

Workshop 3: Functions and modules

FIE463: Numerical Methods in Macroeconomics and Finance using Python

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February 5, 2026

See GitHub repository for notebooks and data:

<https://github.com/richardfoltyn/FIE463-V26>

Exercise 1: Standard deviation of a sequence of numbers

The standard deviation σ characterizes the dispersion of a sequence of data (x_1, x_2, \dots, x_N) around its mean \bar{x} . It is computed as the square root of the variance σ^2 , defined as

$$\sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \bar{x})^2$$

where N is the number of elements (we ignore the degrees-of-freedom correction), and the mean \bar{x} is defined as

$$\bar{x} = \frac{1}{N} \sum_{i=1}^N x_i$$

The above formula for the variance can be rewritten as

$$\sigma^2 = \left(\frac{1}{N} \sum_{i=1}^N x_i^2 \right) - \bar{x}^2$$

This suggests the following algorithm to compute the standard deviation:

1. Compute the mean $\bar{x} = \frac{1}{N} \sum_{i=1}^N x_i$.
2. Compute the mean of squares $S = \frac{1}{N} \sum_{i=1}^N x_i^2$.
3. Compute the variance $\sigma^2 = S - \bar{x}^2$.
4. Compute the standard deviation $\sigma = \sqrt{\sigma^2}$.

In this exercise, you are asked to implement the above algorithm and compare your function with NumPy's implementation `np.std()`.

1. Create a module `my_stats.py` and add the function

```
def my_std(x):  
    """  
    Compute and return the standard deviation of the sequence x.  
    """
```

which implements the above algorithm to compute the standard deviation of a given sequence x (this could be a tuple, list, array, etc.). Your implementation should *only use built-in functions* such as `len()`, `sum()`, and `sqrt()` from the `math` module.

2. Import this function into the Jupyter notebook. Using an array of 11 elements that are uniformly spaced on the interval $[0, 10]$, confirm that your function returns the same value as `np.std()`.
3. Benchmark your implementation against `np.std()` for three different arrays with 11, 101, and 10001 elements that are uniformly spaced on the interval $[0, 10]$.

Hint: Use the cell magic `%timeit` to time the execution of a statement.

You should add the following cell magic so that the contents of `my_stat.py` are automatically reloaded whenever you change the file:

```
[1]: %load_ext autoreload
      %autoreload 2
```

Solution.

Part 1 — Implement `my_std`

You should implement `my_std()` in the separate file `my_stats.py`, but we will implement it directly in the notebook to keep the solution within one file.

```
[2]: from math import sqrt

def my_std(x):
    """
    Compute standard deviation of x using the built-in functions sum()
    and len().

    Parameters
    -----
    x: Sequence of numbers

    Returns
    -----
    sd : float
        Standard deviation of x.
    """

    # Number of observations
    N = len(x)

    # Compute mean
    mean = sum(x) / N

    # Compute the mean of squares
    S = sum(xi**2.0 for xi in x) / N

    # Compute variance
    var = S - mean**2.0

    # Compute standard deviation
    sd = sqrt(var)

    return sd
```

Part 2 — Compare with NumPy std()

```
[3]: import numpy as np

# Create the test data
data = np.linspace(0.0, 10.0, 11)

[4]: # Uncomment this to import my_std from the separate module
# from my_stats import my_std

# Call your own implementation
my_std(data)

[4]: 3.1622776601683795

[5]: # Call NumPy's implementation
np.std(data)

[5]: np.float64(3.1622776601683795)
```

As you can see, both implementations return the same value.

