How-To: Profile in Python

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Profiling in Python is provided by the cProfile module. We additionally use the SnakeViz graphical viewer (pip install snakeviz) to visualise the call stack, but if you do not have access to packages outside the standard Python library or a graphical user interface (SnakeViz is browser based) the pstats module can be used to inspect the profile instead. To profile a Python program contained in myscript.py:

```
python -m cProfile -o myscript.prof myscript.py
```

The output is saved to myscript.prof, and may be viewed with

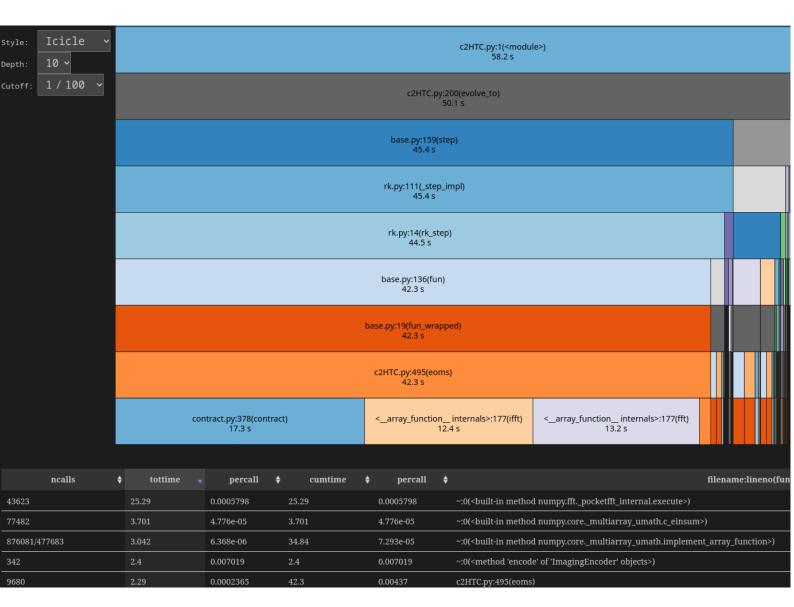
```
snakeviz myscript.prof
```

This should open a view in a new browser window. Below an extract for a script c2HTC.py whose main function evolve_to uses the Runge-Kutta solver (rk.py) from the scipy.integrate submodule to integrate a system of equations eoms over a fixed interval.

¹To print the 10 most significant function calls by cumulative time,

```
import pstats
p = pstats.Stats("myscript.prof")
print(p.sort_stats(pstats.SortKey.CUMULATIVE).print_stats(10))
```

For further details see the pstats. Stats class on the The Python Profiles documentation page.



The main two quantifiers are tottime, the total time spent in a function alone i.e. not including calls to sub-functions and cumtime, the cumulative time spent in this and all subfunctions (from invocation to exit). It is the latter that is shown above using nested bars.

Here we see out of 58 second total runtime the main function evolve_to took approximately 50; there was some initialisation and cleanup around this function. Moving the cursor over any block provides additional information (displayed in the sidebar): doing so for the first generically named base.py:159 reveals this is the base script in the scipy.integrate submodule.

Going down to the final line, what this profiling has revealed is that the majority of the runtime is split between performing tensor contractions with the contract function of opt_einsum and performing Fast Fourier Transforms with numpy's fft/ifft. In general one might check for unexpected timesinks (due to purely written or extraneous code) or at least identify the main computational bottleneck(s) which could be targeted for optimisation e.g. can a more efficient pathing algorithm be chosen for the tensor contractions?

Further considerations Typically you run the profiler on a 'standard' mode for your script. If your Python program has multiple modes of operations, you probably want to profile each 'mode' separately. Another useful strategy is to generate profiles whilst changing a parameter to identity the scaling of different parts of the computation with that parameter to e.g. determine the cause of runaway growth of computation time.