

## **Setup and run automated docking docking**

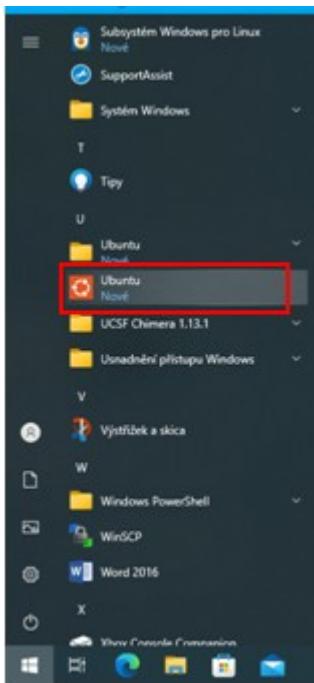
<https://github.com/ci-lab-cz/docking-scripts>

### **Install (on own laptops/computers)**

```
conda create -n vina-c conda-forge python=3.9 numpy=1.20 rdkit scipy  
dask distributed vina  
  
conda activate vina  
  
pip install meeko moldock
```

### **Run on computers:**

Run Ubuntu on Windows machines



The Ubuntu console will be opened and you have to activate the environment by

```
conda activate vina
```

```

student@DESKTOP-IDPFL93:~ 
To run a command as administrator (user "root"), use "sudo <command>". 
See "man sudo_root" for details.

Welcome to Ubuntu 22.04.1 LTS (GNU/Linux 5.15.79.1-microsoft-standard-WSL2 x86_64)

 * Documentation: https://help.ubuntu.com
 * Management: https://landscape.canonical.com
 * Support: https://ubuntu.com/advantage

This message is shown once a day. To disable it please create the
/home/student/.hushlogin file.
(base) student@DESKTOP-IDPFL93:~$ conda activate vina
(vina) student@DESKTOP-IDPFL93:~$ 

```

To run docking on a local machine:

```
vina_dock -i ligands.smi -o out.db -p protein_prepared.pdbqt -s config.txt --sdf -c 3 -v --no_protonation
```

Top 1 docked poses will be stored in `out.sdf` file at the same directory as `out.db` when the docking will finish

To retrieve some other data from db one may use SQLiteBrowser (free cross-platform tool) or by a script:

	<b>id</b>	<b>smi</b>	<b>smi_protonated</b>	<b>rce_mol_blo</b>	<b>nol_block_prc</b>	<b>docking_score</b>	<b>pdb_block</b>	<b>mol_block</b>	<b>time</b>
1	1ke5_ligand	CNS(=O)(=O)c...	NULL	NULL	NULL	-7.882	MODEL 1REMARK ...	1ke5_ligand ...	2023-02-02 09:17:35
2	1ke6_ligand	CNS(=O)(=O)...	NULL	NULL	NULL	NULL	NULL	NULL	NULL
3	1ke7_ligand	O=C1Nc2ccc(-...)	NULL	NULL	NULL	-9.685	MODEL 1REMARK ...	1ke7_ligand ...	2023-02-02 09:17:52

- get top 2 poses for every ligand

```
get_sdf_from_dock_db -i out.db -o 2.sdf --poses 1 2
```

- apply some additional conditions when retrieving of sdf – retrieve top poses of molecules having docking score below -8:

```
get_sdf_from_dock_db -i out.db -o 3.sdf --add_sql 'docking_score < -8'
--fields docking_score
```