Part 3 — Benchmarks

We now compare the runtime for our own vs. NumPy's implementation for increasing sample sizes. For simplicity, we can directly embed the `%timeit` magic into the loop iterating over sample sizes. This is, strictly speaking, not valid Python syntax and only works in Jupyter notebooks. It should be avoided in real applications.

```
[6]: # Sample sizes to benchmark
N = [11, 101, 10001]

# Benchmark our own implementation for various sample sizes
for n in N:
    # Create test data of given size
    data = np.linspace(0.0, 10.0, n)
    print(f'Running own implementation for N={n}')
    # Time the execution
    %timeit my_std(data)
```

Running own implementation for N=11

3.12 μ s \pm 7.37 ns per loop (mean \pm std. dev. of 7 runs, 100,000 loops each)

Running own implementation for N=101

19.1 μ s \pm 172 ns per loop (mean \pm std. dev. of 7 runs, 100,000 loops each)

Running own implementation for N=10001

1.77 ms \pm 13.1 μ s per loop (mean \pm std. dev. of 7 runs, 1,000 loops each)

```
[7]: # Benchmark NumPy implementation for various sample sizes
for n in N:
    # Create test data of given size
    data = np.linspace(0.0, 10.0, n)
    print(f'Running NumPy implementation for N={n}')
    # Time the execution
    %timeit np.std(data)
```

Running NumPy implementation for N=11

9.26 μ s \pm 890 ns per loop (mean \pm std. dev. of 7 runs, 100,000 loops each)

Running NumPy implementation for N=101

8.69 μ s \pm 86.5 ns per loop (mean \pm std. dev. of 7 runs, 100,000 loops each)
Running NumPy implementation for N=10001

16.2 μ s \pm 535 ns per loop (mean \pm std. dev. of 7 runs, 100,000 loops each)

As you can see, our implementation is fast for small arrays but substantially slower for arrays of 10,000 elements (by a factor of around 100, depending on hardware and software).

Exercise 2: Locating maximum values

In this exercise, you are asked to write a function that returns the position of the largest element from a given sequence (list, tuple, array, etc.).

1. Write a function `my_argmax()` that takes as an argument a sequence and returns the (first) index where the maximum value is located. Only use built-in functionality in your implementation (no NumPy).
2. Create an array with 101 values constructed using the sine function,

```
arr = np.sin(np.linspace(0.0, np.pi, 101))
```

and use it to test your function.
3. Compare the result returned by your function to NumPy's implementation `np.argmax()`.

Solution.

Part 1 — Implement `my_argmax()`

```
[8]: def my_argmax(x):  
    """  
    Return the location of the (first) maximum element.  
  
    Parameters  
    -----  
    x : array-like  
        A list or array of numerical values.  
  
    Returns  
    -----  
    int  
        The index of the first occurrence of the maximum value in x.  
    """  
  
    # Initially, the maximum element must be the first one.  
    imax = 0  
    xmax = x[0]  
  
    # Iterate through REMAINING elements to see if there is a larger one  
    for i, xi in enumerate(x[1:]):  
        if xi > xmax:  
            # Update the location of the maximum if a larger value is found  
            imax = i  
            xmax = xi  
  
    return imax
```

Part 2 — Test implementation

```
[9]: import numpy as np

# Construct test array
arr = np.sin(np.linspace(0.0, np.pi, 101))

[10]: # Find maximum location and print the result
i = my_argmax(arr)
print(f'The maximum is located at index {i} with value = {arr[i]}')

The maximum is located at index 49 with value = 0.9995065603657316
```

Part 3 — Compare with NumPy argmax()

```
[11]: # Find location using NumPy's argmax()
j = np.argmax(arr)
print(f'The maximum is located at index {j} with value = {arr[j]}')

The maximum is located at index 50 with value = 1.0
```

Exercise 3: Two-period consumption-savings problem

This exercise asks you to find the utility-maximizing consumption levels using grid search, an algorithm that evaluates all possible alternatives from a given set (the “grid”) to locate the maximum.

Consider the following standard consumption-savings problem over two periods with lifetime utility $U(c_1, c_2)$ given by

$$\begin{aligned} \max_{c_1, c_2} \quad & U(c_1, c_2) = u(c_1) + \beta u(c_2) \\ \text{s.t.} \quad & c_1 + \frac{c_2}{1+r} = w \\ & c_1 \geq 0, c_2 \geq 0 \end{aligned}$$

where β is the discount factor, r is the interest rate, w is initial wealth, and (c_1, c_2) is the optimal consumption allocation to be determined. The second line is the budget constraint which ensures that the chosen consumption bundle (c_1, c_2) is feasible. The per-period CRRA utility function $u(c)$ is given by

$$u(c) = \begin{cases} \frac{c^{1-\gamma}}{1-\gamma} & \text{if } \gamma \neq 1 \\ \log(c) & \text{if } \gamma = 1 \end{cases}$$

where γ is the coefficient of relative risk aversion (RRA) and $\log(\bullet)$ denotes the natural logarithm.

