

Lab 3: Debugging and profiling code

Components

- TA demonstration
 - The TA will show his/her debugging and profiling setup in VS Code.
- 1. Debug code
 - Specifically, we'll have the debugger stop inside someone else's code!
- 2. Profiling code
 - Walk through a short demo
 - Students will then profile code themselves
 - Show a few alternative (hacky) methods

Learning objectives

- Understand that debuggers and profilers exist and make your life easier
- How to use a debugger within a Jupyter notebook
- How to profile code

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Intro

As our example today, we'll look at code for computing the matrix exponential, e^A . You should remember this from taking matrix methods APPM 3310. If A is diagonalizable, then we can compute e^A using eigenvalues; if A is not diagonalizable, it's a bit more complicated. Regardless, computing e^A *efficiently* and *stably* is somewhat challenging and is a [classic problem](#) in numerical analysis.

We'll use the standard `scipy.linalg` package to do this for us, and demonstrate how we can debug and profile code, even when we didn't write the code ourselves!

This lab was tested with SciPy versions 1.11.1 and 1.16.1. We *hope* it works with most other recent versions!

Note: we recommend you download this entire `.ipynb` notebook and run it locally on your computer. You can download it via navigating our github site, or directly from <https://raw.githubusercontent.com/cu-applied-math/appm-4600-numerics/refs/heads/>

```
In [1]: import scipy  
print(scipy.__version__)
```

1.17.0

1. Debugging

Note: we **highly** suggest running this in an editor like VS Code, or at least Jupyter Lab (not a plain Jupyter Notebook). You can debug in colab, but it's actually harder because you have to use some command line debugging skills. See this article [Debugging in Google Colab](#) if you're curious (it uses the `ipdb` package).

Note: If using a local editor, like VSCode, there may be a setting that by default does **not** let you debug code from libraries/packages. To make sure you can debug the scipy code, do the following:

1. Open the VSCode settings (one way to do this is to click on the gear icon in the lower left hand corner).
2. In the settings search bar, search "debug just my code".
3. This should return a setting/checkbox tilted "Jupyter: Debug Just My Code". By default, this is checked/enabled. **Uncheck/disable** this setting.

Now you should be able to debug the scipy code.

There may be issues in Jupyter Lab with debugging external code as well. If you really cannot get it to work, let the TA know, and instead write your own code and step through it with the debugger.

Instructions

Run the following code using your IDE, like VS Code. Inside the `expm` function, there is a line of code `m, s = pick_pade_structure(Am)`.

Deliverable

For the matrix A below, when it is passed into `expm`, what is the value of `m` from the `m, s = pick_pade_structure(Am)` line inside the Scipy code?

```
In [3]: import scipy.linalg as sla
import numpy as np

A = np.arange(int(1e2)).reshape(10,10)/100
print(A)
eA = sla.expm(A)

[[0.    0.01 0.02 0.03 0.04 0.05 0.06 0.07 0.08 0.09]
 [0.1   0.11 0.12 0.13 0.14 0.15 0.16 0.17 0.18 0.19]
 [0.2   0.21 0.22 0.23 0.24 0.25 0.26 0.27 0.28 0.29]
 [0.3   0.31 0.32 0.33 0.34 0.35 0.36 0.37 0.38 0.39]
 [0.4   0.41 0.42 0.43 0.44 0.45 0.46 0.47 0.48 0.49]
 [0.5   0.51 0.52 0.53 0.54 0.55 0.56 0.57 0.58 0.59]
 [0.6   0.61 0.62 0.63 0.64 0.65 0.66 0.67 0.68 0.69]
 [0.7   0.71 0.72 0.73 0.74 0.75 0.76 0.77 0.78 0.79]
 [0.8   0.81 0.82 0.83 0.84 0.85 0.86 0.87 0.88 0.89]
 [0.9   0.91 0.92 0.93 0.94 0.95 0.96 0.97 0.98 0.99]]
```

From debugging, $m = 13$ for this program

2. Profiling Code

Profiling code refers to looking at running code and noticing what parts of the code take a long time, as well as memory usage. For memory, you might look at what parts of the code require a lot of memory, or where memory leaks are, or at the cache miss rate, etc.

