

# **Tutorial of molecular generation and drug modification by MolAICal and deep learning model**

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## 1. Introduction

In this tutorial, two examples are introduced: one is drug-like and FDA-like molecular generation that can be used for drug virtual screening, the other is scaffolds and groups modification based on a known ligand, which can be used to modify or design new drugs. For more detailed MolAICal, please read this paper (<https://doi.org/10.1093/bib/bbaa161>).

## 2. Materials

### 2.1. Software requirement

1) MolAICal: <https://molaical.github.io>

### 2.2. Example files

1) All the necessary tutorial files are downloaded from:

<https://github.com/MolAICal/tutorials/tree/master/016-AImolecularGeneration>

## 3. Procedure

### 3.1 Generate drug-like and FDA-like dataset

1) Generating FDA-like molecules.

```
#> molaical.exe -dock AI -s FDAMol -n 30 -nf 10 -nc 3 -g on -b on -v off
```

It will generate a file named tmpGenVS.dat that contains 30 molecular SMILES strings, and 3 folders named tmpGenMols1, tmpGenMols2, and tmpGenMols3. Users can generate much more molecules according to their needs. The generated 3D molecules are in PDBQT format by default. If users need to change the molecular format, for instance, they can use the below command for format conversion:

1) Adding hydrogen (option)

```
#> molaical.exe -dock addh -i 1.pdbqt
```

2) Changing “pdbqt” to “pdb” format

```
#> molaical.exe -dock pdbqt2pdb -i 1.pdbqt
```

2) Generating drug-like molecules.

```
#> molaical.exe -dock AI -s ZINCMol -n 30 -nf 10 -nc 3 -g on -b on -v off
```

It will generate a file named tmpGenVS.dat that contains 30 molecular SMILES strings, and 3 folders named tmpGenMols1, tmpGenMols2, and tmpGenMols3. Users can generate much more molecules according to their needs. The generated 3D molecules are in PDBQT format by default. If users need to change the molecular format, for instance, they can use the below command for format conversion:

1) Adding hydrogen (option)

```
#> molaical.exe -dock addh -i 1.pdbqt
```

2) Changing “pdbqt” to “pdb” format

```
#> molaical.exe -dock pdbqt2pdb -i 1.pdbqt
```

### 3.2 Generate new similar molecules based on a known ligand

Webserver could supply a convenient way for the researchers to design drugs without any special software and hardware requirements via the browsers. MolAICal supplies this function via webserver which URL can be obtained by the below command:

```
#> molaical.exe -model ligdream
```

```
E:\workdir\MolAICal\tutorial\tutorial>cd ../../
E:\workdir\MolAICal>e:\workdir\MolAICal\create\version\MolAICal-win64\molaical.exe
# Project leader: Qifeng Bai, the official site of MolAICal: https://molaical.github.io
The latest URL is: https://heisenberg.ucam.edu:5000 URL
E:\workdir\MolAICal>
```

1) Input URL <https://heisenberg.ucam.edu:5000> into the browser, you can choose the way for drawing your molecule or input your SMILES string (see Figure 1). Finally, click “Submit and Running”.

Drug design by AI

Job title:  Optional job title      Email:  Optional email

Draw a structure

Option 1: draw your molecule

Option 2: input your molecular SMILES string

Number of generated molecules

☐ Canonical SMILES

SMILES:  C1=CC(C)=CC=C1CC2=CC(CO)=C

Number of generated molecules:  60

Figure 1

2) It will turn to another result web page (see Figure 2).

### The current job status information:

Job ID: f7f6a1f9-7b11-4ae3-858c-a7494eb8499b

Status: finished

Created at: Fri, 08 Oct 2021 15:24:23

Notice: This page will refresh at intervals!!! Please save below url if you donot want to wait for results. You can load this url for checking the final results after you are offline for long time.

<https://heisenberg.ucam.edu:5000/static/upload/210-26-124-238/d113b6f8-284b-11ec-9239-7f4cee7db04.html>

### The generated molecular file:

Download generated files

Download the generated molecules with formats of SMILES and Mol2.

Note: SMILES.dat contains the generated molecules with SMILES format. The molecules with mol2 style use the number as prefix.

### View the generated molecules:

Notice: to improve the display speed, 10 molecules are loaded at the most. If you want to check more molecules, please load them manually (see documentation).

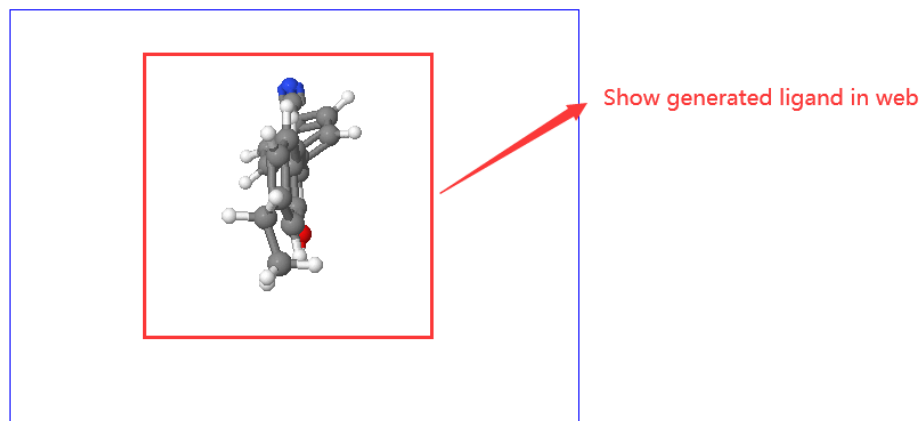


Figure 2