



Writing readable code

Robert Haase



Readable code



Recommendations for readable code

- Every command belongs on its own line
- Insert empty lines to separate important processing steps.
- Put comments to introduce sections
- Print out intermediate results
- Put <u>spaces</u> between operators and operands, because:

This is easier to read thanthat, orisnt'it?

- Keep code blocks it short.
- Consider writing code out instead of keeping it short.

```
# initialize program
                                                  ang")
     b=3
     c=8
     # compute result before we can evaluate it
     d=(a+b)/c
[3]: d
[3]: 1.0
[4]: d = (a + b) / c
[4]: 1.0
[5]: print("Yin" if d==5 else "Yang")
     Yang
[6]: if d == 5:
         print("Yin")
     else:
         print("Yang")
```

Yang

Naming conventions



- Write code so that others can read it. Your future self will thank you.
- Name variables and functions after English words.

```
A = w * h
print(A)

nNucl = cntNcl(img)

number_of_nuclei = count_nuclei(image)
numberOfNuclei = countNuclei(image)

volume = travelled_distance / time

percentage = "Robert"

Snake-case

speed = travelled_distance / time
```

- Name variables using nouns
- Name functions as verb + noun

Don't repeat yourself (DRY)



• Use <u>for-loops</u> to prevent code duplication. It also makes code maintenance easier.

```
image = cle.imread("../../data/BBBC007 batch/17P1 POS0013 D 1UL.tif")
      labels = cle.voronoi otsu labeling(image, spot sigma=3)
      number of nuclei = labels.max()
      number_of_nuclei
[8]: 44.0
     image = cle.imread("../../data/BBBC007_batch/20P1_POS0005_D_1UL.tif")
      labels = cle.voronoi otsu labeling(image, spot sigma=3)
      number of nuclei = labels.max()
      number_of_nuclei
[9]: 41.0
     image = cle.imread("../../data/BBBC007 batch/20P1 POS0007 D 1UL.tif")
      labels = cle.voronoi otsu labeling(image, spot sigma=3)
      number of nuclei = labels.max()
      number of nuclei
[10]: 73.0
```

```
[11]: folder = "../../data/BBBC007_batch/"
      files = ["17P1 POS0013 D 1UL.tif",
              "20P1 POS0005 D 1UL.tif",
              "20P1 POS0007 D 1UL.tif"]
[12]: for file in files:
          image = cle.imread(folder + file)
          labels = cle.voronoi_otsu_labeling(
                          image,
                          spot sigma=3)
          number of nuclei = labels.max()
          print(file, number of nuclei)
      17P1_POS0013_D_1UL.tif 44.0
      20P1 POS0005 D 1UL.tif 41.0
      20P1 POS0007 D 1UL.tif 73.0
```

Don't repeat yourself (DRY)



• Use functions to prevent code duplication. It also makes code more flexible.

Default parameter

```
image = cle.imread("../../data/BBBC007 batch/17P1 POS0013 D 1UL.tif")
                                                                                [13]: def count nuclei(image, spot sigma=3):
      labels = cle.voronoi otsu labeling(image, spot sigma=3)
                                                                                           labels = cle.voronoi otsu labeling(
      number of nuclei = labels.max()
                                                                                                           image,
      number_of_nuclei
                                                                                                           spot_sigma=spot_sigma)
                                                                                           number of nuclei = labels.max()
 [8]: 44.0
                                                                                           return number of nuclei
     image = cle.imread("../../data/BBBC007_batch/20P1_POS0005_D_1UL.tif")
      labels = cle.voronoi otsu labeling(image, spot sigma=3)
                                                                                [14]: count_nuclei(cle.imread(folder + files[0]))
      number of nuclei = labels.max()
      number_of_nuclei
                                                                                [14]: 44.0
[9]: 41.0
                                                                                      count nuclei(cle.imread(folder + files[1]))
     image = cle.imread("../../data/BBBC007 batch/20P1 POS0007 D 1UL.tif")
                                                                                [15]: 41.0
      labels = cle.voronoi otsu labeling(image, spot sigma=3)
      number of nuclei = labels.max()
                                                                                 [16]: count nuclei(cle.imread(folder + files[2]))
      number of nuclei
                                                                                [16]: 73.0
[10]: 73.0
                                                                   [18]: count nuclei(cle.imread(folder + files[2]), spot sigma=5)
                                                                   [18]: 68.0
```

