

Computing for Medicine: Phase 3, Seminar 4 Project

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Based on slides by Michelle Craig

Package Installation

> conda activate C4M (Windows)

> source activate C4M (Mac)

Packages needed for this project:

- numpy
- scipy
- scikit-learn
- matplotlib
- **scikit-image**
- **joblib**

Package Installation

- > conda install -c anaconda numpy
- > conda install -c anaconda scipy
- > conda install -c anaconda scikit-learn
- > conda install -c conda-forge matplotlib

New for this project:

- > conda install -c conda-forge scikit-image
- > conda install -c anaconda joblib

Starter code and data

Starter code

- `image_processing_tutorial.py`
- `nuclei_detection_tutorial.py`
- `project_helpers.py`
- `project.py`

Data

- 100 H&E stained histology images of colorectal adenocarcinomas
- Sirinukunwattana et al., 'Locality Sensitive Deep Learning for Detection and Classification of Nuclei in Routine Colon Cancer Histology Images'

Your tasks

Project goal: automatically detect Nuclei centres in histology images.

- Read and understand the code provided in `image_processing_tutorial.py`.
- Complete functions from by modelling your solutions after the starter code.
 - `project_helpers.py`,
 - `nuclei_detection_tutorial.py`, and
 - `project.py`

Data path

- The starter code assumes that the data directory (crchristophenotypes_2016_04_28) will be in the same directory as the .py files.
- If that is not the right location, you must set the data_path variable to the right directory.

Viewing plots

- In the starter code, there is a constant named VIEW.
- When VIEW is set to False, no images are shown.
- When VIEW is set to True, the images are displayed using show.
- You may change the value of VIEW as you develop your code.
- Example code snippet from starter code:

```
if VIEW:
```

```
    pyplot.show()
```

tuple

- Python has a type tuple, which is used to store ordered collections of data.
- Like lists, tuples can be indexed.
- Unlike lists, tuples are immutable.

Example:

```
>>> t = (1, 2, 3, 4)
```

```
>>> len(t)
```

```
4
```

```
>>> t[1]
```

```
2
```


Numpy's vstack

- Take arrays and stack them vertically to produce a single array.
- Example:

```
>>> a1 = np.array([1, 2, 3])
```

```
>>> a2 = np.array([4, 5, 6])
```

```
>>> result = np.vstack((a1, a2)) # note: two sets of parentheses; the argument is  
the tuple (a1, a2)
```

```
>>> result
```

```
array([[1, 2, 3],  
       [4, 5, 6]])
```

Numpy's dstack

- Take arrays and stack them depthwise to produce a single 3D array.
- Example:

```
>>> a1 = np.array([1, 2, 3])
```

```
>>> a2 = np.array([4, 5, 6])
```

```
>>> result = np.dstack((a1, a2)) # note: two sets of parentheses; the argument is  
the tuple (a1, a2)
```

```
>>> result
```

```
array([[[1, 4],  
        [2, 5],  
        [3, 6]]])
```