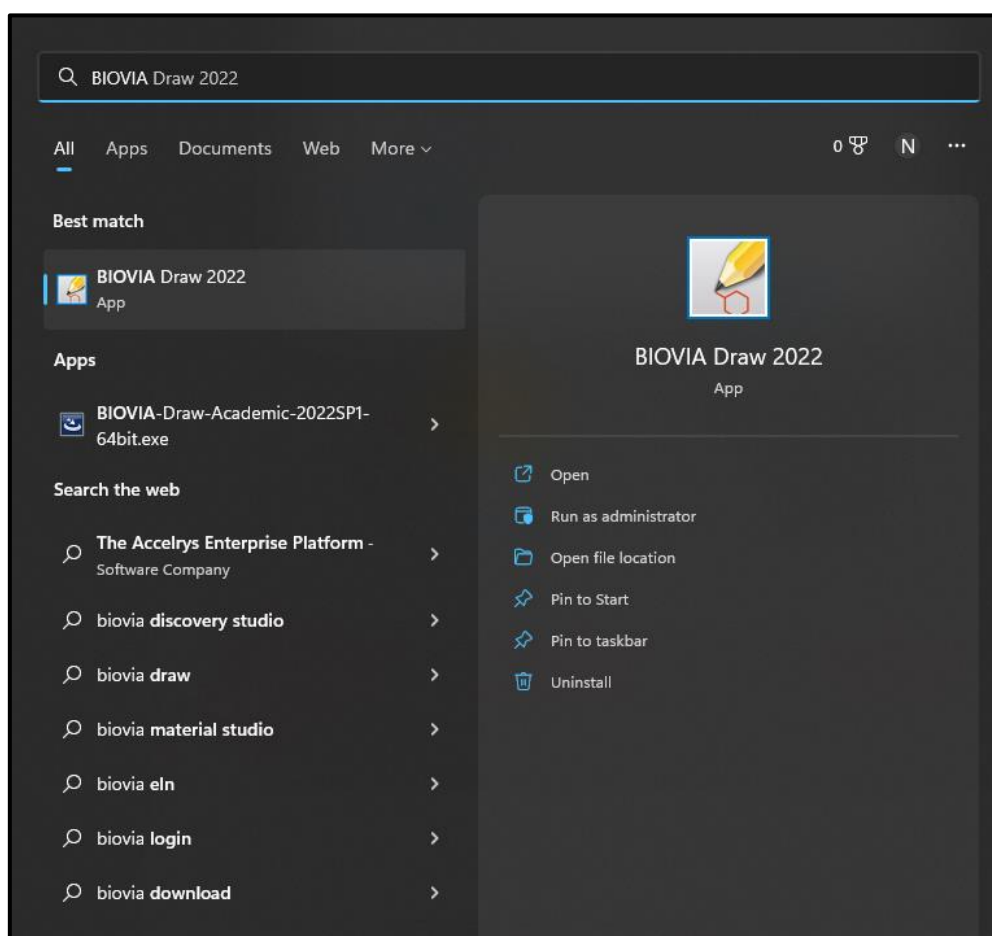




6. The Installation is completed. Click finish to exit the wizard



7. BIOVIA after installation



II. OPEN BABEL Tool

HISTORY:

Open Babel and JOELib were derived from the OELib cheminformatics library. In turn, OELib was based on ideas in the original chemistry program Babel and an unreleased object-oriented programming library called OBabel.

INTRODUCTION:

The development and use of the Open Babel project, a full-featured open chemical toolbox, designed to "speak" the many different representations of chemical data. It allows anyone to search, convert, analyze, or store data from molecular modeling, chemistry, solid-state materials, biochemistry, or related areas. It provides both ready-to-use programs as well as a complete, extensible programmer's toolkit for developing cheminformatics software. It can handle reading, writing, and interconverting over 110 chemical file formats, supports filtering and searching molecule files using Daylight SMARTS pattern matching and other methods, and provides extensible fingerprinting and molecular mechanics frameworks. We will discuss the frameworks for file format interconversion, fingerprinting, fast molecular searching, bond perception and atom typing.


Open Babel has its origin in a version of OELib released as open-source software by OpenEye Scientific under the GPL (GNU Public License). In 2001, OpenEye decided to rewrite OELib in-house as the proprietary OEChem library, so the existing code from OELib was spun out into the new Open Babel project. Since 2001, Open Babel has been developed and substantially extended as an international collaborative project using an open-source development model. It has over 160,000 downloads, over 400 citations, is used by over 40 software projects, and is freely available from the Open Babel website.

FEATURES:

- 1) File Format Support.
- 2) Fingerprints and Fast Searching.
- 3) Bond Perception and Atom Typing.
- 4) Canonical Representation of Molecules.
- 5) Coordinate Generation in 2D and 3D.

Installation Steps:

1. Open Homepage of Open Babel website and then click on download (URL: http://openbabel.org/wiki/Main_Page).



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[Capabilities](#)
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[Develop with Babel](#)
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
Open Babel: The Open Source Chemistry Toolbox

Open Babel is a chemical toolbox designed to speak the many languages of chemical data. It's an open, collaborative project allowing anyone to search, convert, analyze, or store data from molecular modeling, chemistry, solid-state materials, biochemistry, or related areas.

- **Ready-to-use programs, and complete programmer's toolkit**
- Read, write and convert over **110 chemical file formats**
- Filter and search molecular files using **SMARTS** and other methods
- Supports molecular modeling, cheminformatics, bioinformatics
- Organic chemistry, inorganic chemistry, solid-state materials, nuclear chemistry
- Downloaded over **325,000 times** and used by over **40 related projects**
- More about Open Babel
- Open Babel on SourceForge


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To support Open Babel, please cite [J. Cheminf. 2011, 3:33](#)

Navigation


- **Get Open Babel**
(downloads, installation, release notes)
- **Need Help?**
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- **Develop with Open Babel**
(how does it work, examples, developer documentation...)
- **Get Involved**
(tell others, report bugs, suggest features, citing Open Babel...)

News

- **2016-09-21** Open Babel 2.4.0 Released
- **2012-10-11** Open Babel 2.3.2 Released
- **2011-10-17** Open Babel 2.3.1 Released
- **2011-10-07** Open Babel Paper Published
- **2011-03-24** Pre-registration open for 2nd Open Babble
- **2010-10-26** Open Babel 2.3.0 Released
- **2010-07-06** Silicos NV contributes Spectrophore code
- **2009-07-31** Open Babel 2.2.3 Released
- **2008-07-10** Open Babel 2.2.2 Released
- **2009-02-03** Open Babel 2.2.1 Released
- **2008-07-04** Open Babel 2.2.0 Released
- **2008-03-20** Paper on Pybel published
- **2008-02-26** New OpenBabel.org Website
- **2007-07-07** Open Babel 2.1.1 Released
- **2007-04-07** Open Babel 2.1.0 Released

2. Open Babel is available on Windows, Linux and Mac. For Windows install the 64bit version.



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Category
Discussion

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Category:Installation

(Redirected from Get Open Babel)

Open Babel is available for Windows, Linux and MacOSX.

Windows

- **Open Babel GUI**
Provides a graphical user interface for Open Babel, as well as a command-line interface.
Get the [latest installer](#) for 64-bit (recommended) or 32-bit (has "-x86" in name).
[Documentation](#)
- **Python module** (requires OpenBabelGUI above)
Provides access to the Open Babel libraries from Python.
- **Java library** (requires OpenBabelGUI above)
Provides access to the Open Babel libraries from Java.
- **OBDotNet assembly** (requires OpenBabelGUI above)
Provides access to the Open Babel libraries from .NET languages.

The following options are only recommended for experienced developers.

- **Compile from source**
Download the source for the latest release
or Get latest development code
[Compile instructions](#)
Open Babel can be compiled using any of MSVC++, Cygwin or MinGW

Linux

- **Compile from source**
Compile Open Babel:
[Download the latest release](#)
or Get latest development code (today)
[How to compile](#)
[How to use obabel](#)
[How to develop with Open Babel](#)

Scripting language modules:
[Perl](#), [Python](#), [Ruby](#), [Java](#), [Mono](#)
or

- **Install a binary package**
Several Linux distributions provide binary packages. For scripting languages, the package may be named like 'openbabel-perl' or 'python-openbabel'
Alternatively, with Conda: `conda install openbabel -c conda-forge`
Also available as a snap package: `snap install openbabel`

MacOSX

There are several ways to install Open Babel on MacOSX.

- With Conda, `conda install -c conda-forge openbabel`
- With HomeBrew, `brew install open-babel`
- **Compile the source code**
Compile Open Babel using clang or gcc

Once installed, you may wish to try [iBabel](#), a graphical interface to Open Babel.

This category has the following 3 subcategories, out of 3 total.

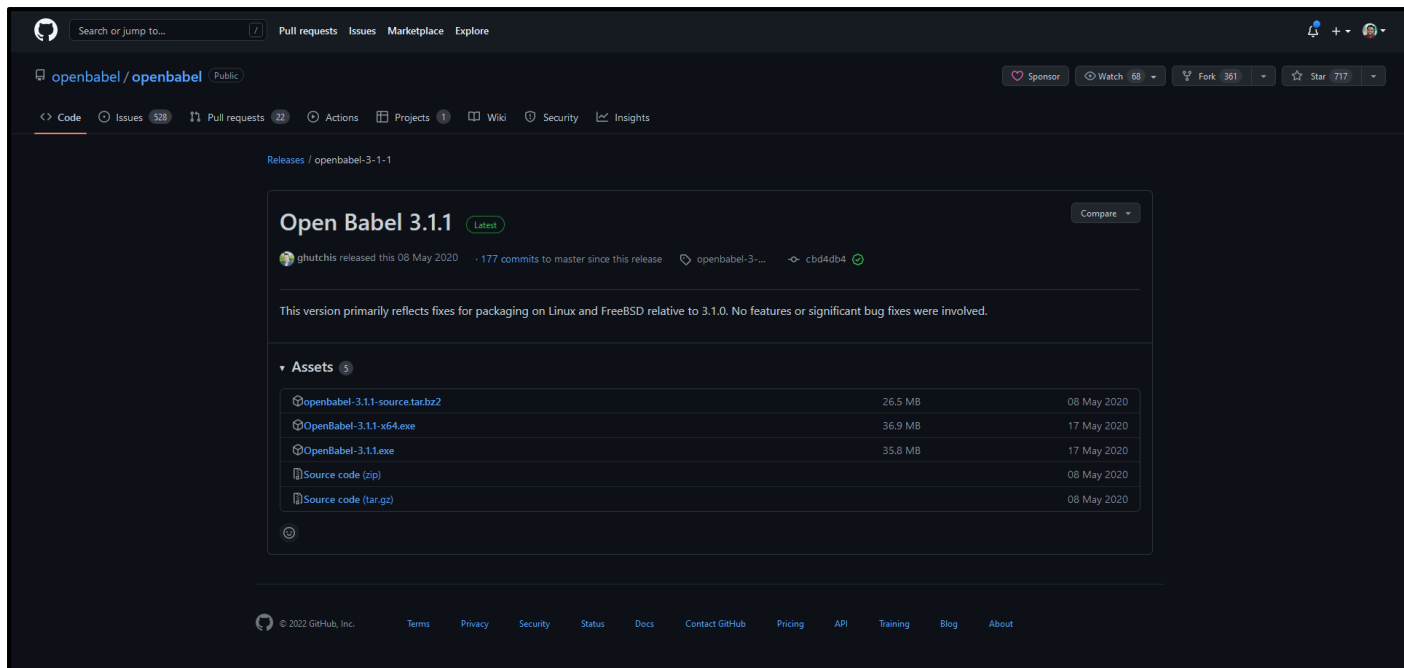
M

- [Macintosh](#)

