Week 10 Laboratory Exercises

Objectives

- Writing and use your own Python modules
- Exploring modules in Python

Preparation

Before the lab you should re-read the relevant lecture slides and their accompanying examples.

Getting Started

Set up for the lab by creating a new directory called lab10 and changing to this directory.

\$ mkdir lab10

\$ cd lab10

There are some provided files for this lab which you can fetch with this command:

\$ 2041 fetch lab10

If you're not working at CSE, you can download the provided files as a zip file or a tar file.

EXERCISE:

DNA analysis in Python

You have been given a file dna.py

dna.py is a python module that performs DNA analysis.

Compleate the functions in dna.py as described in the docstrings.

```
def read_dna(dna_file):
   Read a DNA string from a file.
   the file contains data in the following format:
   G <-> C
   G <-> C
   C <-> G
   G <-> C
   T <-> A
   Output a list of touples
       ('A', 'T'),
       ('G', 'C'),
       ('G', 'C'),
       ('C', 'G'),
       ('G', 'C'),
       ('T', 'A'),
   Where either (or both) elements in the string might be missing:
   <-> T
   G <->
   G <-> C
   <->
   <-> C
   T <-> A
   ....
   pass
def is_rna(dna):
   Given DNA in the aforementioned format,
   return the string "DNA" if the data is DNA,
   return the string "RNA" if the data is RNA,
   return the string "Invalid" if the data is neither DNA nor RNA.
   DNA consists of the following bases:
   Adenine ('A'),
   Thymine ('T'),
   Guanine ('G'),
   Cytosine ('C'),
   RNA consists of the following bases:
   Adenine ('A'),
   Uracil ('U'),
   Guanine ('G'),
   Cytosine ('C'),
   The data is DNA if at least 90% of the bases are one of the DNA bases.
   The data is RNA if at least 90% of the bases are one of the RNA bases.
   The data is invalid if more than 10% of the bases are not one of the DNA or RNA bases.
   Empty bases should be ignored.
   pass
def clean_dna(dna):
   Given DNA in the aforementioned format,
   If the pair is incomplete, ('A', '') or ('', 'G'), ect
   Fill in the missing base with the match base.
   In DNA 'A' matches with 'T', 'G' matches with 'C'
   In RNA 'A' matches with 'U', 'G' matches with 'C'
   If a pair contains an invalid base the pair should be removed.
   Pairs of empty bases should be ignored.
   pass
def mast_common_base(dna):
   Given DNA in the aforementioned format,
   return the most common first base:
   eg. given:
   A <-> T
```

```
G <-> C
   G <-> C
   C <-> G
   G <-> C
   T <-> A
   The most common first base is 'G'.
   Empty bases should be ignored.
   pass
def base_to_name(base):
   Given a base, return the name of the base.
   The base names are:
   Adenine ('A'),
   Thymine ('T'),
   Guanine ('G'),
   Cytosine ('C'),
   Uracil ('U'),
   return the string "Unknown" if the base isn't one of the above.
   pass
```

Download dna.py, or copy it to your CSE account using the following command:

```
$ cp -n /web/cs2041/22T2/activities/DNA/files.cp/dna.py
```

You have been given the file test_dna.py that uses the functions in dna.py to test your code.

```
$ ./test_dna.py data1
the file data1 is DNA
there are 98 pairs in the file
first 10 pairs:
G <-> C
T <-> A
A <-> T
G <-> C
A <-> T
G <-> C
T <-> A
C <-> G
T <-> A
A <-> T
last 10 pairs:
C <-> G
T <-> A
G <-> C
T <-> A
G <-> C
G <-> C
T <-> A
C <-> G
A <-> T
T <-> A
the most common base is Guanine
```

```
NOTE:
```

Your answer must be Python only. You can not use other languages such as Shell, Perl or C.

You must not use subprocess, or otherwise run external programs.

When you think your program is working, you can use autotest to run some simple automated tests:

```
$ 2041 autotest DNA
```

When you are finished working on this exercise, you must submit your work by running give:

```
$ give cs2041 lab10_DNA dna.py
```

before Monday 08 August 12:00 to obtain the marks for this lab exercise.

CHALLENGE EXERCISE:

Bashful Python

We have some Shell (Bash) scripts that do arithmetic calculations that we need to translate to Python.

Write a Python program bashpy.py which takes such a Bash script on stdin and outputs an equivalent Python program.

The scripts use the arithmetic syntax supported by Bash (and several other shells). Fortunately, the scripts only use a very limited set of shell features.

You can assume all the features you need to translate are present in the following 4 examples.

sum.sh sums a series of integers:

```
1=>((1 + 1))
done
echo $sum
$ ./sum.sh
$ ./bashpy.py < sum.sh</pre>
#!/usr/bin/env python3
# sum the integers $start .. $finish
start = 1
finish = 100
sum = 0
i = 1
while i <= finish:
    sum = sum + i
    i = i + 1
print(sum)
$ ./bashpv.pv < sum.sh | pvthon3
```

double.sh prints powers of two:

```
j=1
while ((i < 31))
   j=\$((j + j))
    i=$((i + 1))
    echo $i $j
done
$ ./double.sh
2 2
3 4
4 8
5 16
6 32
7 64
8 128
9 256
10 512
11 1024
12 2048
13 4096
```

• pythagorean_triple.sh searches for Pythagorean triples:

```
while c < max:
            if a * a + b * b == c * c:
               print(a, b, c)
            c = c + 1
        b = b + 1
    a = a + 1
$ ./bashpy.py < pythagorean_triple.sh | python3</pre>
3 4 5
5 12 13
6 8 10
7 24 25
8 15 17
9 12 15
9 40 41
10 24 26
12 16 20
12 35 37
15 20 25
15 36 39
16 30 34
18 24 30
```

collatz.sh prints an interesting series:

```
$ cat collatz.sh
#!/bin/bash
# https://en.wikipedia.org/wiki/Collatz_conjecture
# https://xkcd.com/710/
n=65535
while ((n != 1))
   if ((n % 2 == 0))
       n=$((n / 2))
    else
       n=\$((3 * n + 1))
    fi
    echo $n
done
$ ./bashpy.py <collatz.sh</pre>
#!/usr/bin/env python3
# https://en.wikipedia.org/wiki/Collatz_conjecture
```

HINT:

Don't worry about the many possible features that can occur in shell scripts.

