

# Computing for Medicine: Phase 3, Seminar 4 Project

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# Seminar 4 Project

- The project handout is posted:
  - <http://c4m.cdf.toronto.edu/cohort1/phase3/>
- Two approaches for doing your work:
  - Use the Computer Science Teaching Labs computing network.
  - Use your personal computer.

# Software installation

- New required packages:
  - skimage (<http://scikit-image.org/docs/dev/api/skimage.html>)
  - joblib (<https://pythonhosted.org/joblib/>)
- Required packages that were also used by the previous projects:
  - numpy, pylab, scipy, matplotlib, sklearn

# OVERVIEW

# 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’, IEEE Transactions on Medical Imaging, 2016. (in press)

# 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
  - `project_helpers.py`,
  - `nuclei_detection_tutorial.py`, and
  - `project.py`by modeling your solutions after the provided tutorials and starter code.

# 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 the data is in a different 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 `matplotlib`'s `show` function.
- You may change the value of `VIEW` as you develop your code.
- Example code snippet from starter code:

```
if VIEW:  
    pyplot.show()
```

# PYTHON TOOLS

# 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 depth wise 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]]])
```

# UPCOMING SEMINARS

# Seminar 5: Dr. Periklis Andritsos

- Tuesday December 13 2016 6-8pm
- Location: DCS Innovation Lab
- Topic: Database Systems
- <https://www.ischool.utoronto.ca/faculty/periklis-andritsos>

**FEEDBACK**

# Phase 3, Seminar 4 Survey

- You will receive an email with the subject “C4M: Phase 3, Seminar 4 Feedback Survey”.
- Please complete that survey now:
  - <https://www.surveymonkey.com/r/C4MSeminar4>