R

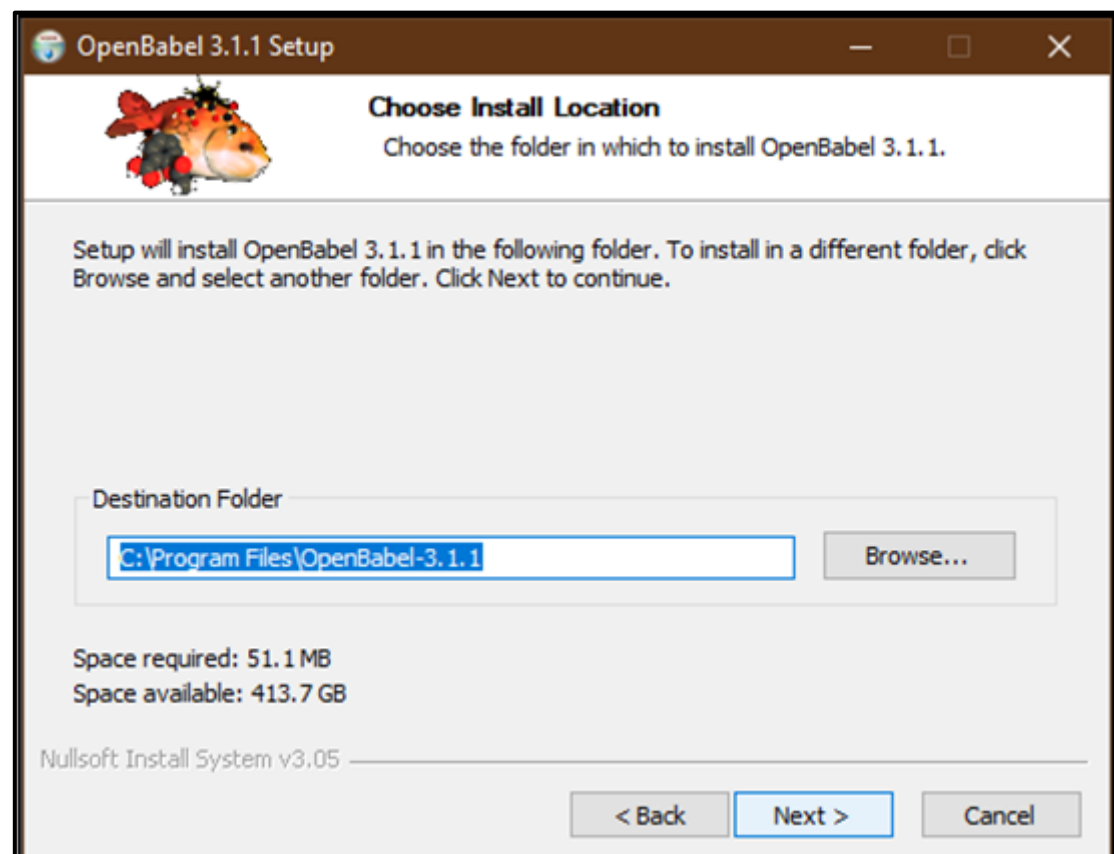
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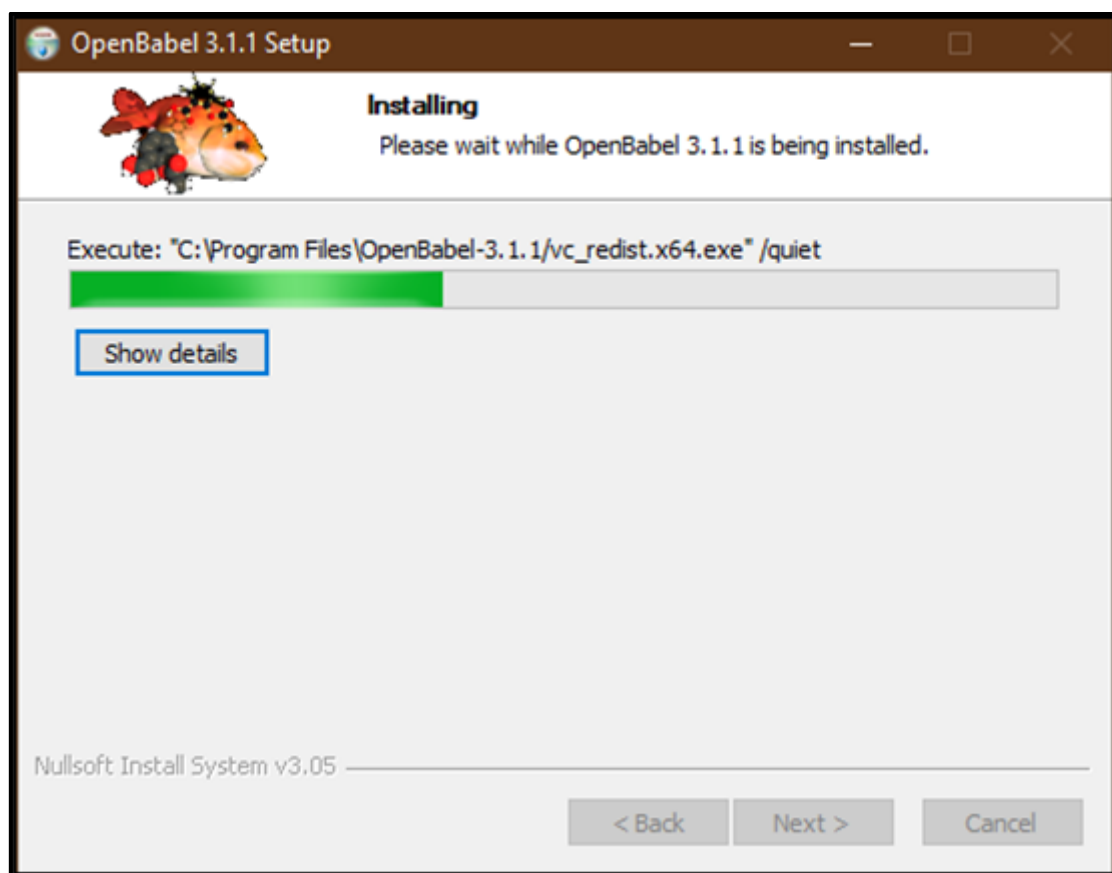
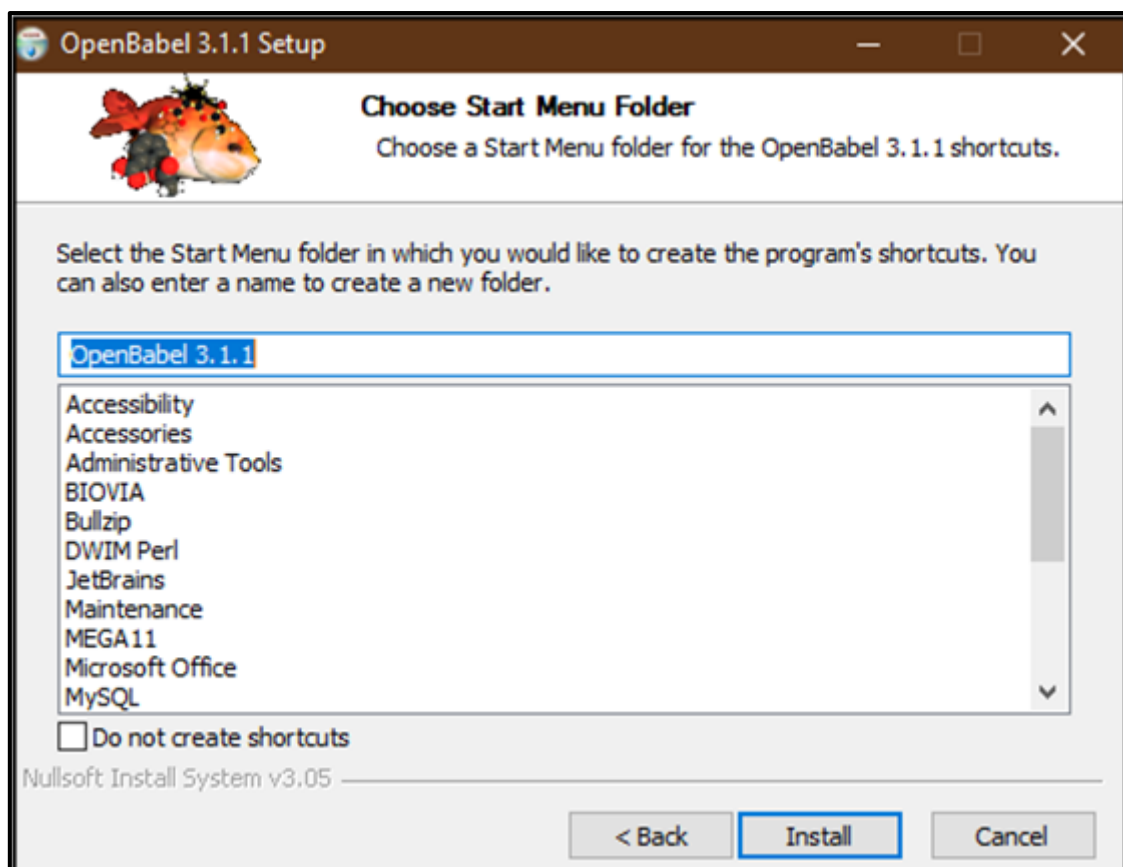
3. Click on OpenBabel-3.1.1.-64.exe



4. GO through the installation procedure

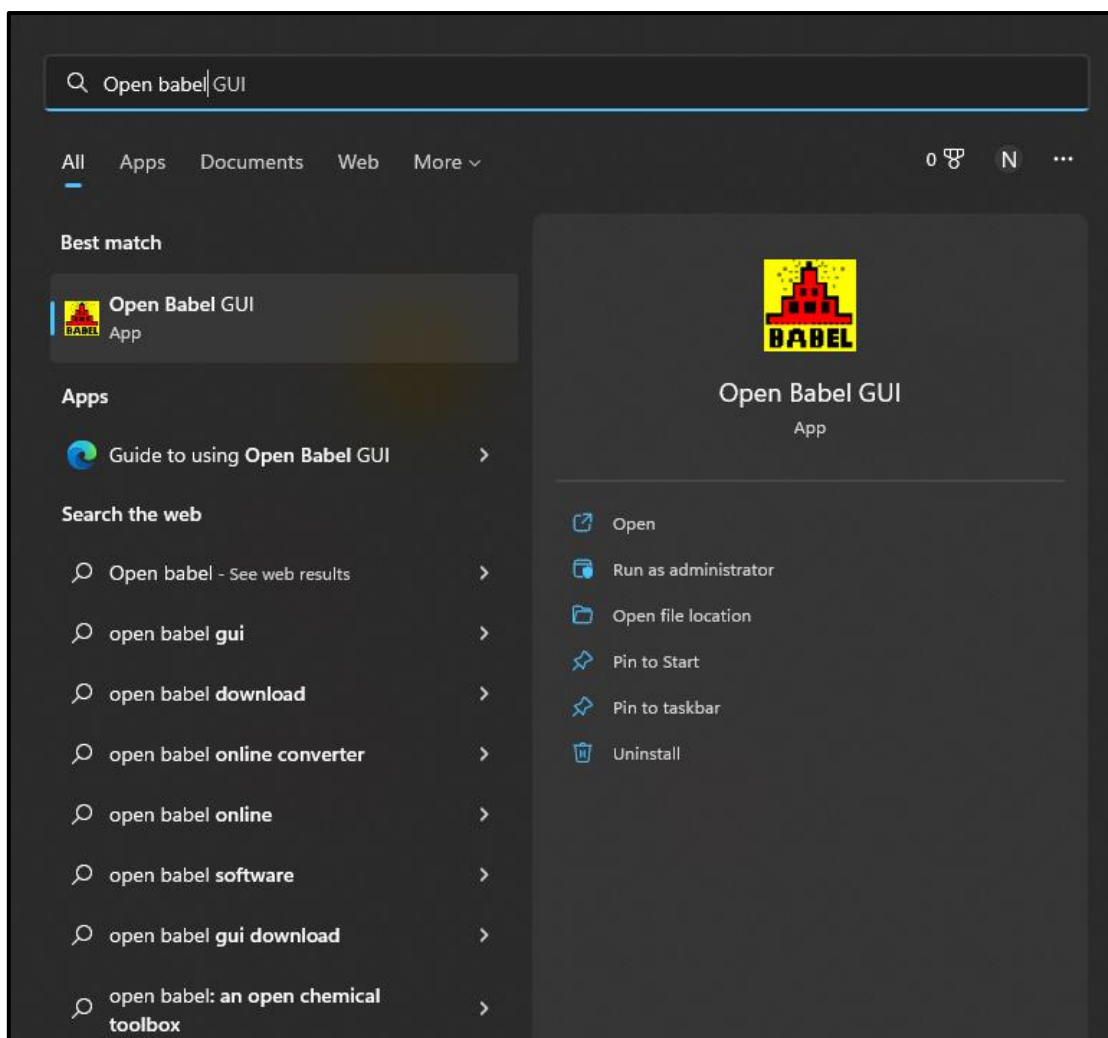








5. Icon of Open Babel GUI after installation



References:

1. Biovia draw. (n.d.). BIOVIA Draw. Retrieved September 7, 2022, from https://lib.cnu.edu.tw/3_2_resource/doc/biovia-draw.pdf
2. Price Waterhouse Coopers (PWC) From Vision to Decision. Pharma 2020 Report. PWC; London, UK: 2012. [(accessed on 16 November 2015)].
3. Weininger, D. (1988). SMILES, a chemical language and information system. 1. Introduction to methodology and encoding rules. J Chem Inf Comput Sci. 28, 31-36.

WEBLEM: 2A

BIOVIA DRAW SOFTWARE

(URL:<https://www.3ds.com/products-services/biovia/products/scientific-informatics/biovia-draw/>).

Aim:

To perform drawing, editing & manipulation of 2D & 3D structures for query “Penicillin” using BIOVIA DRAW software.

Introduction:

Penicillin V is a member of the penicillin family exhibiting broad-spectrum antibiotic property. Penicillin V binds to penicillin binding proteins (PBP), the enzymes that catalyze the synthesis of peptidoglycan, which is a critical component of the bacterial cell wall. This leads to the interruption of cell wall synthesis, consequently leading to bacterial cell growth inhibition and cell lysis.

Phenoxymethylpenicillin is a narrow spectrum antibiotic also commonly referred to as Penicillin V or Penicillin VK. It is a phenoxymethyl analog of Penicillin G, or [benzylpenicillin]. An orally active naturally penicillin, phenoxymethylpenicillin is used to treat mild to moderate infections in the respiratory tract, skin, and soft tissues caused by penicillin G--sensitive microorganisms. Phenoxymethylpenicillin has also be used in some cases as prophylaxis against susceptible organisms. While there have been no controlled clinical efficacy studies that were conducted, phenoxymethylpenicillin has been suggested by the American Heart Association and the American Dental Association for use as an oral regimen for prophylaxis against bacterial endocarditis in patients with congenital heart disease or rheumatic or other acquired valvular heart disease when they undergo dental procedures and surgical procedures of the upper respiratory tract, except for those who are at an elevated risk for endocarditis.

With the use of BIOVIA Draw, scientists can quickly and easily sketch and edit complex compounds, chemical reactions, and biological sequences, easing group searches, viewing, sharing, and archiving of scientific data. With the ability to save and retrieve peptides, oligonucleotides, and oligosaccharides, among other complicated biological things, BIOVIA Draw provides scientists with specialised management skills. A biological sequence editor that enables the definition of unique residues and linkers, Markush structure tools, and haptic and hydrogen bond tools are just a few of the features available to scientists. Developers can build unique add-ins and drop them in as tools, buttons, or menu items using BIOVIA Draw's published API. Tools for enumeration, bioavailability, isotopomer distribution, stoichiometry calculations, and many other tasks are available as add-ins.

Methodology:

1. Open BIOVIA Draw software.
2. Draw the Chemical Structure using various tools.
3. In the chemistry option use Clean the Structure.
4. Apply different filter options such as Calculator, Atom Number, Generate Text from Structure (IUPAC, SMILE), Show sequence view, etc.
5. Observe and save the results.