Prevent magic numbers



- Put all parameters of your workflow on top so that one can easily spot them.
- Give them reasonable names

```
[4]: # enter the image filename to be processed here
                                                            file to process = "../../data/BBBC007_batch/17P1_POS0013_D_1UL.tif"
                                                            # enter the expected radius of nuclei here, in pixel units
                                                            approximate nuclei radius = 3
[3]: image = imread("../../data/BBBC007 batch/17P1 POS [5]:
                                                            image = imread(file to process)
                                                            # noise removal
     # noise removal
     blurred = gaussian(image, )
                                                            blurred = gaussian(image, approximate nuclei radius)
                                                            # instance segmentation
     # instance segmentation
     binary = blurred > threshold otsu(blurred)
                                                            binary = blurred > threshold_otsu(blurred)
     labels = label(binary)
                                                            labels = label(binary)
                                                            # quantitative measurement
     # quantitative measurement
     labels.max()
                                                            labels.max()
                                                       [5]: 37
```

Divide and rule



Prevent long, complicated macros ("spaghetti code")

```
[2]: image = imread("../../data/blobs.tif")
    footprint = disk(15)
    background_subtracted = white_tophat(image,
                                          footprint=footprint)
    particle radius = 5
    denoised = gaussian(background subtracted,
                         sigma=particle radius)
    binary = denoised > threshold_otsu(denoised)
    labels = label(binary)
    requested_measurements = ["label", "area", "mean_intensity"]
    regionprops = regionprops table(image,
                                     labels.
                                     properties=requested measurements)
    table = pd.DataFrame(regionprops)
    mean total intensity = np.mean(table["area"] * table["mean intensity"])
    mean total intensity
```

At least: Write sections of code with headlines

```
[3]: # configuration
     file to analyze = "../../data/blobs.tif"
     background subtraction radius = 15
     particle radius = 5
     requested measurements = ["area", "mean intensity"]
     # Load data
     image = imread(file to analyze)
     # preprocess image
     footprint = disk(background subtraction radius)
     background_subtracted = white_tophat(image,
                                          footprint=footprint)
     denoised = gaussian(background subtracted,
                         sigma=particle radius)
     # segment image
     binary = denoised > threshold otsu(denoised)
     labels = label(binary)
     # extract features
     regionprops = regionprops table(image,
                                     labels,
                                     properties=requested measurements)
     table = pd.DataFrame(regionprops)
     # descriptive statistics
     mean total intensity = np.mean(table["area"] * table["mean intensity"])
     mean total intensity
```



Prevent long, complicated macros ("spaghetti code")

```
[4]: # reusable functions
    def preprocess_image(image, background_subtraction_radius, particle_radius):
         """Apply background removal and denoising"""
         footprint = disk(background_subtraction_radius)
         background_subtracted = white_tophat(image, footprint=footprint)
         denoised = gaussian(background_subtracted, sigma=particle_radius)
         return denoised
     def segment_image(image):
         """Apply thresholding and connected component analysis"""
         binary = image > threshold otsu(image)
         labels = label(binary)
         return labels
```

- At least: Write sections of code with headlines
- Even better: Write custom functions