We're not going to focus on memory profiling, but instead on time (aka speed) profiling. A profiler tells us which parts of the code took a long time. Some profiling tells us which **functions** take a while, other types of profiling tell us which **lines** of code take a while. Today we'll do the latter.

There are some caveats: to do profiling, the profiler adds some extra overhead, so everything runs a bit slower, and you don't get a 100% accurate picture of exactly how long each part of the code takes. Usually it's accurate enough to give you an idea of where the slowish parts are.

Some programmers avoid formal profiling but just adding timing statements into their code (see section 2c for a few ways to do this). Sometimes this is easy to do and sufficient, but other times it's not systematic enough and takes more effort than you need; it's also a problem if you want to profile some library code where you don't want to have to edit the files.

2a. Example usage

We're going to use the `line_profiler` package, so let's first check if it is installed (and if not, we'll install it):

```
In [5]: try:
```

```

import line_profiler
print(f"The package 'line_profiler' is installed.")
except ModuleNotFoundError:
    print(f"The package 'line_profiler' is NOT installed.")
    # To install it via PIP, we can do the following:
    import subprocess
    subprocess.check_call(['pip', 'install', 'line_profiler'])
    # or, just run: !pip install line_profiler      from within jupyter
    # or, just run:  pip install line_profiler      from a command line
    # or, if using conda, run:  conda install conda-forge::line_profiler      from a command line

```

The package 'line_profiler' is installed.

Now let's use it. There are different ways to use it. We'll use it in one manner which works well with jupyter notebooks. The following is an ipython/jupyter specific command (you only need to run this once per session):

```
In [6]: %load_ext line_profiler
```

Now let's have some code to profile. Below is some silly code that does the computation

$$f(n) = \sum_{k=1}^n k$$

in a few different ways.

```

In [22]: def sum_python(n):
          sum = 0.
          for k in range(n+1):
              sum += k
          return sum

def sum_numpy(n):
    nList = np.arange(n+1)
    sum = np.sum(nList)
    return sum

def my_sum(n):
    """ returns the sum of k from k = 1 ... n """
    s1 = sum_python(n)
    s2 = sum_numpy(n)
    s3 = n*(n+1)/2 # use the closed-form formula, cf. https://en.wikipedia.org/wiki/1_%2B_2_%2B_3_%2B_4_%2B_%E2%8B%AF
    assert np.isclose(s1,s2), "The first two methods gave different values :-( )"
    assert np.isclose(s2,s3), "The second two methods gave different values :-( )"

    return s1

```

```
my_sum(1000)
```

```
Out[22]: 500500.0
```

Now let's see how to use the line profiler, which will tell us which lines are taking up most of the time.

Note: this works better in an IDE like VS Code. It *kinda* works in Colab (it may only tell you the line number)

We'll use it in the following form: `%lprun -f name_of_fcn big_fcn(...)` where `big_fcn(...)` is the code that you are going to call, and `name_of_fcn` is the *part* of that code that you want to investigate line-by-line.

For example:

```
In [18]: %lprun -f my_sum my_sum(1000)
```

```
Timer unit: 1e-09 s
```

```
Total time: 0.000917 s
```

```
File: /var/folders/40/26f5s8qj42z2mxw1gfrnh5rm0000gn/T/ipykernel_50763/4186063441.py
```

```
Function: my_sum at line 12
```