Observations:

Shortcut

Header

Template

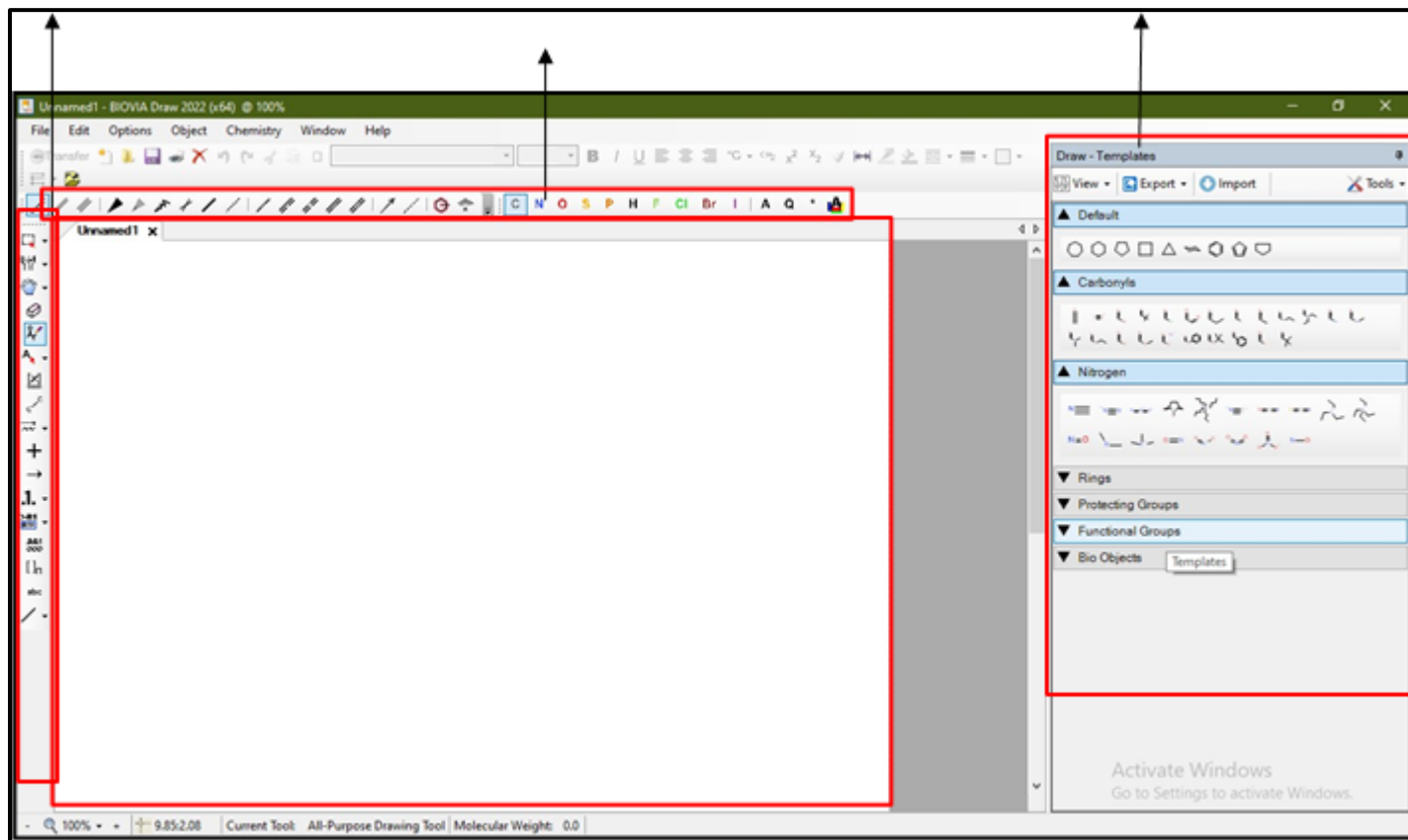


Fig 1. Homepage of BIOVIA DRAW SOFTWARE

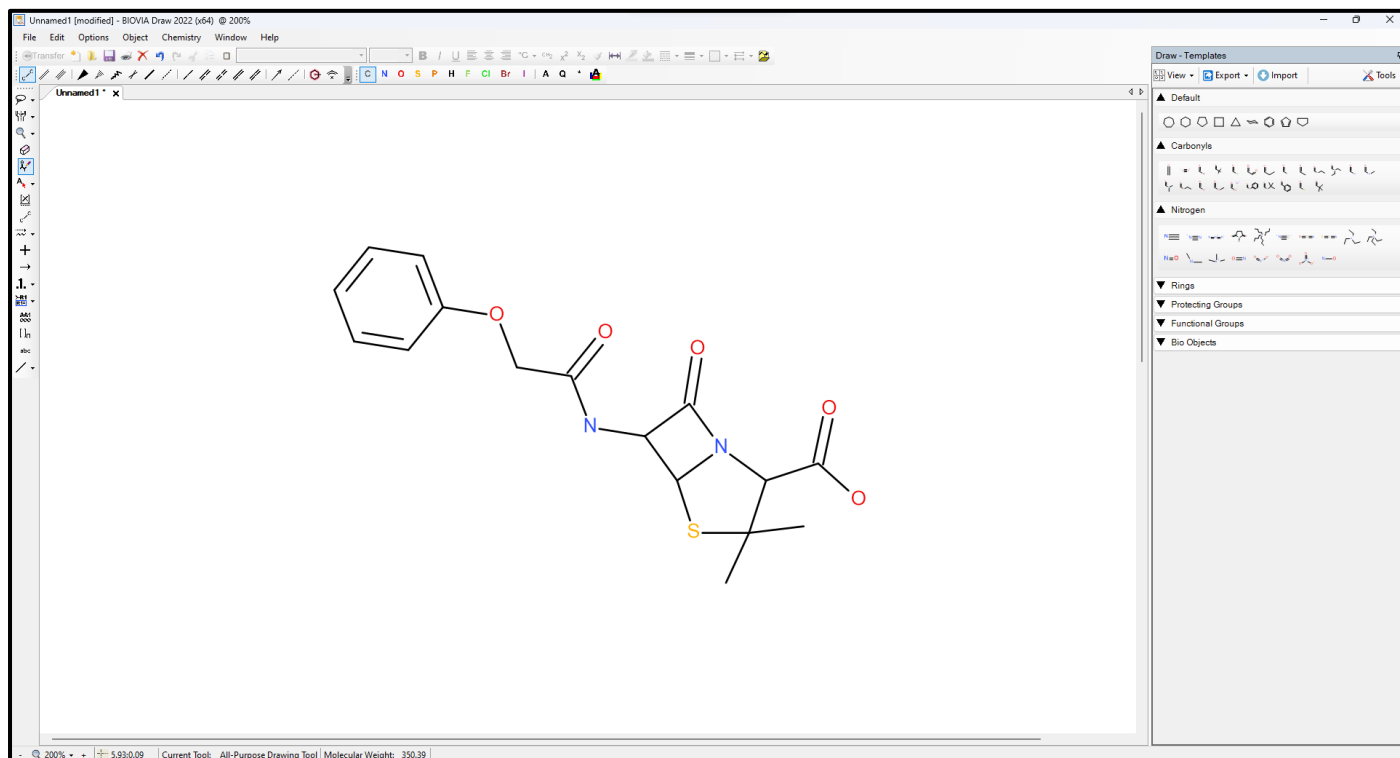


Fig 2. Structure of Penicillin chemical compound

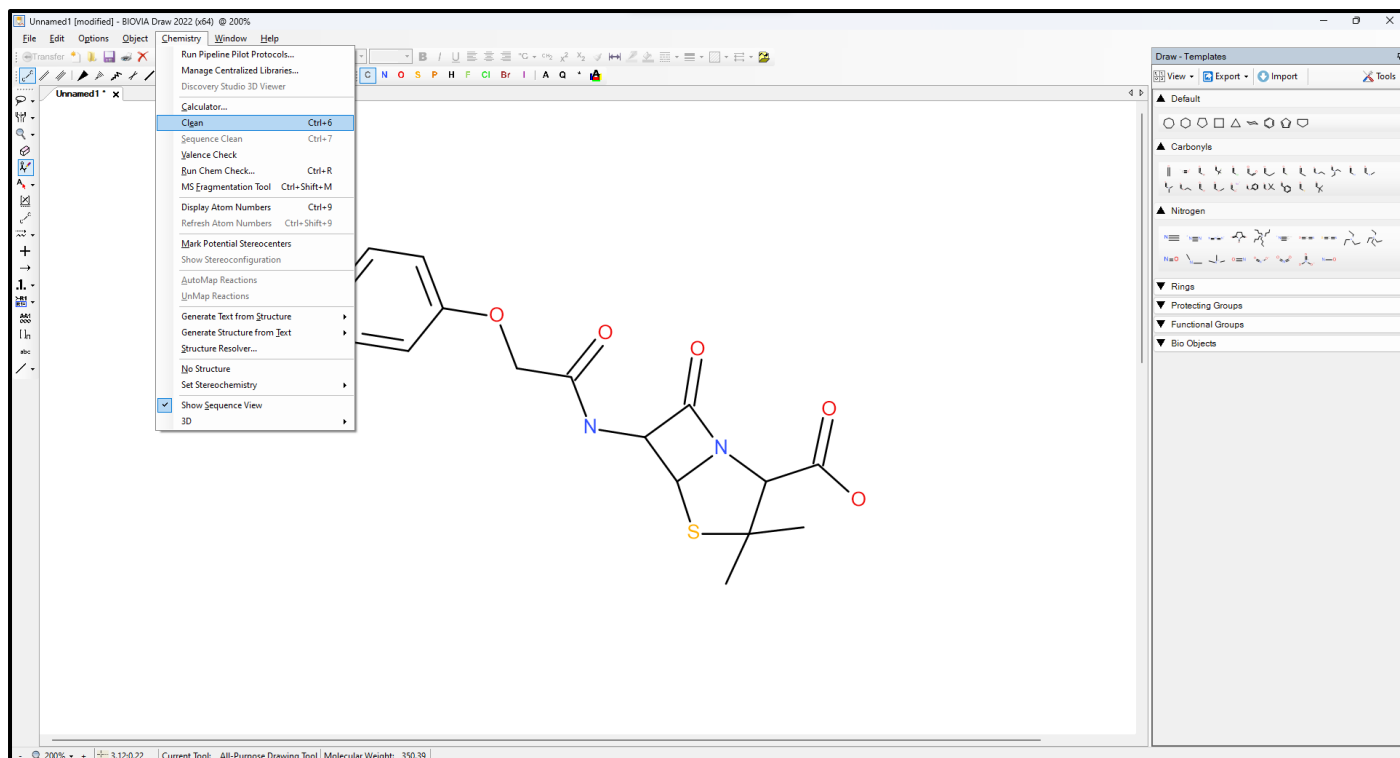


Fig 3. Option applied “Clean structure”

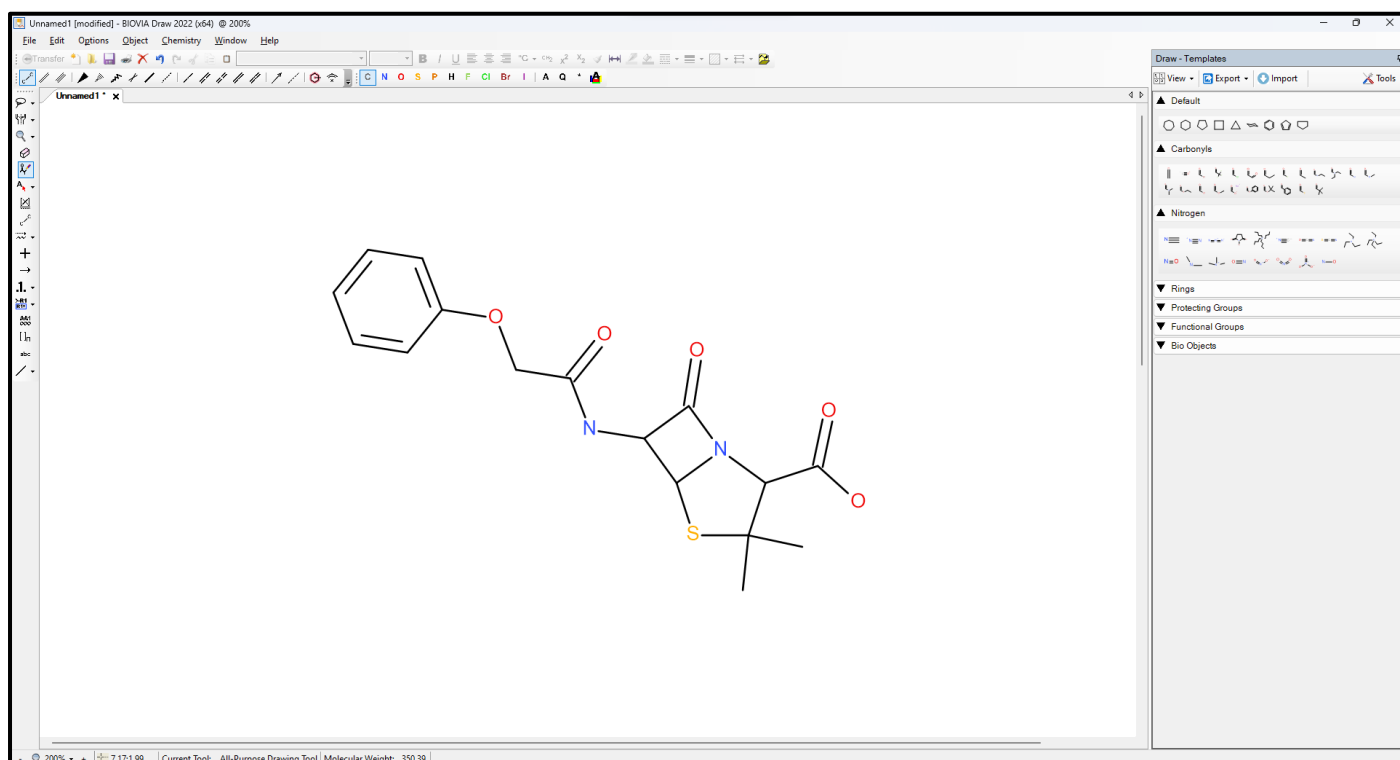


Fig 4. Clean structure of Penicillin

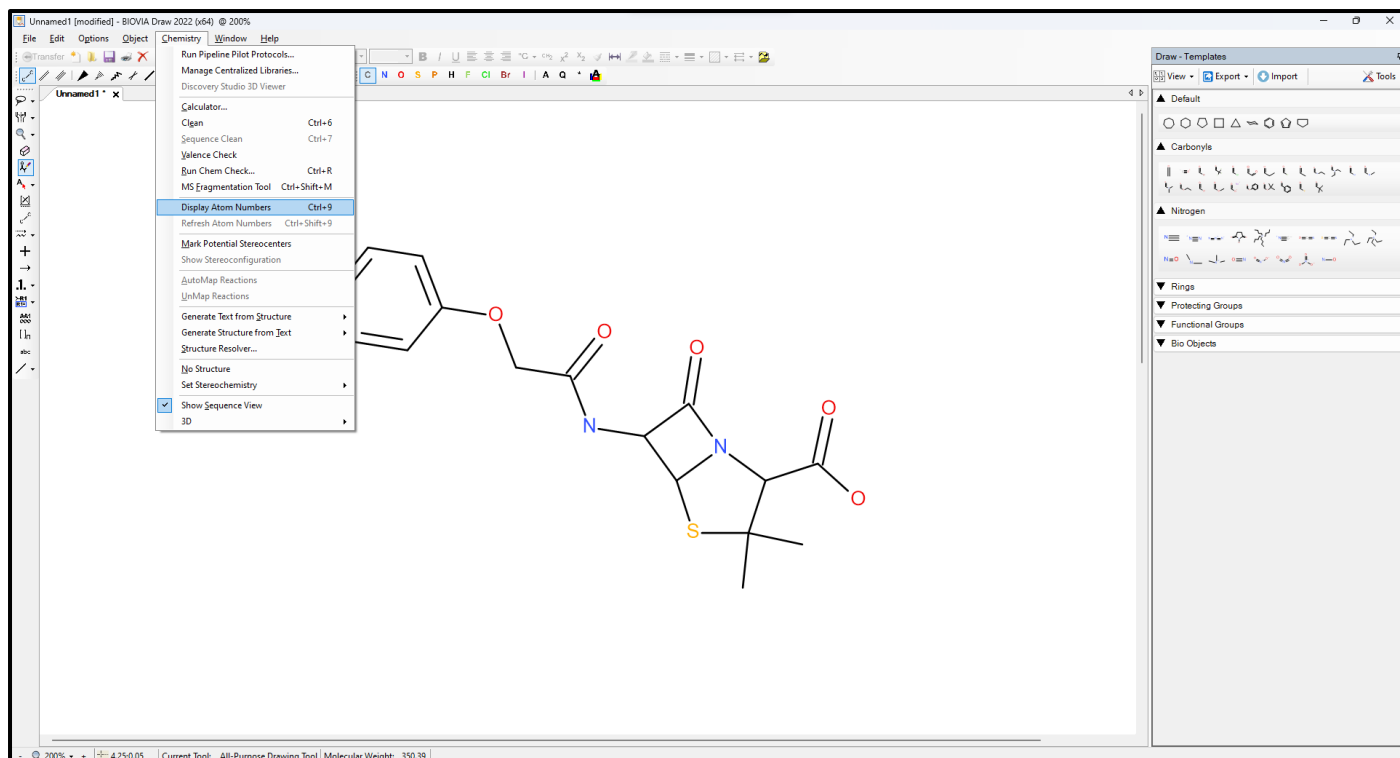


Fig 5. Option applied “Display Atom Numbers”