Focus on translating correctly the very limited set of Shell in the examples you are given.

For example, the Shell scripts you are given don't contain strings (quoted or unquoted), so don't worry about translating these.

You can translate each line of the shell script individually using simple string operations or research and resub.

NOTE:

You can assume the Shell scripts are always indented in the same manner as the example you've given

in other words the indenting will be compatible with Python.

You must translate the bash scripts to Python not just embed the shell of the script in a Python system call.

Your answer must be Python only. You can not use other languages such as Shell, Perl or C.

You must not use subprocess, or otherwise run external programs.

When you think your program is working, you can use autotest to run some simple automated tests:

\$ 2041 autotest bashpy

When you are finished working on this exercise, you must submit your work by running give:

```
$ give cs2041 lab10_bashpy bashpy.py
```

before Monday 08 August 12:00 to obtain the marks for this lab exercise.

CHALLENGE EXERCISE:

When Regular Expressions Aren't Regular

Write a regular expression that validates a JSON file.

In other words, write a regex that matches a string iff (if and only if) that string is valid JSON.

Here is a test program assist you in doing this:

```
#! /usr/bin/env python3
from sys import argv, stderr
import regex
regex.DEFAULT_VERSION = regex.V1
assert len(argv) == 3, f"Usage: {argv[0]} <json file> <regex file>"
json_file, regex_file = argv[1], argv[2]
try:
   with open(json_file) as json_data, open(regex_file) as regex_data:
       if regex.search(regex_data.read(), json_data.read(), timeout=5):
            # In the test suite, all files that start with "y " should be valid.
           print(f"Valid JSON file: {json_file}")
            # In the test suite, all files that start with "n " should be invalid.
           print(f"Invalid JSON file: {json_file}")
except TimeoutError as e:
   # Allow a timeout error to signal that the jason file is not valid
   print(f"Invalid JSON file: {json_file}")
   # This is printed to stderr so that it is not captured by the test
   print(f"5 second time limit reached while reading {json_file}", file=stderr)
```

Download <u>test_regex_json.py</u>, or copy it to your CSE account using the following command:

```
$ cp -n /web/cs2041/22T2/activities/regex_json/test_regex_json.py
test_regex_json.py
```

You have been given a directory JSONTestSuite containing a number of JSON files.

There are two types of files in this directory:

Files starting with y are valid JSON files.

Files starting with n_ are invalid JSON files.

Put your solution in regex_json.txt:

For example to test the regex ^.+\$

If your solution is correct, all files in the JSONTestSuite starting with y_ should be labelled valid, and all files starting with y_ should be labelled invalid,

NOTE:

This exercise is not possible with true regular expression, i.e using |*() alone,

You will need to use additional regular expression syntax to achieve the same result.

Not even Python regular expression syntax enough.

You will need to use the PCRE regular expression library regex .

Autotest will use install ans use this module for you.

But you may need to install it manually for local testing.

The regex module extends the Python re module.

So the Python regular syntax documentation will still be useful.

You will additionally need to read the documentation for the regex module.

This is (another) (in)?famous problem to try and solve with regex.

There will be answers easily found online.

Don't google for other people solutions - see if you can come up with your own.

HINT:

You are not able to change the flags = argument of the regex.search function.

But there are ways to enable and disable flags within the regex itself.

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Python supports the <u>regex.VERBOSE</u> <u>flag</u>.

This flag allows you to add comments to your regular expression.

Plus whitespace within the pattern is ignored, except when in a character class, or when preceded by an unescaped backslash This allows you to write much more readable regular expressions.

When you think your program is working, you can use autotest to run some simple automated tests:

\$ 2041 autotest regex_json

When you are finished working on this exercise, you must submit your work by running give:

\$ give cs2041 lab10_regex_json regex_json.txt

before Monday 08 August 12:00 to obtain the marks for this lab exercise.

Submission

When you are finished each exercises make sure you submit your work by running give .

You can run give multiple times. Only your last submission will be marked.

Don't submit any exercises you haven't attempted.

If you are working at home, you may find it more convenient to upload your work via give's web interface.

Remember you have until Week 11 Monday 12:00:00 to submit your work.

You cannot obtain marks by e-mailing your code to tutors or lecturers.

You check the files you have submitted here.

Automarking will be run by the lecturer several days after the submission deadline, using test cases different to those autotest runs for you. (Hint: do your own testing as well as running autotest.)

After automarking is run by the lecturer you can view your results here. The resulting mark will also be available via give's web interface.

Lab Marks

When all components of a lab are automarked you should be able to view the the marks <u>via give's web interface</u> or by running this command on a CSE machine:

\$ 2041 classrun -sturec

the <u>school of compater science and Engineering</u>

at the <u>University of New South Wales</u>, Sydney.

For all enquiries, please email the class account at cs2041@cse.unsw.edu.au

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