Line #	Hits	Time	Per Hit	% Time	Line Contents
12					def my_sum(n):
13					""" returns the sum of k from k = 1 ... n """
14	1	657000.0	657000.0	71.6	s1 = sum_python(n)
15	1	70000.0	70000.0	7.6	s2 = sum_numpy(n)
16	1	0.0	0.0	0.0	s3 = n*(n+1)/2 # use the closed-form formula, cf. https://en.wikipedia.org/wiki/1_2B_2_2B_3_2B_4_2B_5_2B_6_2B_7_2B_8_2B_9_2B_10_2B_11_2B_12_2B_13_2B_14_2B_15_2B_16_2B_17_2B_18_2B_19_2B_20_2B_21_2B_22_2B_23_2B_24_2B_25_2B_26_2B_27_2B_28_2B_29_2B_30_2B_31_2B_32_2B_33_2B_34_2B_35_2B_36_2B_37_2B_38_2B_39_2B_40_2B_41_2B_42_2B_43_2B_44_2B_45_2B_46_2B_47_2B_48_2B_49_2B_50_2B_51_2B_52_2B_53_2B_54_2B_55_2B_56_2B_57_2B_58_2B_59_2B_60_2B_61_2B_62_2B_63_2B_64_2B_65_2B_66_2B_67_2B_68_2B_69_2B_70_2B_71_2B_72_2B_73_2B_74_2B_75_2B_76_2B_77_2B_78_2B_79_2B_80_2B_81_2B_82_2B_83_2B_84_2B_85_2B_86_2B_87_2B_88_2B_89_2B_90_2B_91_2B_92_2B_93_2B_94_2B_95_2B_96_2B_97_2B_98_2B_99_2B_100
17	1	122000.0	122000.0	13.3	assert np.isclose(s1,s2), "The first two methods gave different values :-()"
18	1	67000.0	67000.0	7.3	assert np.isclose(s2,s3), "The second two methods gave different values :-()"
19					
20	1	1000.0	1000.0	0.1	return s1

The output of that tells us that `sum_python` takes about 50% of the total time (this varies... run the above cell a few times to get an idea of the average), and `sum_numpy` takes about 10% of the time, and the rest of the time is spent on the `assert` statements.

So let's take a deep dive on the `sum_python` code:

```
In [15]: %lprun -f sum_python my_sum(1000)
```

Timer unit: 1e-09 s

Total time: 0.000937 s

File: /var/folders/40/26f5s8qj42z2mxw1gfrnh5rm0000gn/T/ipykernel_50763/4186063441.py

Function: sum_python at line 1

Line #	Hits	Time	Per Hit	% Time	Line Contents
=====					
1					def sum_python(n):
2	1	2000.0	2000.0	0.2	sum = 0.
3	1002	496000.0	495.0	52.9	for k in range(n+1):
4	1001	437000.0	436.6	46.6	sum += k
5	1	2000.0	2000.0	0.2	return sum

The output of this tells is that the line `for k in range(n+1):` takes 40% of the time, and the `sum += k` takes another 60%. This means that the actual loop itself is slow, not the computation in the loop. That's not uncommon for interpreted languages like Matlab and Python (hence the emphasize on "vectorized" code).

We can also take a deep dive on the `sum_numpy` code:

```
In [16]: %lprun -f sum_numpy my_sum(1000)
```

Timer unit: 1e-09 s

Total time: 0.000131 s

File: /var/folders/40/26f5s8qj42z2mxw1gfrnh5rm0000gn/T/ipykernel_50763/4186063441.py

Function: sum_numpy at line 7

Line #	Hits	Time	Per Hit	% Time	Line Contents
=====					
7					def sum_numpy(n):
8	1	34000.0	34000.0	26.0	nList = np.arange(n+1)
9	1	96000.0	96000.0	73.3	sum = np.sum(nList)
10	1	1000.0	1000.0	0.8	return sum

2b: Your task

Deliverable

Run the code below, and find the **3 most expensive lines of code** from within the `scipy.linalg.expm` function

```
In [23]: rng = np.random.default_rng(123456)
A = rng.standard_normal( (3000,3000) )/1e2
eA = sla.expm(A)

%lprun -f sla.expm sla.expm(A)
```

Timer unit: 1e-09 s

Total time: 1.39894 s

File: /Library/Frameworks/Python.framework/Versions/3.14/lib/python3.14/site-packages/scipy/linalg/_matfuncs.py