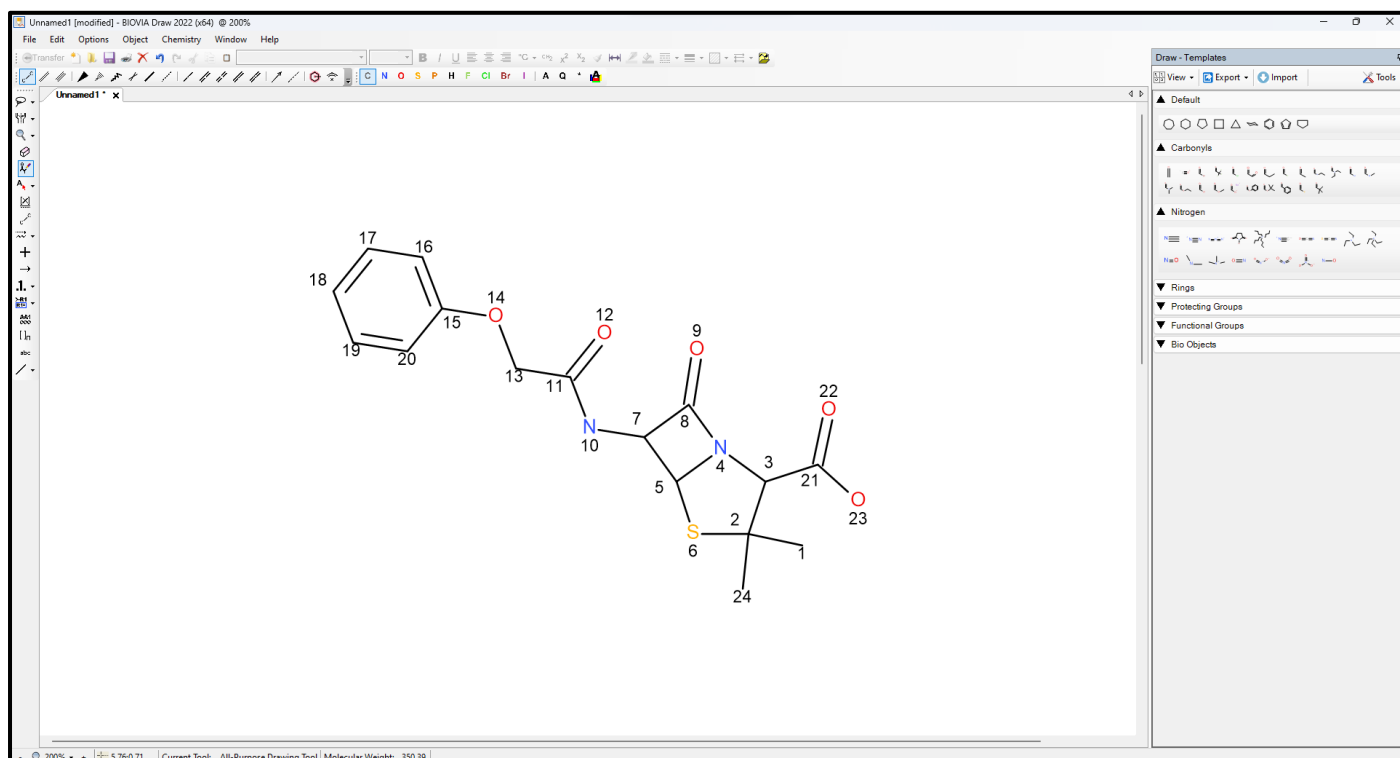


Fig 6. Atom numbering of Penicillin structure

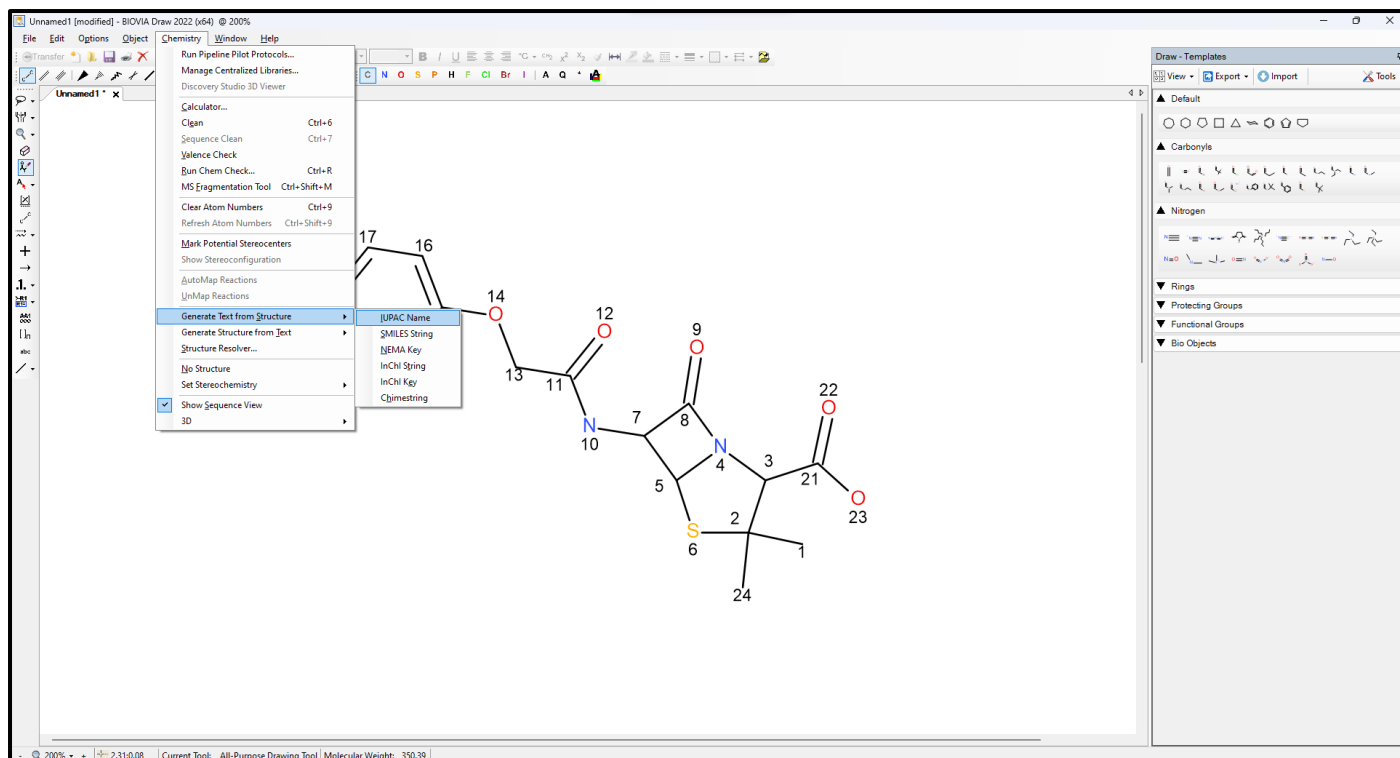


Fig 7. Option applied: To generate IUPAC name

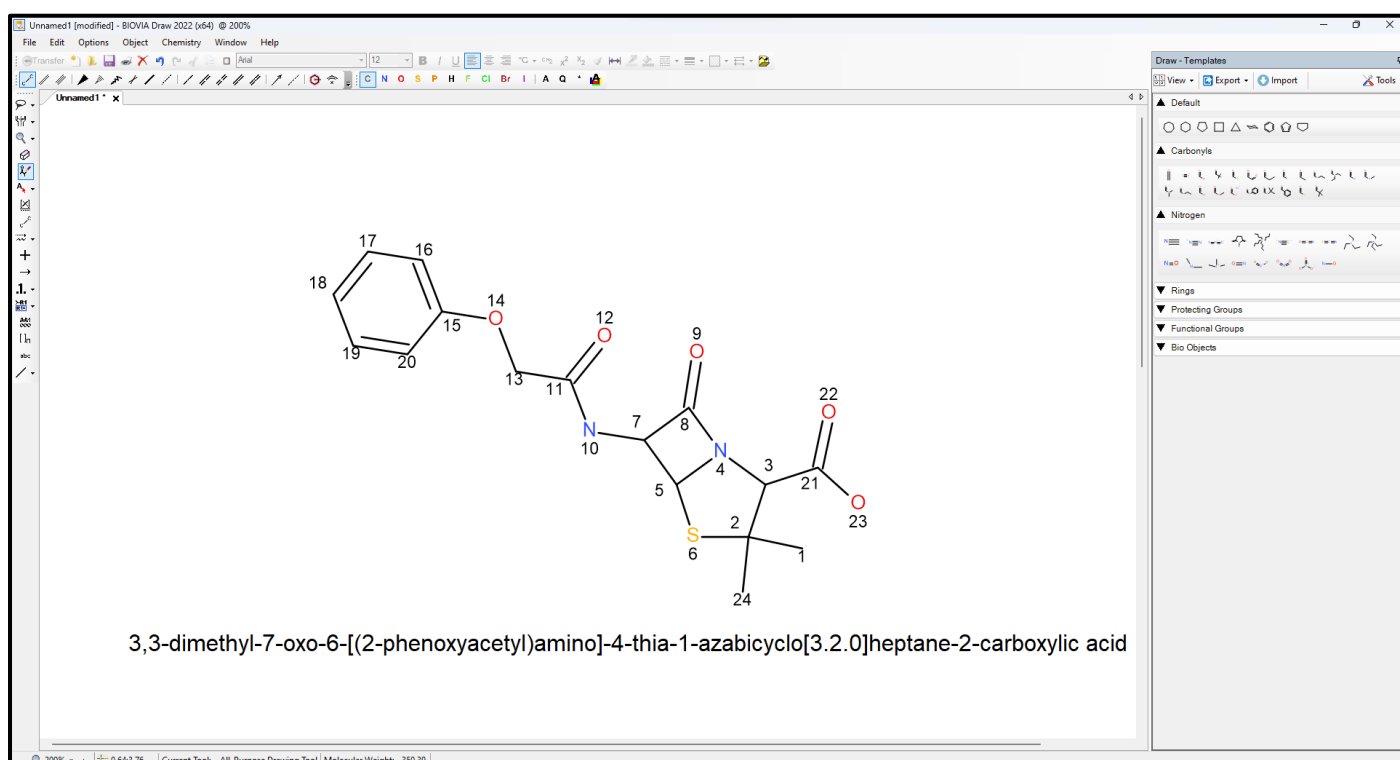


Fig 8. IUPAC name of Penicillin

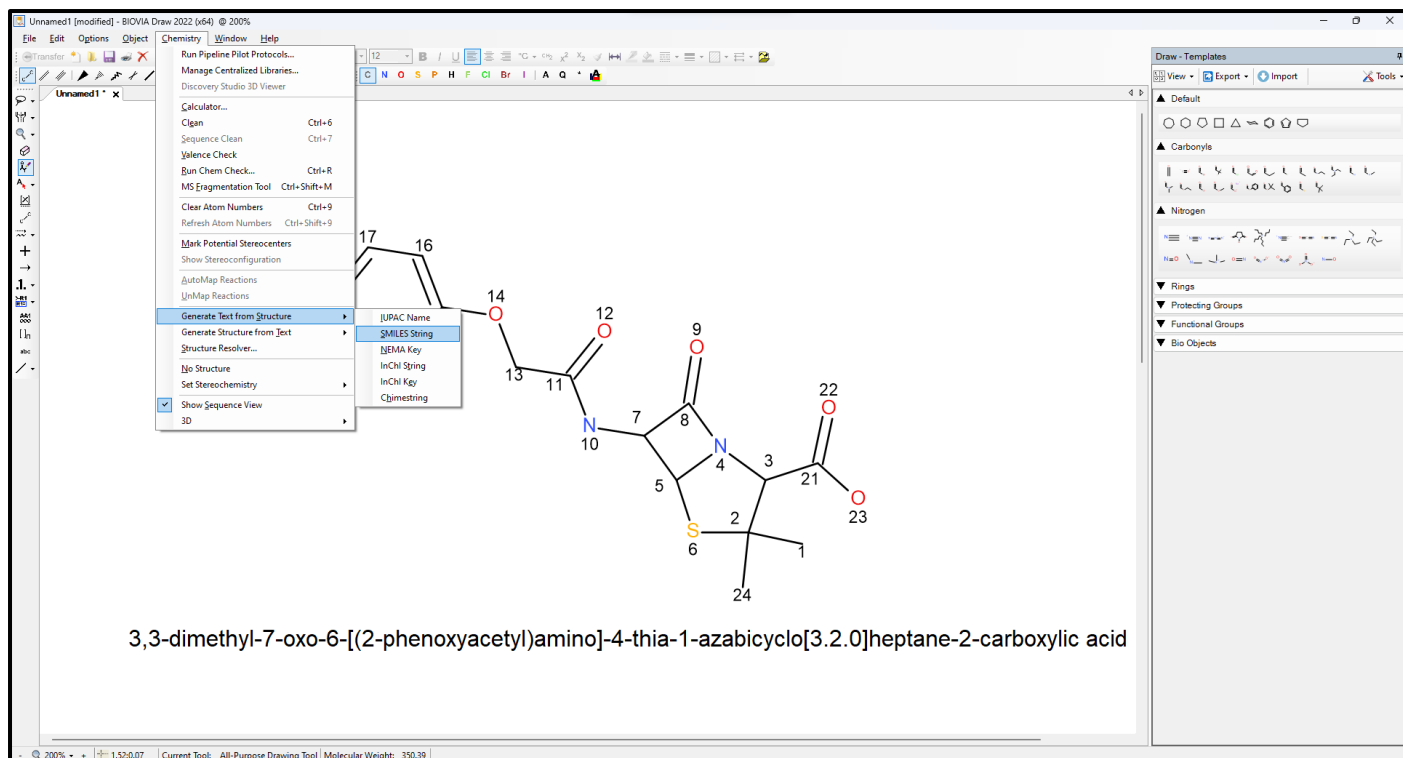


Fig 9. Option applied: To generate SMILES string

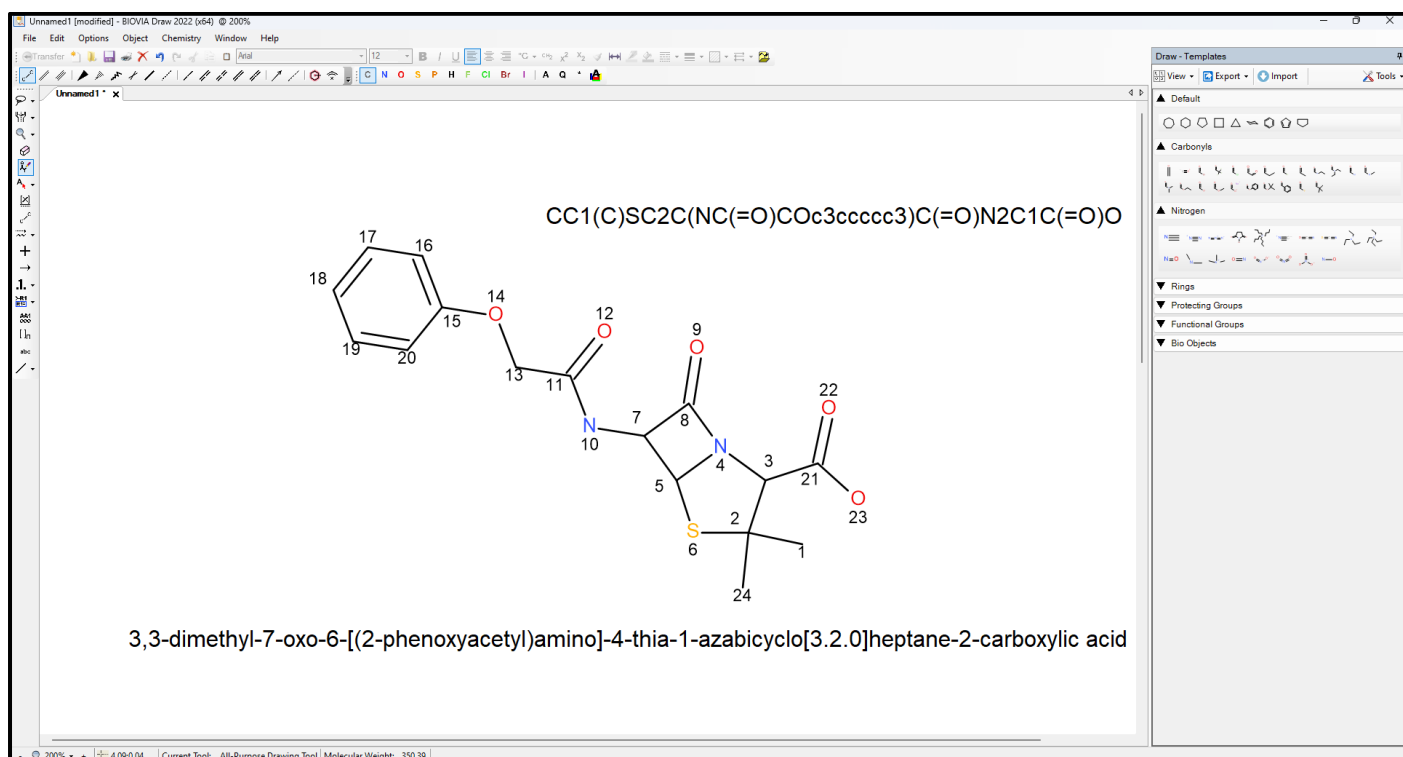


Fig 10. SMILES string of Penicillin

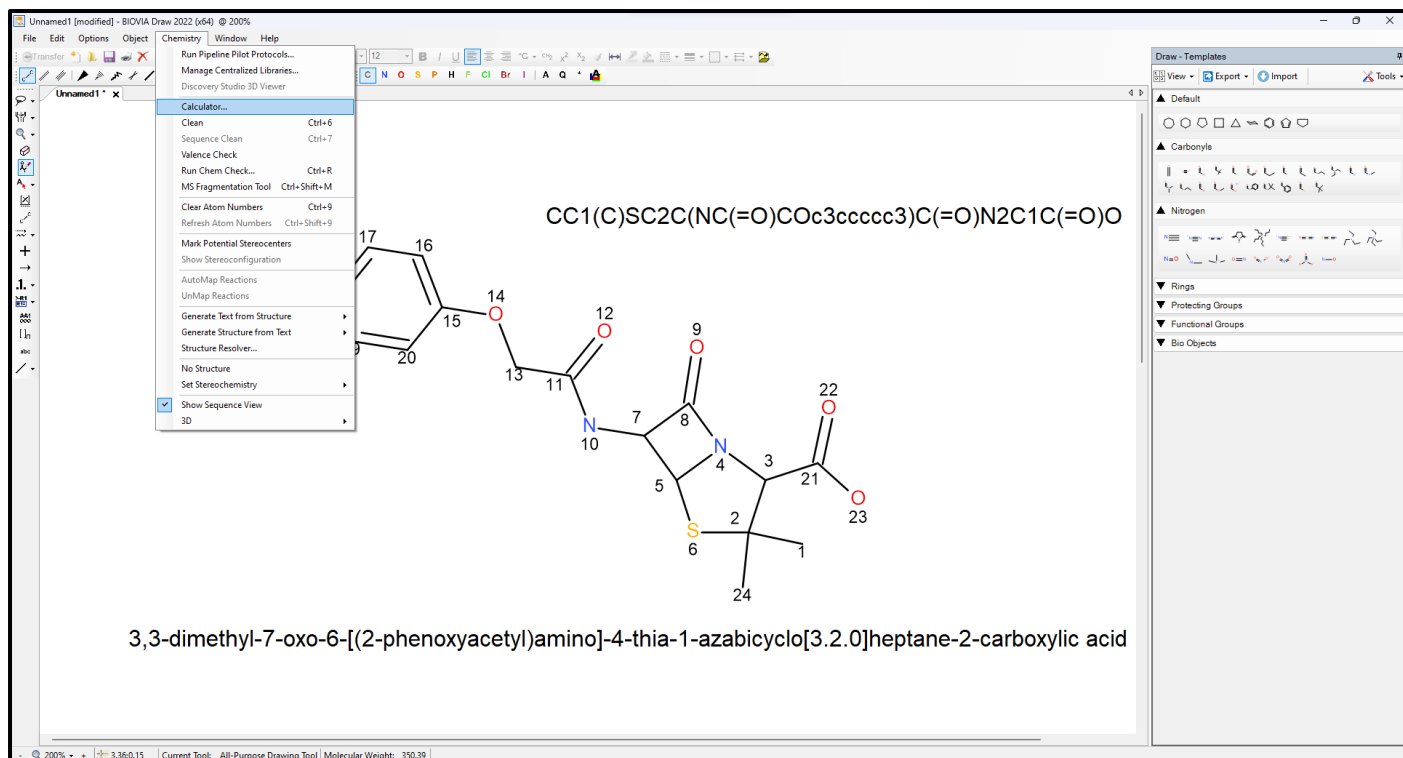


Fig 11. Option applied: To calculate Physicochemical properties of Penicillin

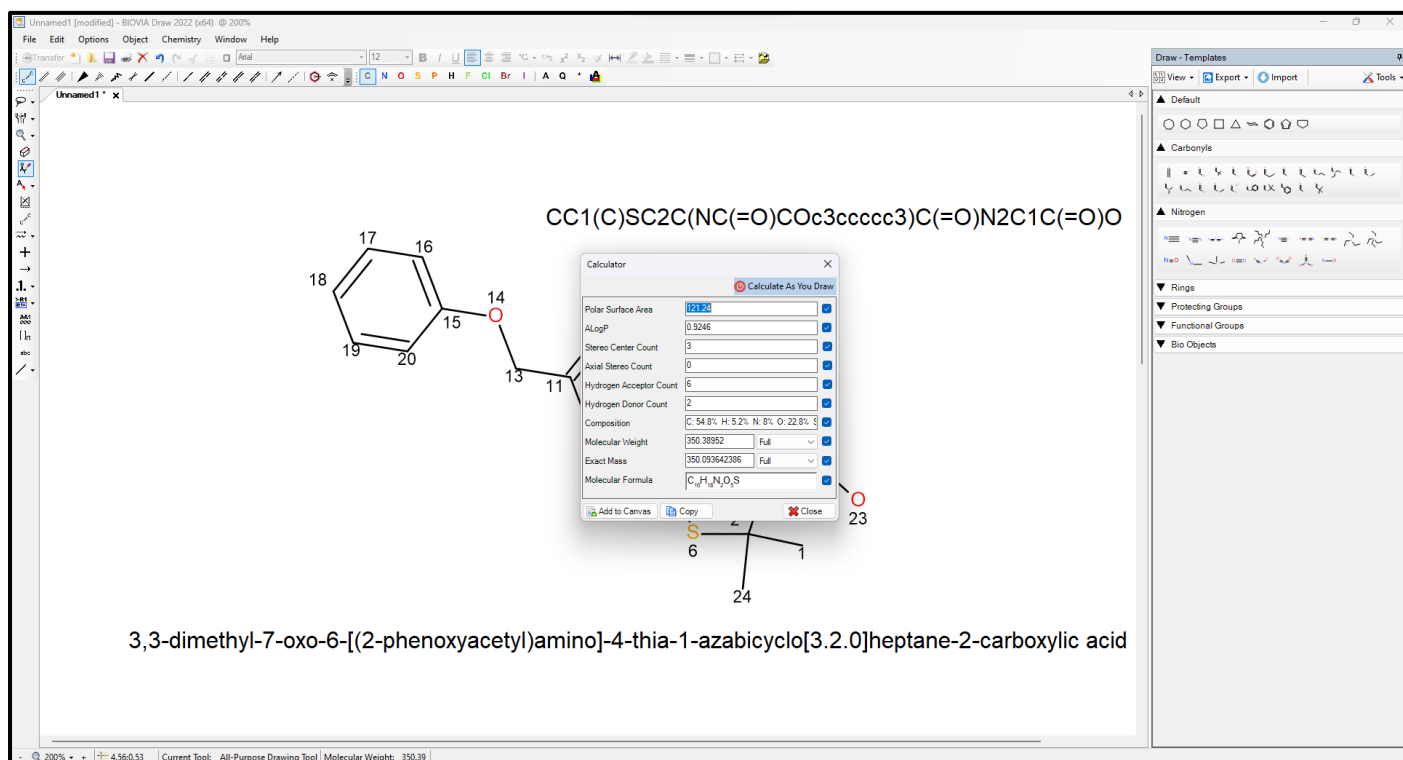


Fig 12. Different Physicochemical properties available for Penicillin

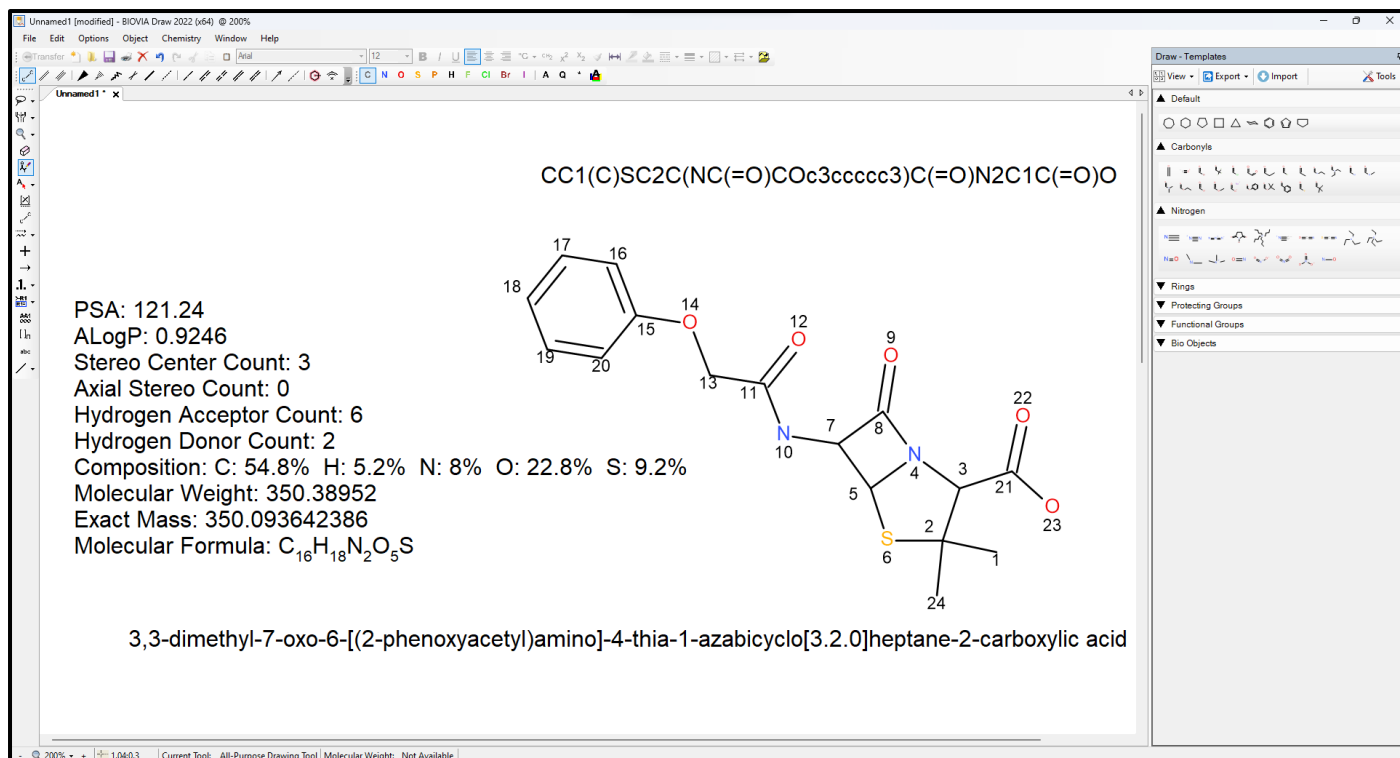


Fig 13. Calculated Physicochemical properties of Penicillin

Results:

The structure of Penicillin was drawn using the BIOVIA DRAW structure. Following are the information retrieved for the structure using software in tabulation format

SR NO	Name of Option	Results
1.	Physicochemical Properties	PSA: 121.24 ALogP: 0.9246 Stereo Center Count: 3 Axial Stereo Count: 0 Hydrogen Acceptor Count: 6 Hydrogen Donor Count: 2 Composition: C: 54.8% H: 5.2% N: 8% O: 22.8% S: 9.2% Molecular Weight: 350.38952 Exact Mass: 350.093642386 Molecular Formula: C ₁₆ H ₁₈ N ₂ O ₅ S
2.	IUPAC name	3,3-dimethyl-7-oxo-6-[(2-phenoxyacetyl)amino]-4-thia-1-azabicyclo[3.2.0]heptane-2-carboxylic acid
3.	SMILES string	CC1(C)SC2C(NC(=O)COC3=CC=CC=C3)C(=O)N2C1C(=O)O

Conclusion:

BIOVIA Draw enables the drawing and editing of complex biologics, molecules and chemical reactions. It provides structure and query drawing, allows registering, searching and reporting on chemically modified peptide or nucleotide sequences and to add structure drawing and display to user's applications, and customizing according to their organizational workflows. The software can facilitate the collaborative searching, viewing, communicating and archiving of scientific information as well.

