

For DL and NLP demos

***ETI – Estimated Time to Install when tested on a 16GB RAM, i7 11th gen laptop with ~30 Mbps internet connection speed**

To install PyTorch (ETI <15 min - downloads about 2.7 GB of contents)

```
>> pip3 install torch torchvision torchaudio --index-url https://download.pytorch.org/whl/cu118
```

To create a new environment in conda:

```
>> conda create --name nlp_projects
```

```
>> conda activate nlp_projects
```

To install spaCy:

Option 1: Using conda (ETI < 12 min)

```
>> conda install -c conda-forge spacy
```

```
>> python -m spacy download en_core_web_sm
```

OR

Option 2: Using pip

```
>> pip install -U pip setuptools wheel
```

```
>> pip install -U spacy
```

```
# To download models for using trained pipelines:
```

```
>> python -m spacy download en_core_web_sm
```

For more info: <https://spacy.io/usage#installation>

Troubleshooting guidelines:

- (1) If installing packages through conda is taking a very long time i.e. stuck at solving environment step, it is very likely that there are too many packages in base environment. In general, it's always a best practice to have the base environment loaded with minimal number of packages and use miniconda as against conda.

Enter 'Ctrl + C' or 'Ctrl + Z' to forcefully exit and then try following to deinstall unnecessary packages from base environment:

```
>> conda install --rev 0
>> conda update conda
>> conda update conda-build          # Optional
```

If removing dependencies is not an option, one could try the following command:

```
>> conda install conda=22.9.0
```

Alternatively, one could use libmamba solver (a faster option by Anaconda):

```
# install mamba
>> conda install -n base conda-forge::mamba
>> conda config --set solver libmamba

# use mamba
>> mamba install <package-name>
```

If installing mamba is not an option (for corporate issued laptops that use Z-scanner), try:

```
>> conda config --set channel_priority flexible
```

A faster alternative to mamba is to use micromamba that can be installed via homebrew on MacOS or using PowerShell on Windows OS:

<https://mamba.readthedocs.io/en/latest/installation/micromamba-installation.html>

- (2) If you encounter Microsoft C++ build error during installation of scispaCy or medspaCy (error screenshot enclosed below):

```
error: Microsoft Visual C++ 14.0 or greater is required. Get it with "Microsoft C++ Build Tools": https://visualstudio.microsoft.com/visual-cpp-build-tools/
[end of output]
```

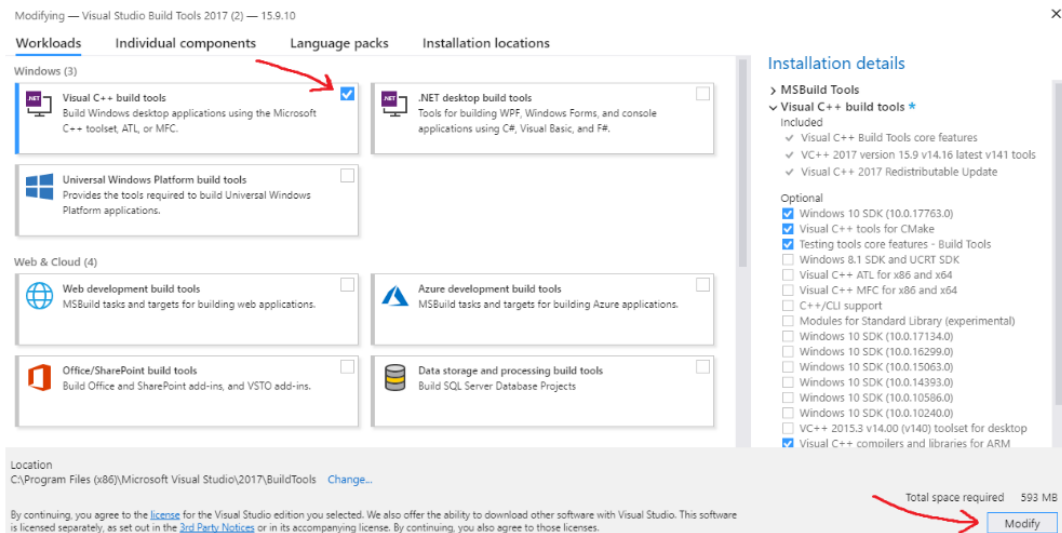
[ETI: < 15

min]

- (a) Go to <https://visualstudio.microsoft.com/visual-cpp-build-tools/> and click 'Download Build Tools'. Once the download is complete, run the .exe file by double clicking.



- (b) Just choose the 'Visual C++ build tools' option by pressing the checkmark and then click 'Modify' or 'Install' at bottom right corner of the screen to begin installation.



- (c) After the C++ tools finish installing, reboot the computer
- (d) Now go to the anaconda prompt, activate the conda environment, and run the pip command again. It should work this time!