Function: expm at line 234

Line #	Hits	Time	Per Hit	% Time	Line Contents
234					def expm(A):
235					"""Compute the matrix exponential of an array.
236					
237					Array argument(s) of this function may have additional
238					"batch" dimensions prepended to the core shape. In this case, the
array is treated					
239					as a batch of lower-dimensional slices; see :ref:`linalg_batch` fo
r details.					
240					
241					Parameters
242					-----
243					A : ndarray
244					Input with last two dimensions are square ``(..., n, n)``.
245					
246					Returns
247					-----
248					eA : ndarray
249					The resulting matrix exponential with the same shape of ``A``
250					
251					Notes
252					-----
253					Implements the algorithm given in [1]_, which is essentially a Pad
e					
254					approximation with a variable order that is decided based on the a
rarray					
255					data.
256					
257					For input with size ``n``, the memory usage is in the worst case i
n the					
258					order of ``8*(n**2)``. If the input data is not of single and doub
le					
259					precision of real and complex dtypes, it is copied to a new array.
260					
261					For cases ``n >= 400``, the exact 1-norm computation cost, breaks
even with					
262					1-norm estimation and from that point on the estimation scheme giv
en in					
263					[2]_ is used to decide on the approximation order.
264					
265					References

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1	11000.0	11000.0	0.0
1	5000.0	5000.0	0.0
1	4000.0	4000.0	0.0

```
-----  
.. [1] Awad H. Al-Mohy and Nicholas J. Higham, (2009), "A New Scal  
and Squaring Algorithm for the Matrix Exponential", SIAM J.  
Anal. Appl. 31(3):970-989, :doi:`10.1137/09074721X`  
.. [2] Nicholas J. Higham and Francoise Tisseur (2000), "A Block A  
for Matrix 1-Norm Estimation, with an Application to 1-Norm  
Pseudospectra." SIAM J. Matrix Anal. Appl. 21(4):1185-1201,  
:doi:`10.1137/S0895479899356080`  
  
Examples  
-----  
>>> import numpy as np  
>>> from scipy.linalg import expm, sinm, cosm  
  
Matrix version of the formula  $\exp(0) = 1$ :  
  
>>> expm(np.zeros((3, 2, 2)))  
array([[[1., 0.],  
        [0., 1.]],  
<BLANKLINE>  
        [[1., 0.],  
        [0., 1.]],  
<BLANKLINE>  
        [[1., 0.],  
        [0., 1.]])  
  
Euler's identity ( $\exp(i*\theta) = \cos(\theta) + i*\sin(\theta)$ )  
applied to a matrix:  
  
>>> a = np.array([[1.0, 2.0], [-1.0, 3.0]])  
>>> expm(1j*a)  
array([[ 0.42645930+1.89217551j, -2.13721484-0.97811252j],  
       [ 1.06860742+0.48905626j, -1.71075555+0.91406299j]])  
>>> cosm(a) + 1j*sinm(a)  
array([[ 0.42645930+1.89217551j, -2.13721484-0.97811252j],  
       [ 1.06860742+0.48905626j, -1.71075555+0.91406299j]])  
  
""""  
a = np.asarray(A)  
if a.size == 1 and a.ndim < 2:  
    return np.array([[np.exp(a.item())]])  
  
if a.ndim < 2:  
    raise LinAlgError('The input array must be at least two-dimens
```

```

ional')
311         1         5000.0   5000.0       0.0       if a.shape[-1] != a.shape[-2]:
312                                     raise LinAlgError('Last 2 dimensions of the array must be square')
313
314                                     # Empty array
315         1         4000.0   4000.0       0.0       if min(*a.shape) == 0:
316                                     dtype = expm(np.eye(2, dtype=a.dtype)).dtype
317                                     return np.empty_like(a, dtype=dtype)
318
319                                     # Scalar case
320         1         5000.0   5000.0       0.0       if a.shape[-2:] == (1, 1):
321                                     return np.exp(a)
322
323         1         18000.0  18000.0       0.0       if not np.issubdtype(a.dtype, np.inexact):
324                                     a = a.astype(np.float64)
325         1         8000.0   8000.0       0.0       elif a.dtype == np.float16:
326                                     a = a.astype(np.float32)
327
328                                     # An explicit formula for 2x2 case exists (formula (2.2) in [1]).
However, without
329                                     # Kahan's method, numerical instabilities can occur (See gh-1958
4). Hence removed
330                                     # here until we have a more stable implementation.
331
332         1         4000.0   4000.0       0.0       n = a.shape[-1]
333         1         10000.0  10000.0       0.0       eA = np.empty(a.shape, dtype=a.dtype)
334                                     # working memory to hold intermediate arrays
335         1         5000.0   5000.0       0.0       Am = np.empty((5, n, n), dtype=a.dtype)
336
337                                     # Main loop to go through the slices of an ndarray and passing to
expm
338         2         18000.0   9000.0       0.0       for ind in product(*[range(x) for x in a.shape[:-2]]):
339         1         4000.0   4000.0       0.0         aw = a[ind]
340
341         1         87000.0   87000.0       0.0         lu = bandwidth(aw)
342         1         5000.0   5000.0       0.0         if not any(lu): # a is diagonal?
343                                     eA[ind] = np.diag(np.exp(np.diag(aw)))
344                                     continue
345
346                                     # Generic/triangular case; copy the slice into scratch and send
d.
347                                     # Am will be mutated by pick_pade_structure
348                                     # If s != 0, scaled Am will be returned from pick_pade_structu
re.
349         1         7285000.0 7.28e+06       0.5       Am[0, :, :] = aw
350         1        363975000.0 3.64e+08      26.0       m, s = pick_pade_structure(Am)
351         1         17000.0   17000.0       0.0       if (m < 0):

