1) Open the Terminal app. Determine the path to the GliomaToolV1 folder in File Explorer. Paste the following command into the terminal to navigate to the folder.

cd path/to/GliomaToolV1

2) To check if you are in the right folder, type the following command.

ls

The result should show the same files listed below.

avnith@Avniths-MacBook-Air GliomaToolV1 - Copy % ls app.py models requirements.txt static templates testImages avnith@Avniths-MacBook-Air GliomaToolV1 - Copy %

3) You will need to have Python 3 and pip already installed on your computer. To run GlioGrade, install all necessary packages with

pip install -r requirements.txt

4) Paste the following command to start running the Flask app

flask --app app.py run

This may take a minute, but the result should look like this:



5) Copy the URL that is given in the output of the previous command. In the image above, it is <a href="https://l27.0.0.1:5000">https://l27.0.0.1:5000</a>. Open a new browser tab and navigate to this page. You should see this screen.

## **Glioma Detection Tool**

Upload T1 and T2 MRI scans to detect glioma types and grades



- 6) To use this tool, upload an axial slice of an MRI scan. Note that you will need to upload both a T1 and T2 MRI image. Select Analyze Scans to view the results.
- 7) There are sample images given in the testImages folder which you can use. Any uploaded MRI images and before and after processing will be stored in the static folder.
- 8) To exit the app, go back to the terminal and press Ctrl + C to stop running the Flask app. All uploaded MRI images will be deleted from the static folder.