• Divide complex issues into smaller, accessible issues like

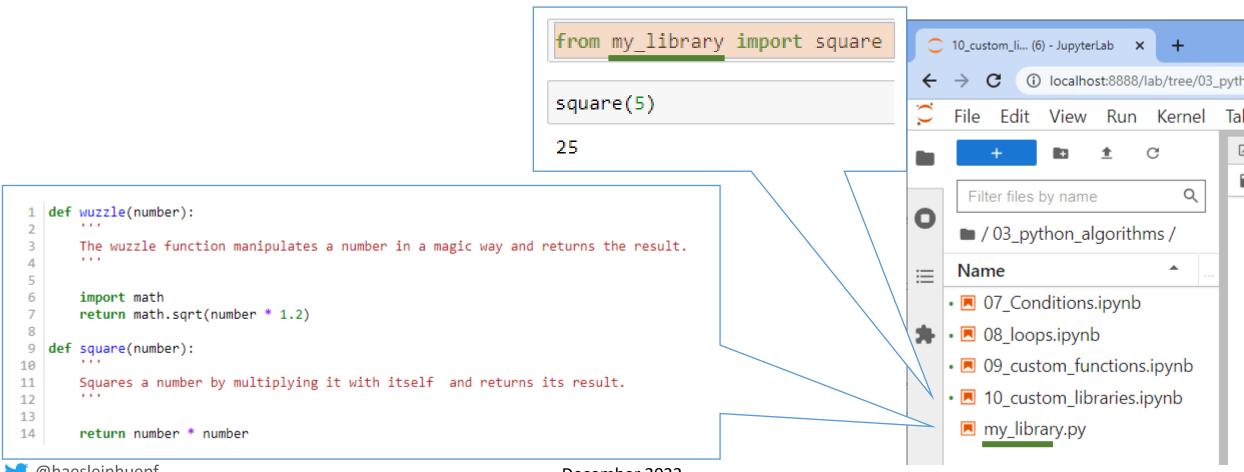
```
def analyse average total intensity(filename,
                                    background subtraction radius = 15,
                                    particle radius = 5):
    """Load an image, segment objects and measure their mean total intensity."""
    image = imread(filename)
    denoised = preprocess image(image,
                                background subtraction radius,
                                particle radius)
    labels = segment_image(denoised)
    requested_measurements = ["area", "mean_intensity"]
    table = extract_features(image,
                             labels,
                             requested measurements)
   # descriptive statistics
   mean total intensity = np.mean(table["area"] * table["mean intensity"])
    return mean total intensity
```

- It's easier to get an overview about small functions
- It forces programmers to organise their macros

Reusing code from custom modules



- For re-using functions between notebooks / projects, use libraries. -> Sustainability
- Simple python libraries are .py files containing multiple functions.
- The import statement allows you to import python files from the same folder.



Keep it short and simple(KISS)



What does this program do?

```
image = imread("../../data/blobs.tif")

# define a list of functions and a corresponding list of arguments
functions = [gaussian_blur, threshold_otsu, label]
argument_lists = [[.5], [], []]

# go through functions and argument lists pair-wise
for function, argument_list in zip(functions, argument_lists):
    # execute function with given arguments
    image = function(image, *argument_list)

result1 = image
imshow(result1)
```

- There are two ways design software:
 - make it <u>so simple</u> that there are <u>obviously no issues</u> or
 - make it <u>so complicated</u> that there are <u>no obvious issues</u>.

(based loosly on Tony Hoare, "The Emperor's Old Clothes," CACM Feb. 1981)

• Take home: Keep it short and simple!



```
image = imread("../../data/blobs.tif")
blurred = gaussian_blur(image, 5)
binary = threshold_otsu(blurred)
labels = label(binary)

result2 = labels
imshow(result2)
```







Exercise: modularization



Go through the code from yesterday and identify a workflow that worked well and contained these steps:

- Image preprocessing
- Image segmentation

Copy & paste code from the notebooks you worked with yesterday into a new custom module my_analysis.py. Write a new notebook that demonstrates how to use the functions of your my_analysis module.

```
def my_workflow(input_image, ...):
     ...
     return label_image
```