```

```

352 raise MemoryError("scipy.linalg.expm could not allocate su
fficient"
353 " memory while trying to compute the Pad
e "
354 f"structure (error code {m}).")
355 info = pade_UV_calc(Am, m)
356 if info != 0:
357     if info <= -11:
358         # We raise it from failed mallocs; negative LAPACK cod
es > -7
359         raise MemoryError("scipy.linalg.expm could not allocat
e "
360 "sufficient memory while trying to compu
te the "
361 f"exponential (error code {info}).")
362 else:
363     # LAPACK wrong argument error or exact singularity.
364     # Neither should happen.
365     raise RuntimeError("scipy.linalg.expm got an internal
LAPACK "
366 "error during the exponential compu
tation "
367 f"(error code {info})")
368 1      8000.0   8000.0   0.0      eAw = Am[0]
369
370 1      4000.0   4000.0   0.0      if s != 0: # squaring needed
371
372 1      4000.0   4000.0   0.0      if (lu[1] == 0) or (lu[0] == 0): # lower/upper triangular
373         # This branch implements Code Fragment 2.1 of [1]_
374
375         diag_aw = np.diag(aw)
376         # einsum returns a writable view
377         np.einsum('ii->i', eAw)[:]= np.exp(diag_aw * 2*(-s))
378         # super/sub diagonal
379         sd = np.diag(aw, k=-1 if lu[1] == 0 else 1)
380
381         for i in range(s-1, -1, -1):
382             eAw = eAw @ eAw
383
384             # diagonal
385             np.einsum('ii->i', eAw)[:]= np.exp(diag_aw * 2.**
(-i))
386             exp_sd = _exp_sinch(diag_aw * (2.**(-i))) * (sd *
2.**(-i))
387
388             if lu[1] == 0: # lower
389                 np.einsum('ii->i', eAw[1:, :-1])[:] = exp_sd
390             else: # upper
391                 np.einsum('ii->i', eAw[:-1, 1:])[:] = exp_sd

```

```

391
392                                     else: # generic
393         4         43000.0  10750.0      0.0         for _ in range(s):
394         3  323507000.0  1.08e+08      23.1         eAw = eAw @ eAw
395
396                                     # Zero out the entries from np.empty in case of triangular inp
ut
397         1         33000.0  33000.0      0.0         if (lu[0] == 0) or (lu[1] == 0):
398                                     eA[ind] = np.triu(eAw) if lu[0] == 0 else np.tril(eAw)
399         else:
400         1         1776000.0  1.78e+06      0.1         eA[ind] = eAw
401
402         1         6000.0   6000.0      0.0         return eA

