NAME: ASIF ERFAN KHAN

ROLL NUMBER: 546

COURSE: MSc CS

SUBJECT: BIOINFORMATICS

TOPIC: RETRIEVING 3D

STRUCTURE FROM PDB

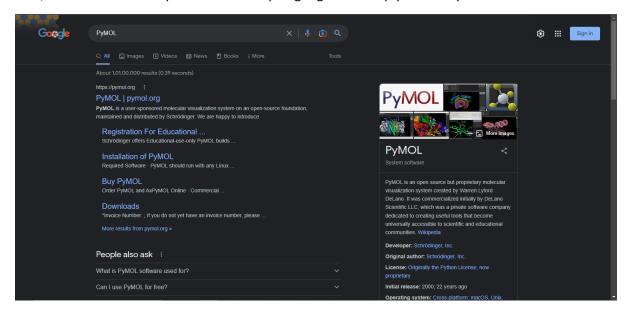
Aim: Retrieving 3D structure from PDB

To perform the current practical, you'll be needing two things

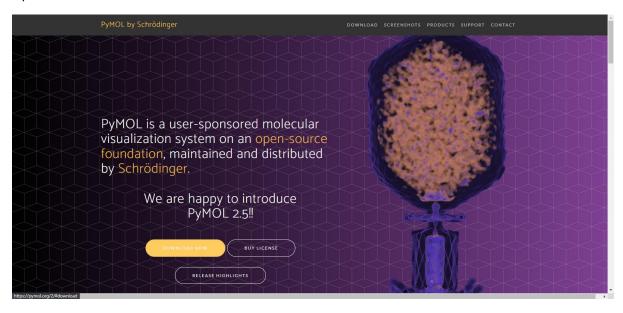
- PyMOL (software)
- II. Protein in .pdb format

Installing PyMOL Software

First, we need to install PyMOL. To do so open google and simply search PyMOL



Open the first link and click Download Now



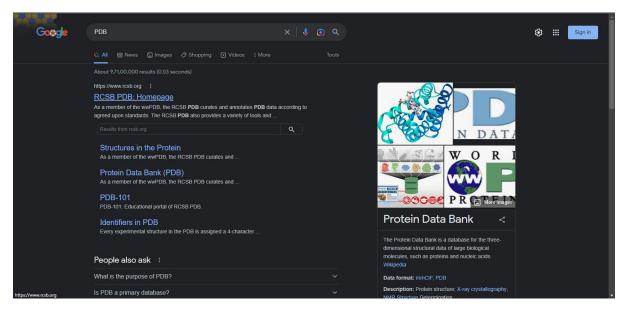
When done downloading install the software.

This is how the interface of the software looks like.

```
PAPOL (TIO 2,5.4 - Incentive Product
Copyright (T) Schrödinger, Lic
This Executable Build integrates and extends Open-Source PytOL,
This Executable Open-Source PytOL,
This Executable
```

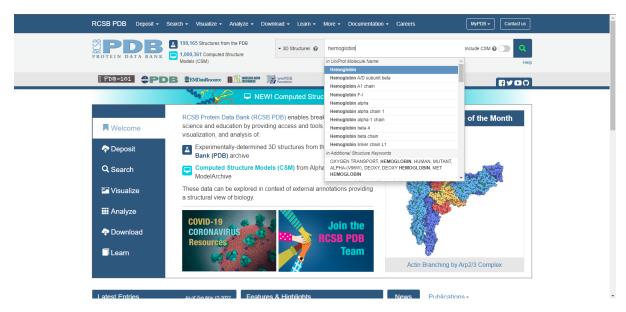
Downloading Protein in .pdb format

To download Protein open, google and search pdb or simply visit https://www.rcsb.org/

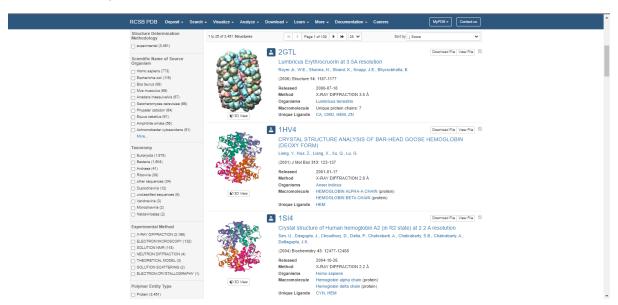


Open the first link named "RCSB PDB: Homepage"

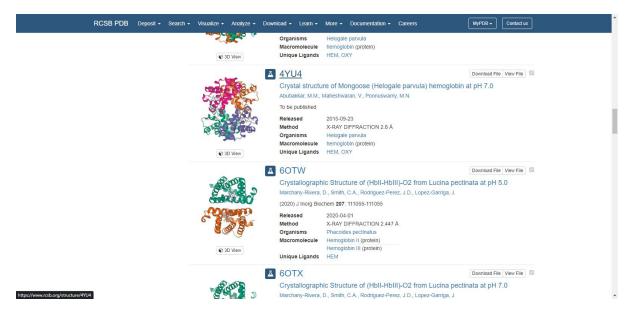
On the search bar search for "hemoglobin"



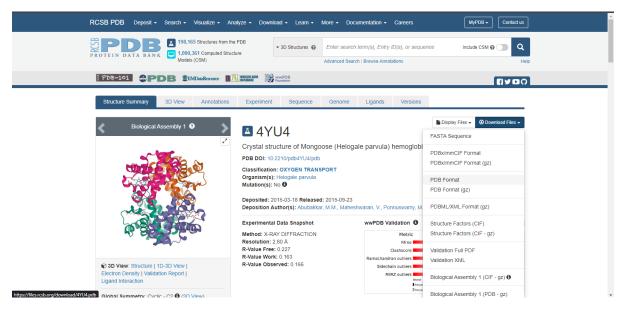
You'll see all the proteins listed below like this



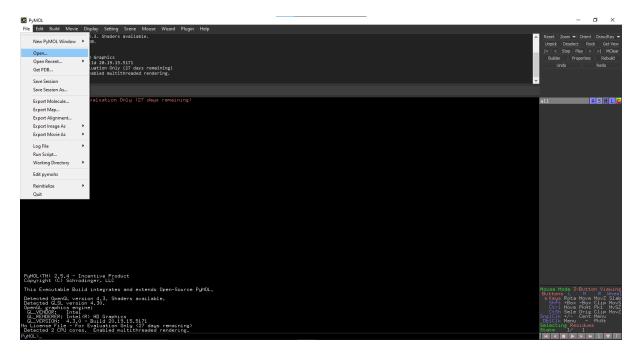
Scroll down until you find "4YU3" and open it or you can directly search for "4yu4"



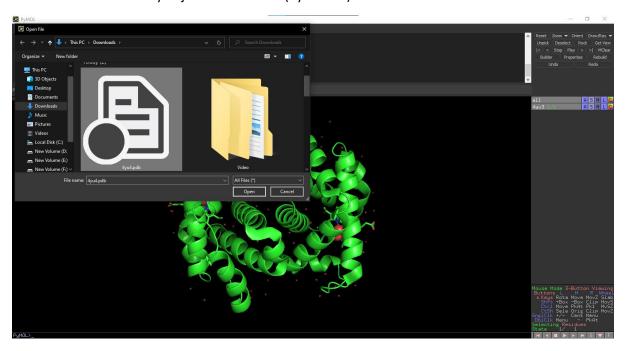
On the right-hand side, you'll the download option. Click on it and download as PDB format



Now that protein is downloaded open PyMOL and on top left corner click File > open



Now browse for the file you just downloaded (4yu4.PDB)



This is how the screen of imported file looks like

