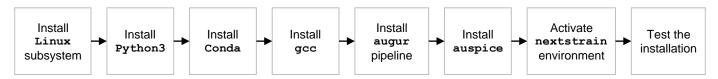
## **Nextstrain installation**

## Yale school of public health

This tutorial is designed to assist you in installing nextstrain and its dependencies. The workflow below explains the main steps to accomplish that. Windows users must install a Linux subsystem first (see step 1).



(1) nexstrain only operates on UNIX systems (MacOS or Linux). As a workaround, Windows users must install Linux (Ubuntu) as a subsystem. Visit the link below ans check the instructions "For windows users" to install and set up Linux on Windows:

https://github.com/grubaughlab/nextstrain course

(2) Access the Terminal in your system, and run the commands below to check if Python3 is installed (see box):

python --version

If no version information is shown, it means the software is not installed. Please **access** the sites below to install it:

 $Python 3 \rightarrow https://www.python.org/downloads/$ 

(3) While it may be available on MacOS, Linux subsystems may not come with a gcc compiler. **Test** if gcc is available by typing:

gcc --version

**(4)** If such command does not tell you which gcc version is installed, Linux users must **install** it by typing the commands below, one line at a time (MacOS users, please see note):

sudo apt-get update
sudo apt install gcc
gcc --version

Note: apt (Advanced Package Tool) is a software used for installing and removing other software in Linux only. MacOS users should **use** brew instead (first **type** brew -h to check if it's installed. If not installed, **access** https://brew.sh/)

brew update
brew upgrade
brew install gcc

(5) Check if conda is installed by typing conda list in your Terminal. If that returns a list of software and packages, proceed to step 7. Otherwise, visit the link below, **download** a Python3 Miniconda installer (in bash format .sh, or in executable format .pkg) compatible with your operating system (Linux or Mac), and proceed to step 6 to install Miniconda.

https://docs.conda.io/en/latest/miniconda.html

(6) Access the directory where the Miniconda installer is located. Execute the file (if in .pkg format), or type bash followed the name of the file you downloaded (if in .sh format), as shown below:

bash Miniconda3-latest-<YourOperatingSystem>-x86\_64.sh
conda list

(7) If Python3, gcc and conda are installed, run the commands below, one line at a time, to install augur:

python3 -m pip install nextstrain-cli
curl http://data.nextstrain.org/nextstrain.yml --compressed -o nextstrain.yml
conda env create -f nextstrain.yml

(8) Activate the nextstrain environment using the command below. It will change the prefix of the command prompt (please see the 'nextstrain' prefix in your Terminal):

conda activate nextstrain

(9) Now let's install auspice. First, check if npm is available (see box):

npm --version

If npm is not installed, type the command below to **install** it:

sudo apt install nodejs

(10) With npm installed, run the following command to install auspice, the software we are going to use to visualize the results generated by augur:

npm install --global auspice

(11) That's it. Now **run** the commands below to test the software installation. These commands will print information about how to use each software:

augur -h
auspice -h
nextstrain -h
conda deactivate

All set! 🏂