References:

1. Klein, F., Mouquet, H., Dosenovic, P., Scheid, JF., Scharf, L., Nussenzweig, MC. (2013). Antibodies in HIV-1 vaccine development and therapy. *Science*, 341(6151), 1199-204. doi: 10.1126/science.1241144
2. Pirard, D., Vereecken, P., Mélot, C., Heenen, M. (2005). Three percent Thalidomide in 2.5% hyaluronan gel in the treatment of actinic keratoses: a meta-analysis of the recent studies. *Arch DermatolRes*, 297(5), 185-9.

WEBLEM 2B

Open Babel Tool

Aim:

To retrieve, convert & store structure for “Penicillin” query and study various file formats using Open Babel Tool

Introduction:

Penicillin V is a member of the penicillin family exhibiting broad-spectrum antibiotic property. Penicillin V binds to penicillin binding proteins (PBP), the enzymes that catalyze the synthesis of peptidoglycan, which is a critical component of the bacterial cell wall. This leads to the interruption of cell wall synthesis, consequently leading to bacterial cell growth inhibition and cell lysis.

Phenoxymethylpenicillin is a narrow spectrum antibiotic also commonly referred to as Penicillin V or Penicillin VK. It is a phenoxymethyl analog of Penicillin G, or [benzylpenicillin]. An orally active naturally penicillin, phenoxymethylpenicillin is used to treat mild to moderate infections in the respiratory tract, skin, and soft tissues caused by penicillin G--sensitive microorganisms. Phenoxymethylpenicillin has also be used in some cases as prophylaxis against susceptible organisms. While there have been no controlled clinical efficacy studies that were conducted, phenoxymethylpenicillin has been suggested by the American Heart Association and the American Dental Association for use as an oral regimen for prophylaxis against bacterial endocarditis in patients with congenital heart disease or rheumatic or other acquired valvular heart disease when they undergo dental procedures and surgical procedures of the upper respiratory tract, except for those who are at an elevated risk for endocarditis.

Open Babel is a free, open-source version of the Babel chemistry file translation program. Open Babel is a project designed to pick up where Babel left off, as a cross-platform program and library designed to interconvert between many file formats used in molecular modeling, computational chemistry, and many related areas. Open Babel includes two components, a command-line utility and a C++ library. The command-line utility is intended to be used as a replacement for the original babel program, to translate between various chemical file formats. The C++ library includes all of the file-translation code as well as a wide variety of utilities to foster development of other open source scientific software.

Methodology:

1. Open homepage for Pubchem database
2. Search for compound “Penicillin”
3. Download the compound in various file formats such as SMILES, sdf and mol file formats
4. Open the OPEN BABEL Tool and upload the structure file derived from pubchem
5. Convert the file format into the various structural file formats
6. Observe the penicillin in different file formas and interpret the results

Observations:

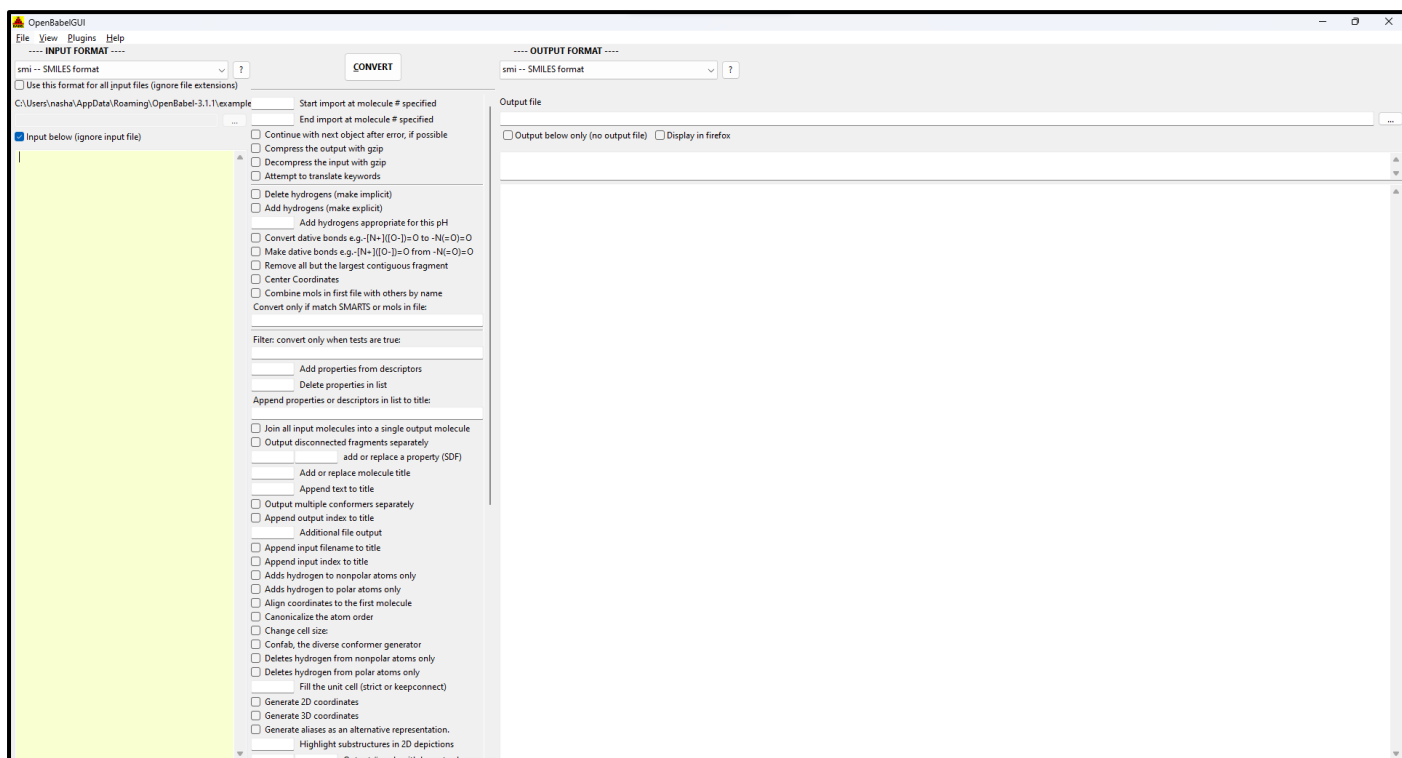


Fig 1. Homepage of OpenBabel tool

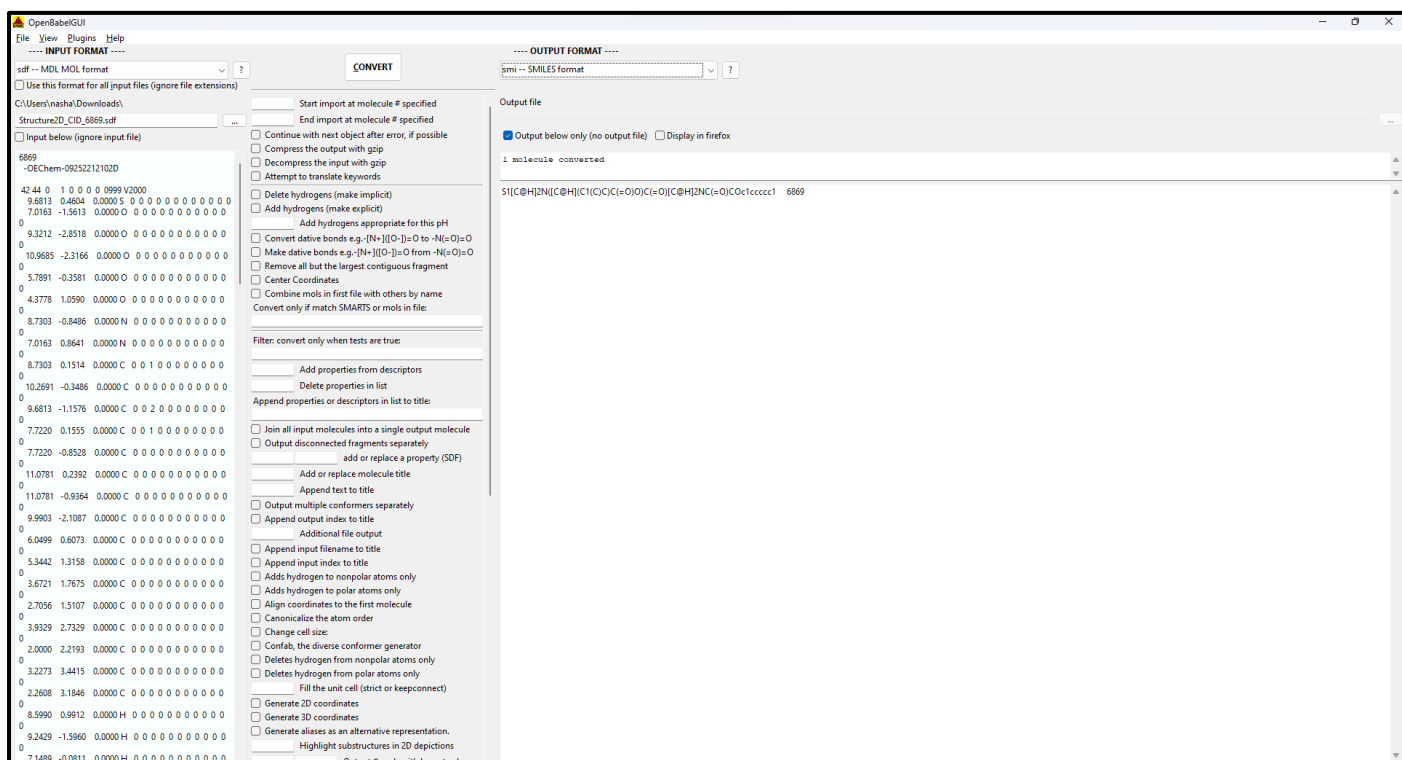


Fig 2. Convert sdf file into smiles for Penicillin

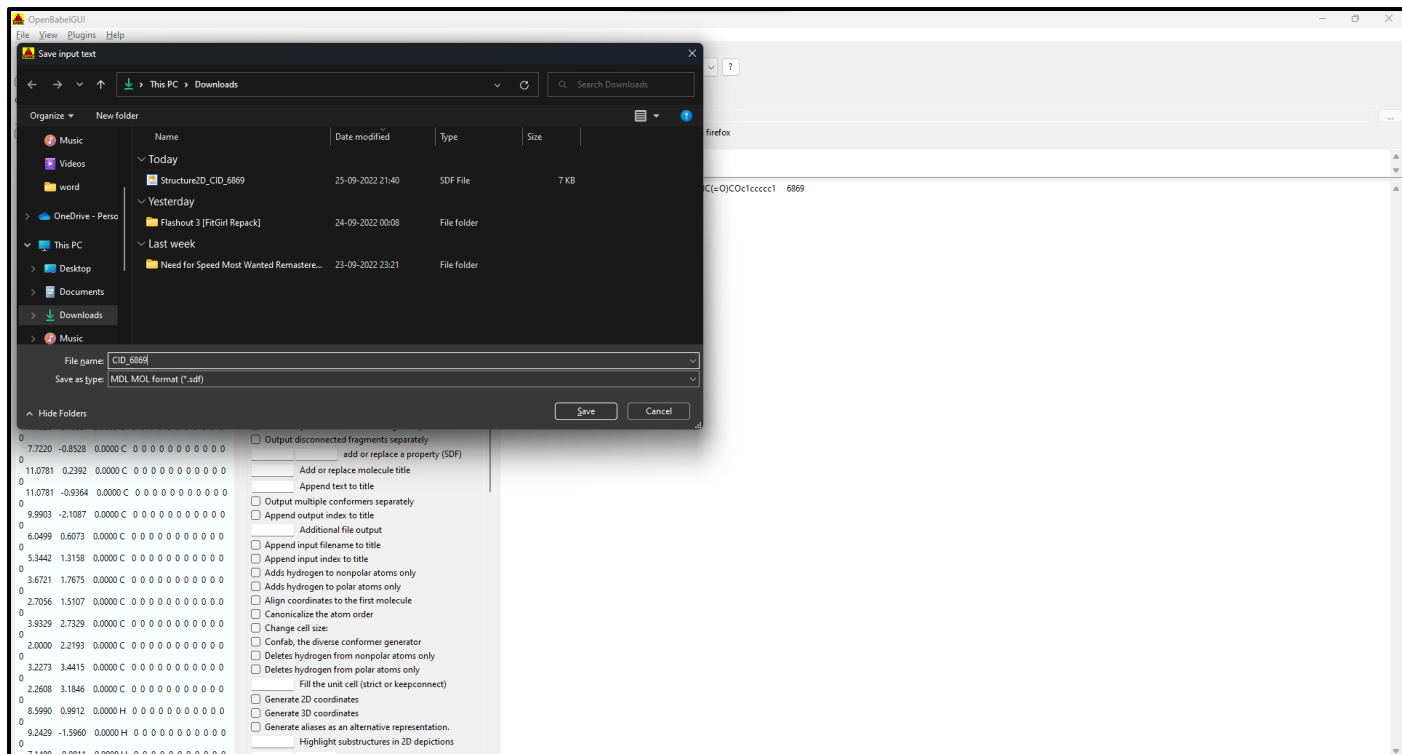


Fig 3. Storage Option for SMILES under Open Babel

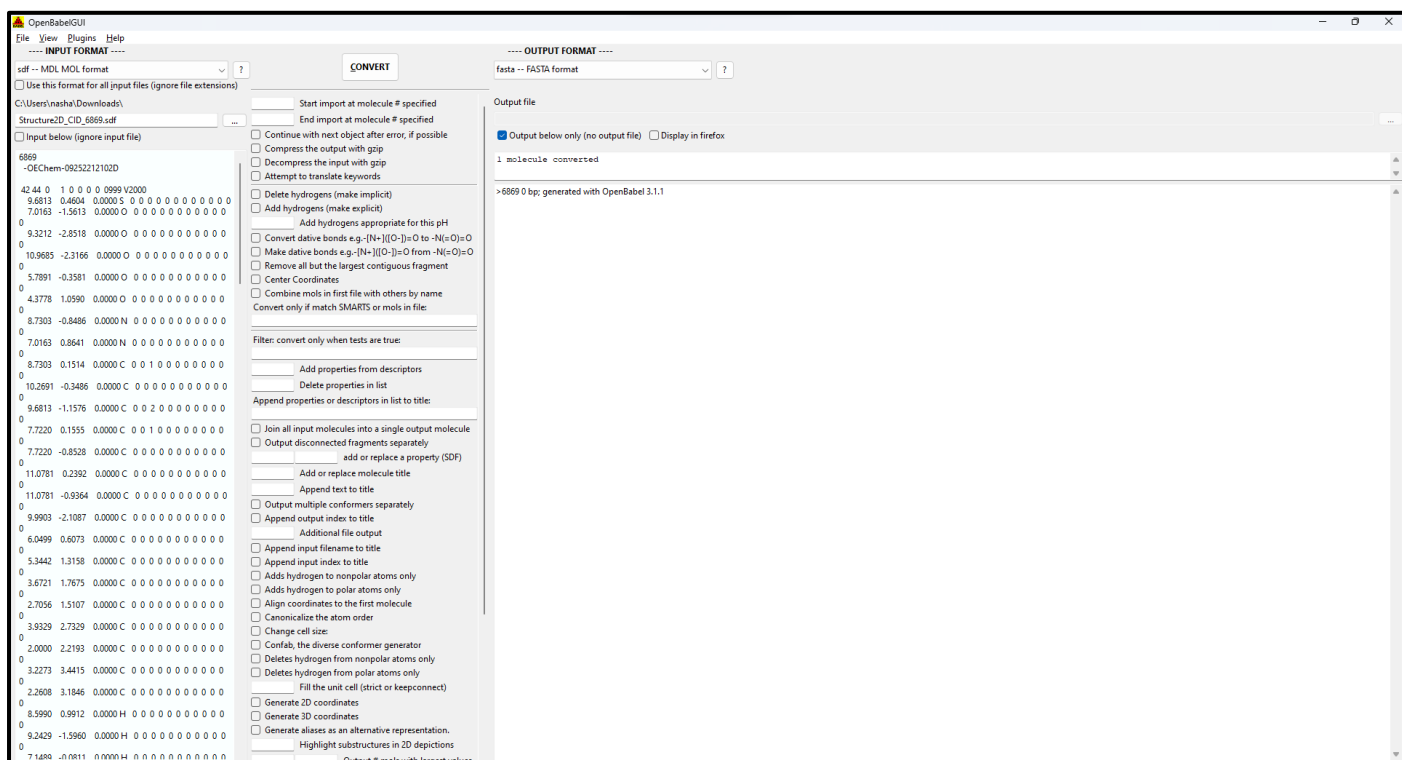


Fig 4. Convert sdf file into fasta file for Penicillin

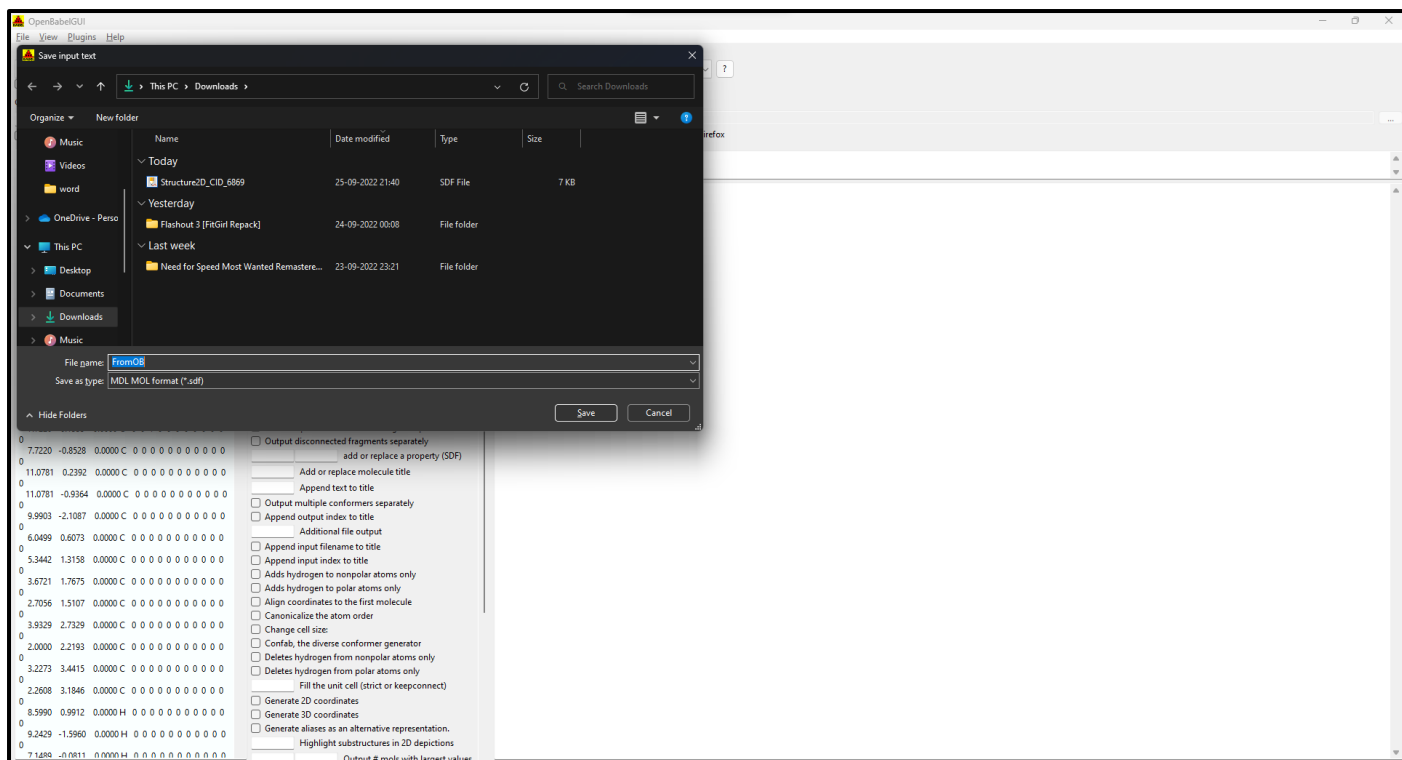


Fig 5. Storage option for fasta under Open Babel

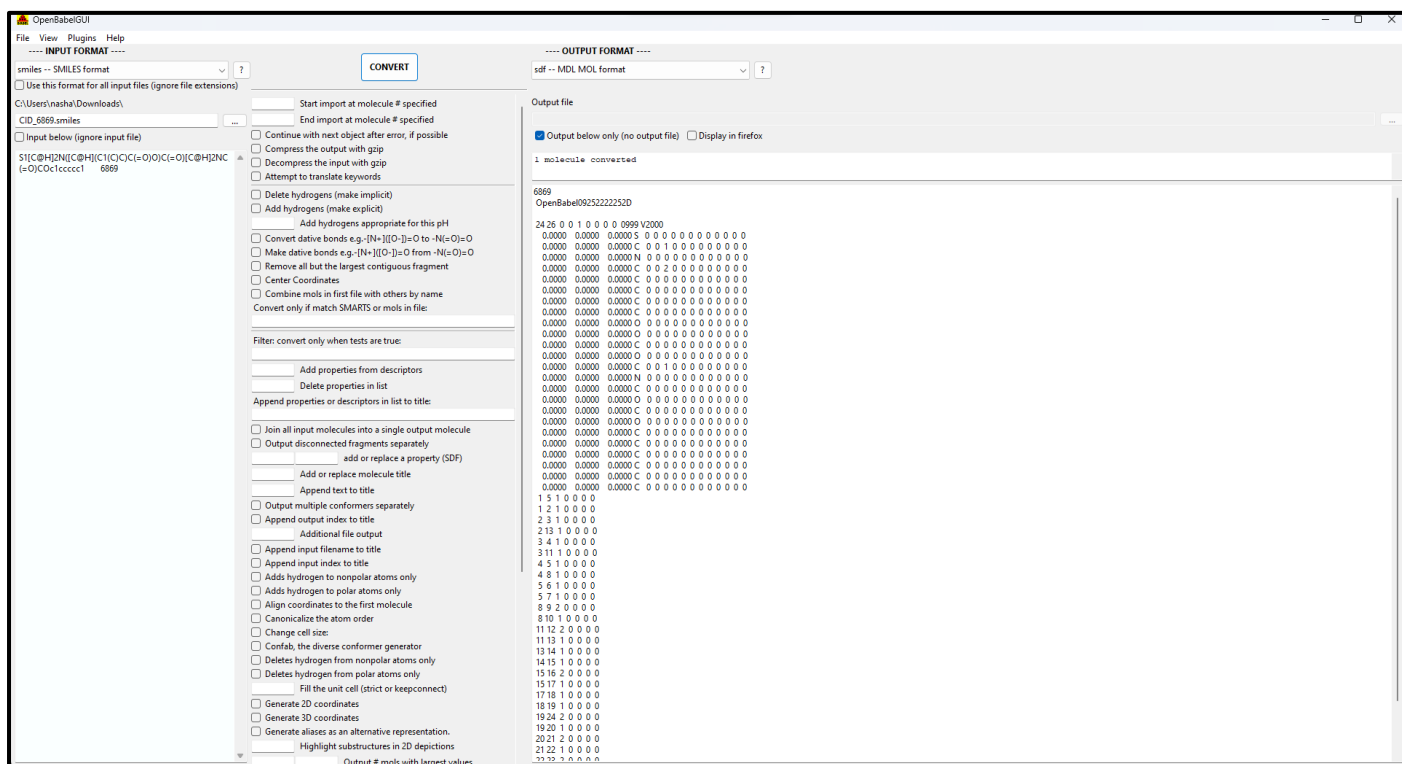


Fig 6. Convert smiles file into SDF file for Penicillin

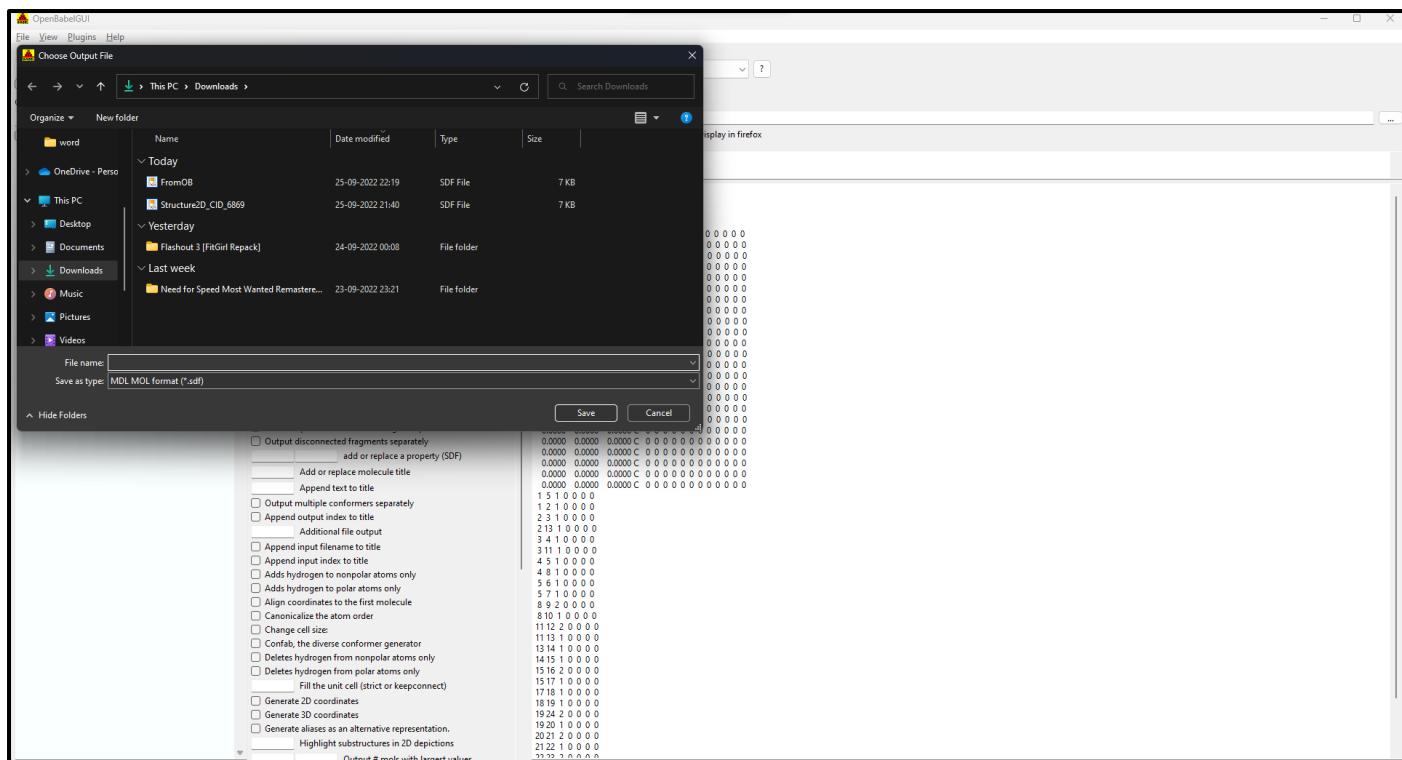


Fig 7. Storage options for sdf under Open Babel

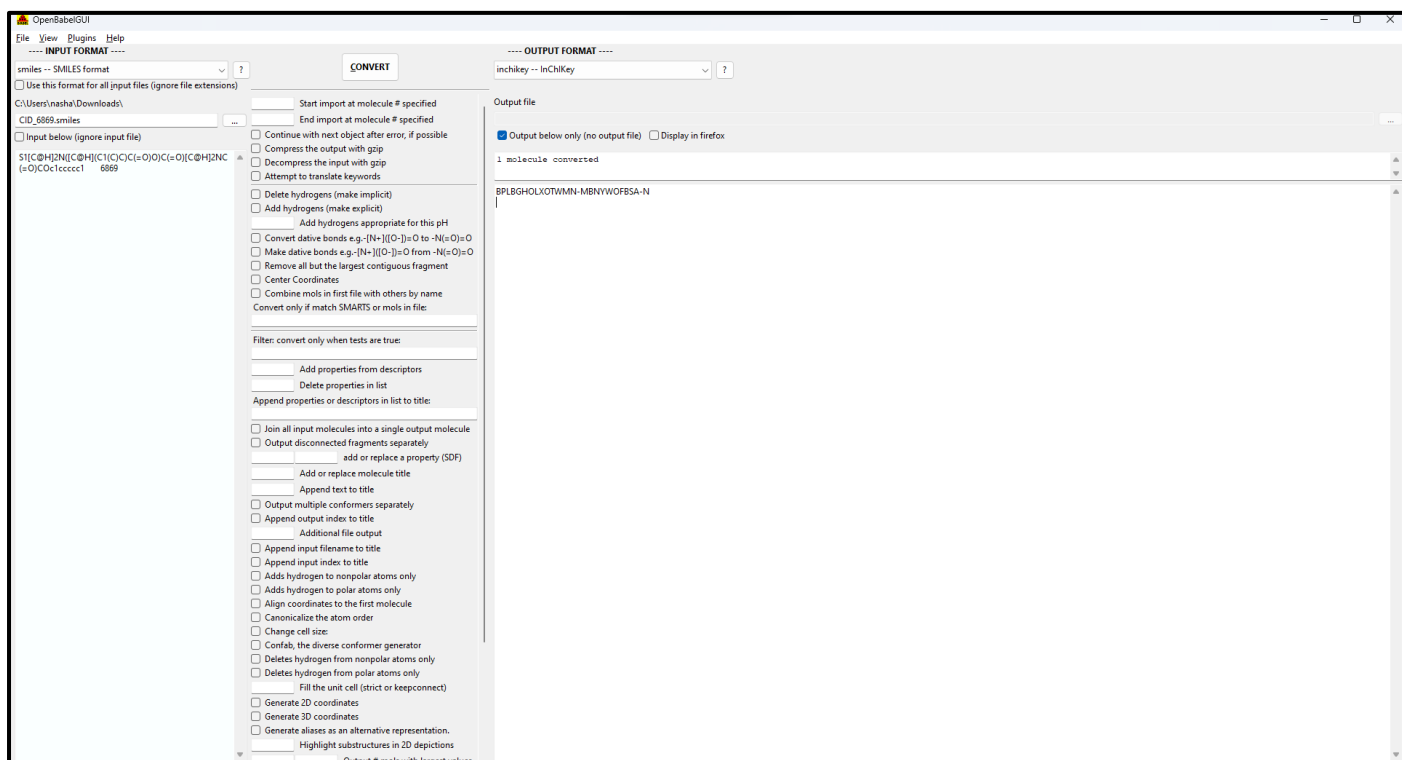


Fig 8. Convert smiles file into InChIKey file for penicillin

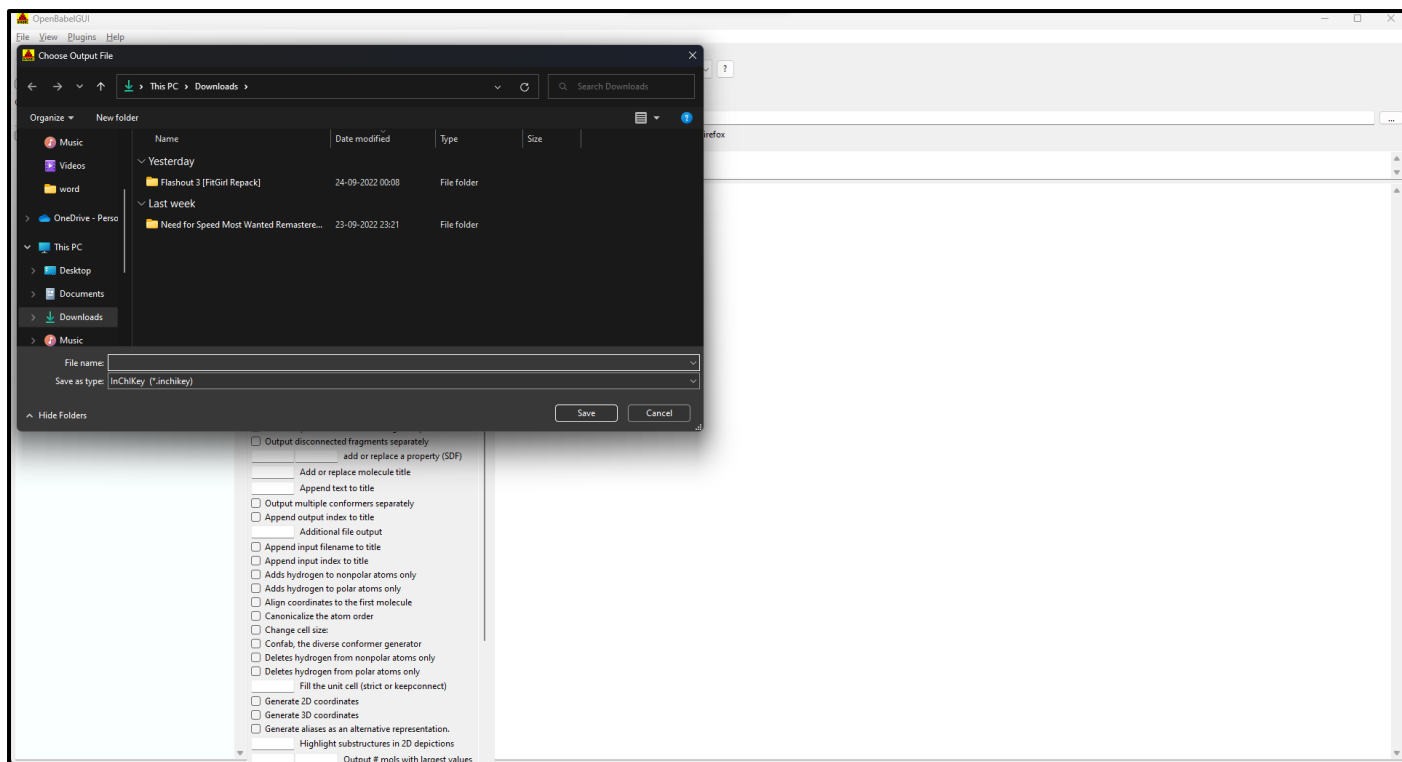


Fig 9. Storage options for InChIKey under open babel

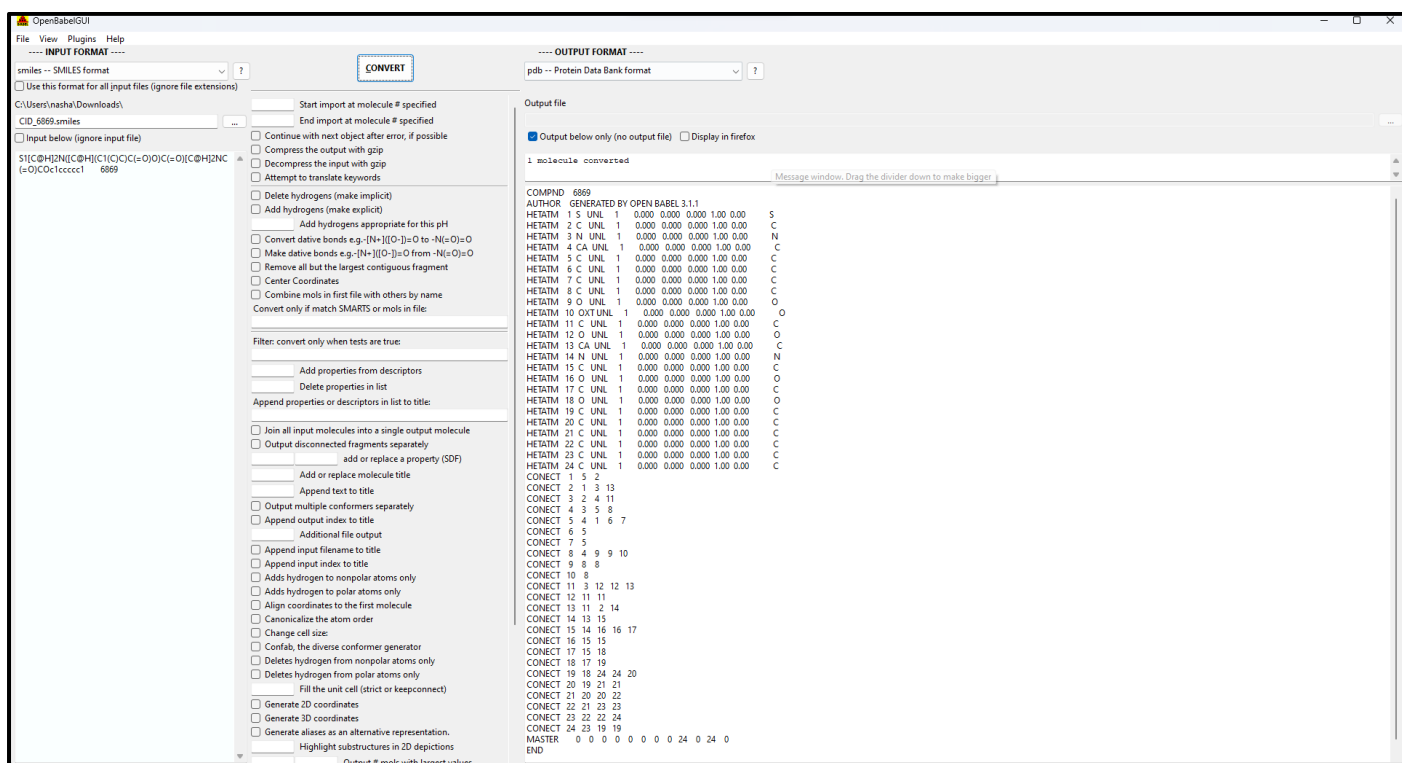


Fig 10. Convert smiles file into pdb file for penicillin

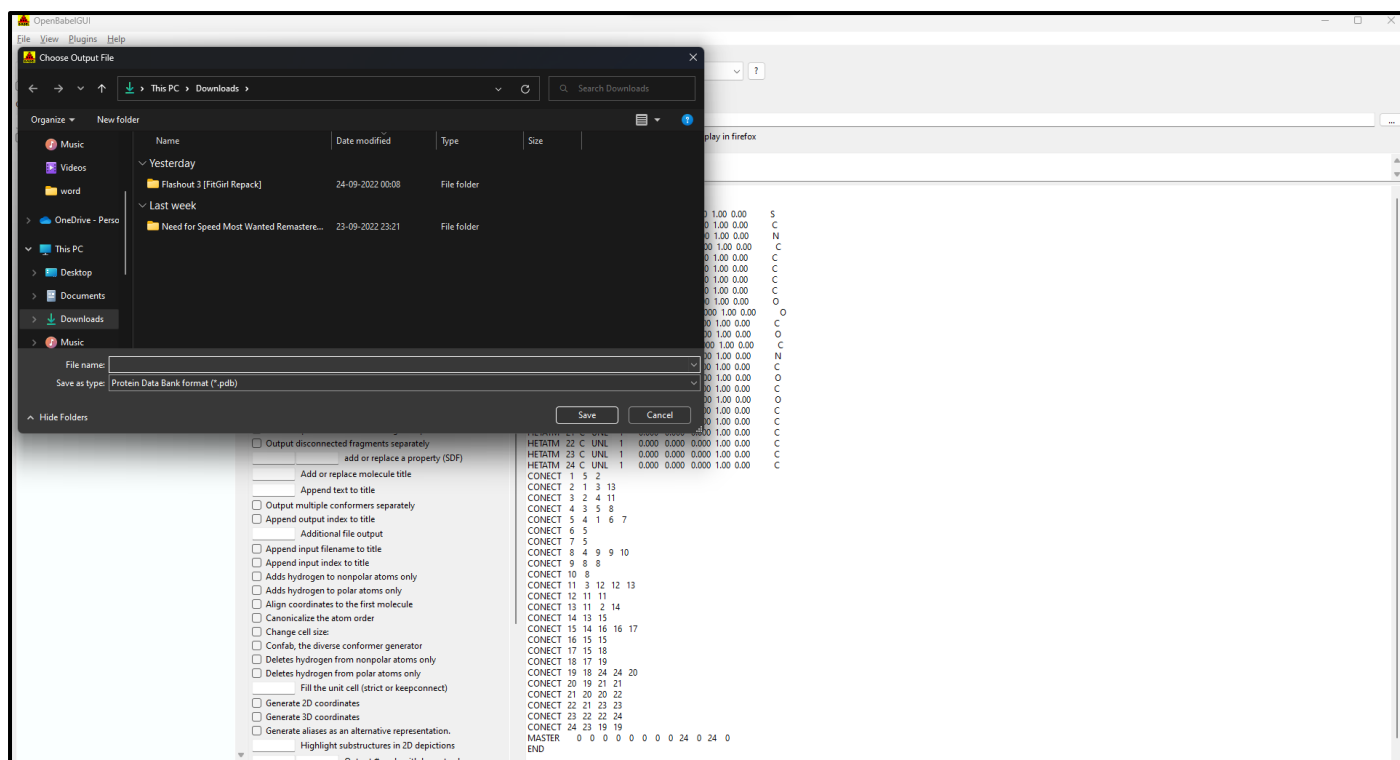


Fig 11. Storage Option for pdb under Open babel

Results:

The structure for Penicillin was retrieved from PUBCHEM in sdf file format. By using Open Babel tool, it was possible to convert various structural file formats into various other formats such as SMILES, MDL MOL, fasta, pdb, InChI key and the output is saved in the desired format for further studies

Conclusion:

The data of the chemical compound “Penicillin” was studied using different file formats. Therefore, the Open Babel tool presents a solution to the increase of multiple chemical file formats. It gains by way of its users, contributors, developers, related projects, and the general chemical community. It allows searching, converting, analyzing, or storing data from molecular modeling, chemistry, solid-state materials, biochemistry, or related areas. It provides both ready-to-use programs as well as a complete, extensible programmer's toolkit for developing cheminformatics software.

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2. Phytochemistry, Pharmacology and Medicinal Uses of Plants of the Genus Salix: An Updated Review. Front Pharmacol, 12(593856).
3. Weininger, D. (1988). SMILES, a chemical language and information system. 1. Introduction to methodology and encoding rules. J Chem Inf Comput Sci. 28, 31-36.