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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-scaler), 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

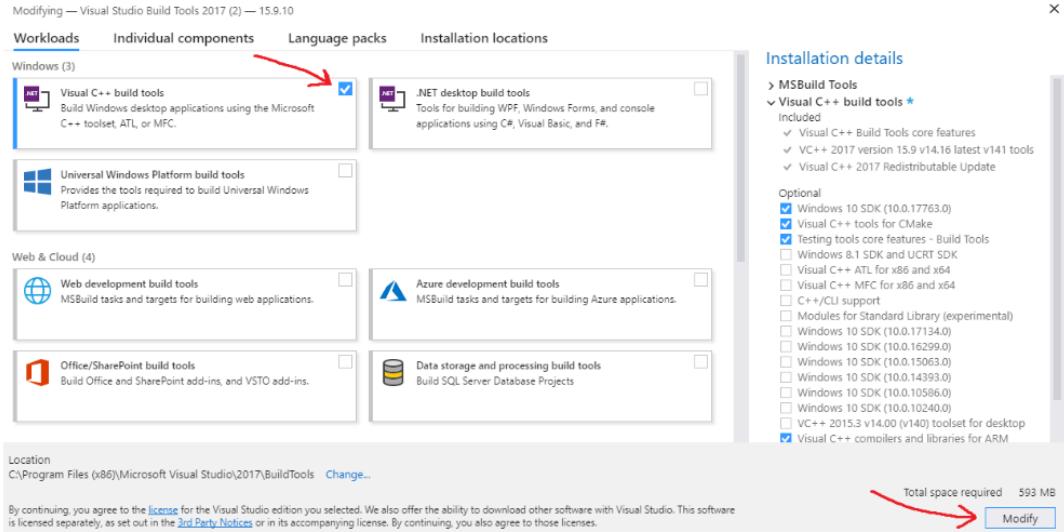
min]

- (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.



The Microsoft C++ Build Tools provides MSVC toolsets via a scriptable, standalone installer without Visual Studio. Recommended if you build C++ libraries and applications targeting Windows from the command-line (e.g. as part of your continuous integration workflow). Includes tools shipped in Visual Studio 2015 Update 3, Visual Studio 2017, Visual Studio 2019, and latest version of Visual Studio 2022.

- (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!