1. Write a function `util(c, gamma)` which evaluates the per-period utility $u(c)$ for a given consumption level c and the parameter γ . Make sure to take into account the log case!

Hint: You can use the `np.log()` function from NumPy to compute the natural logarithm.

2. Write a function `util_life(c_1, c_2, beta, gamma)` which uses `util()` from above to compute the lifetime utility $U(c_1, c_2)$ for given consumption levels (c_1, c_2) and parameters.
3. Assume that $r = 0.04$, $\beta = 0.96$, $\gamma = 1$, and $w = 1$.

- Create a candidate array (grid) of period-1 consumption levels with 100 grid points that are uniformly spaced on the interval $[\epsilon, w - \epsilon]$ where $\epsilon = 10^{-5}$.

Note that we enforce a minimum consumption level ϵ , as zero consumption yields $-\infty$ utility for the given preferences, which can never be optimal.

- Compute the implied array of period-2 consumption levels from the budget constraint.
 - Given these candidate consumption levels, use the function `util_life()` you wrote earlier to evaluate lifetime utility for each bundle of consumption levels (c_1, c_2) .
4. Use the function `np.argmax()` to locate the index at which lifetime utility is maximized. Print the maximizing consumption levels (c_1, c_2) as well as the associated maximized utility level.

Solution.

Part 1 — Implement util function

```
[12]: import numpy as np

def util(c, gamma):
    """
    Return per-period utility for a given consumption level c.
    """
    if gamma == 1:
        # Utility for log preferences
        u = np.log(c)
    else:
        # Utility for general CRRA preferences
        u = c**(1.0 - gamma) / (1.0 - gamma)

    return u
```

Part 2 — Compute lifetime utility

```
[13]: def util_life(c1, c2, beta, gamma):
    """
    Return lifetime utility for given consumption levels.
    """

    # Utility in period 1
    u1 = util(c1, gamma)

    # Utility in period 2
    u2 = util(c2, gamma)

    # Lifetime utility
    U = u1 + beta * u2

    return U
```

Part 3 — Set parameters and evaluate utility

```
[14]: # Parameters
r = 0.04
beta = 0.96
gamma = 1.0

# Initial wealth
wealth = 1.0
```

We can now create the candidate grid for period-1 consumption. Period-2 consumption then follows from the budget constraint. The candidate grid is created on the interval $[\epsilon, w - \epsilon]$ for a small value of epsilon. The reason for this is twofold:

1. With CRRA preferences, zero consumption yields $-\infty$ utility, which can never be optimal.
2. Moreover, trying to evaluate `np.log(0)` generates warnings which we wish to avoid.

We avoid these complications by creating a grid of candidate consumption levels on the interval $[\epsilon, w - \epsilon]$ instead of $[0, w]$.

```
[15]: # Grid size
      N = 100

      # Minimum consumption level
      epsilon = 1.0e-5

      # Candidate grid for period-1 consumption
      c1_grid = np.linspace(epsilon, wealth - epsilon, N)

      # Candidate grid for period-2 consumption (from budget constraint)
      c2_grid = (1+r) * (wealth - c1_grid)
```

With the consumption grids at hand, we can evaluate the lifetime utility for each alternative.

```
[16]: # Evaluate lifetime utility for each (c1, c2)
      u_grid = util_life(c1_grid, c2_grid, beta, gamma)
```

Part 4 — Find optimal consumption

```
[17]: # Find maximum
      imax = np.argmax(u_grid)

      # Recover the maximizing consumption levels and utility
      c1_max = c1_grid[imax]
      c2_max = c2_grid[imax]
      u_max = u_grid[imax]

      # Report consumption levels and utility at maximum
      print(f'Utility is maximized at c1={c1_max:.3f}, c2={c2_max:.3f} with u={u_max:.5e}')
```

```
Utility is maximized at c1=0.515, c2=0.504 with u=-1.32060e+00
```
