


Execution Of Project

Step 1: Download Anaconda

1. Go to the official Anaconda website:
 <https://www.anaconda.com/download>
 2. Select your operating system:
 - Windows (64-bit installer)
 - Mac (Apple Silicon or Intel)
 - Linux
 3. Click Download and wait for the file to be saved.
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Step 2: Install Anaconda

Windows

1. Open the downloaded .exe installer.
2. Click Next and accept the license agreement.
3. Select "Just Me" (recommended for personal use) and click Next.
4. Choose an installation location (default is recommended).
5. Check the box:
 - "Add Anaconda to my PATH environment variable" (Optional but useful for command-line access)
6. Click Install and wait for it to finish.
7. Click Finish and launch Anaconda Navigator.

Mac

1. Open the .pkg file.
2. Follow the on-screen instructions, accepting the defaults.
3. After installation, open a terminal and verify the installation by running:

bash

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conda --version

4. Open Anaconda Navigator from Applications.

Linux

1. Open a terminal and go to the download location.
2. Run the following command:

```
bash
```

```
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```

```
bash Anaconda3-*.sh
```

3. Follow the installation steps and accept the defaults.
4. After installation, run:

```
bash
```

```
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```
source ~/.bashrc
```

```
conda --version
```

5. If the version appears, Anaconda is installed correctly.
6. Launch Anaconda Navigator using:

```
bash
```

```
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```

```
anaconda-navigator
```

Step 3: Open Jupyter Notebook from Anaconda Navigator

1. Launch Anaconda Navigator
 - Open Anaconda Navigator from the Start Menu (Windows) or Applications (Mac/Linux).
2. Select Your Environment
 - Go to the Environments tab.
 - Choose the environment where your Patient Routing RAG project is installed.
 - Click Open Terminal (or click the Play button and select Open with Jupyter Notebook).
3. Launch Jupyter Notebook

- If Jupyter Notebook is installed in your environment, click Launch under Jupyter Notebook.
 - If not installed, go to the Home tab, search for Jupyter Notebook, and click Install.
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Running .py Files Using the Run Button in Anaconda Navigator

1. Open Anaconda Navigator
 - Launch Anaconda Navigator from the Start Menu (Windows) or Applications (Mac/Linux).
2. Select the Correct Environment
 - Go to the Environments tab.
 - Choose the environment where your patient_routing.py script is installed.
 - Click on Open with Terminal (or Jupyter Notebook if needed).
3. Run the Python Script from the Navigator
 - In the Home tab, under the environment where your script is located, click Run.
 - Browse for Patient Routing RAG Generation and Recommendation.py.
 - Click Run.