```

From inspecting the output, these three lines took the longest:

Instruction	Hits	Time	Per hit	% Time
info = pade_UV_calc(Am, m)	1	702075000.0	7.02e+08	50.2
m,s = pick_pade_structure(Am)	1	363975000.0	3.64e+08	26.0
eAw = eAw @ eAw	3	323507000.0	1.08e+08	23.1

If you're curious and wanted to look at even lower level functions, it doesn't always work, because some functions are not implemented in Python, so you can't see their source code.

2c. Alternatives

Instead of doing a formal profiler, you can add in manual timing statements. This is less systematic, but sometimes (due to its simplicity) it's enough. Below are some examples.

Instructions

Go through the examples below

timeit

First, let's use the **timeit** package. You can load this like `import timeit` and use it explicitly, but in a jupyter/ipython notebook, the easiest way to use it is with **line magics** or **cell magics** (i.e., lines that start with `%` or `%%`, respectively).

The idea of **timeit** is that it runs your code a few times to get a good average value

Below is an example usage of **line magics**

```
In [24]: n = int(1e4)
%timeit sum_python(n)
%timeit sum_numpy(n)
```

266 μ s \pm 6.7 μ s per loop (mean \pm std. dev. of 7 runs, 1,000 loops each)
4.77 μ s \pm 14.4 ns per loop (mean \pm std. dev. of 7 runs, 100,000 loops each)

... and here's a **cell magic**. It will time the entire cell (so in this case, it doesn't tell us which line was slower)

```
In [25]: %%timeit
n = int(1e4)
sum_python(n)
sum_numpy(n)
```

274 μ s \pm 5.03 μ s per loop (mean \pm std. dev. of 7 runs, 1,000 loops each)

You can get fancy if you want, and give it flags. `r` tells is how many times to repeat the full run, and `n` is how many reps per run. (It generally guesses good values for these based on how fast the program runs)

```
In [26]: %%timeit -o -r 2 -n 1
# n for number of times to run the code, r to repeat the full runs
sum_numpy(n)

tmStructure = _ # this _ is the result of the previous computation
tmStructure
```

41.8 μ s \pm 22.7 μ s per loop (mean \pm std. dev. of 2 runs, 1 loop each)

```
Out[26]: <TimeitResult : 41.8  $\mu$ s  $\pm$  22.7  $\mu$ s per loop (mean  $\pm$  std. dev. of 2 runs, 1 loop each)>
```

time

We can also use the **time** package, which doesn't do any averaging for us:

```
In [27]: import time
```

```

start_time = time.time() # Record the start time
sum_numpy(n)
end_time = time.time()   # Record the end time

execution_time = end_time - start_time
print(f"Execution time: {execution_time} seconds")

# or...
start_perf_counter = time.perf_counter()
sum_numpy(n)
end_perf_counter = time.perf_counter()

execution_time_perf = end_perf_counter - start_perf_counter
print(f"Execution time (perf_counter): {execution_time_perf} seconds")

```

Execution time: 0.0002799034118652344 seconds

Execution time (perf_counter): 0.00032449999707750976 seconds

and like the `timeit` package, the `time` package also has **line magics** and **cell magics**:

```

In [28]: # Line magics
%time sum_python(n)
%time sum_numpy(n)

```

CPU times: user 477 μ s, sys: 215 μ s, total: 692 μ s

Wall time: 501 μ s

CPU times: user 70 μ s, sys: 17 μ s, total: 87 μ s

Wall time: 82 μ s

```

Out[28]: np.int64(50005000)

```

```

In [29]: %%time
sum_python(n)
sum_numpy(n)

```

CPU times: user 454 μ s, sys: 98 μ s, total: 552 μ s

Wall time: 466 μ s

```

Out[29]: np.int64(50005000)

```

Deliverables for 2c

None. Just make sure to turn in your work for parts 1 and 2b