

# Sequences II

## Warm-up

Last class, we looked at `for`-loops. Test your knowledge by answering the following questions:

1. Write a function called `get_int` that takes in a list of digits, calculates and returns the number corresponding to the digits in the list. For example, `get_int([3,7,1])` must return 371.

```
1 def get_int(digits: list) -> int:
2     number=0
3     for digit in digits:
4         number = number * 10 + digit
5     return number
```

2. Provide testcases that you would use to test the `get_int` function.

```
get_int([]) must return 0 and get_int([1]) must return 1
get_int([1,2]) must return 12 and get_int([2,1]) must return 21
get_int([0,0,1]) must return 1 and get_int([1,0,0]) must return 100
```

3. Write a function called `draw_pyramid` that takes in an integer and draws a pyramid of the size equal to the integer parameter. For example, for `draw_pyramid(1)` the function must print:

```
*
* *
```

For `draw_pyramid(2)` the function must print:

```
  *
 * *
*  *
```

For `draw_pyramid(3)` the function must print:

```
   *
  * *
 *  *
*   *
```

```
1 def draw_pyramid(size: int) -> None:
2     print(' '*size + '*')
3     for i in range(1,size+1):
4         print(' '*(size-i) + '*' + ' '*(2*i-1) + '*')
```

## Model 1 Indexing and Slicing

A string is a sequence of characters in single quotes ('') or double quotes (""). Depending on the application, we can treat a string as a single value (e.g., dna), or we can access individual characters using square brackets (e.g., dna[0]). We can also use *slice notation* (e.g., dna[4:8]) to refer to a range of characters. In fact, all types of sequences (including list) support indexing and slicing.

4. Complete the table below to further explore how strings work:

Python code	Output
dna = 'CTGACGACTT'	
dna[5]	'G'
dna[10]	IndexError: index out of range
len(dna)	10
dna[:5]	'CTGAC'
dna[5:]	'GACTT'
dna[5:10]	'GACTT'
triplet = dna[2:5]	
print(triplet)	GAC
dna[-5]	'G'
dna[-10]	'C'
dna[:-5]	'CTGAC'
dna[-5:]	'GACTT'
triplet = dna[-4:-1]	
print(triplet)	'ACT'

5. What is the *positive* index of each character in the dna string? Check your answers above.

Character:	C	T	G	A	C	G	A	C	T	T
Index:	0	1	2	3	4	5	6	7	8	9

6. What is the *negative* index of each character in the dna string? Check your answers above.

Character:	C	T	G	A	C	G	A	C	T	T
Index:	-10	-9	-8	-7	-6	-5	-4	-3	-2	-1

7. Based on the previous questions, what are `dna[2]` and `dna[-2]`? Explain your answers.

They are G and T, respectively. Index 2 means to the third from the left, and index -2 means the second from the right.

8. Explain the `IndexError` you observed. What is the range of indexes for the `dna` string?

Because the length of the string is 10, the indexes range from 0 to 9. Therefore, `dna[10]` is out of range.

9. Consider the notation of the operator `[m:n]` for slicing the string.

a) Is the value at the start of the resulting string the same as the value at index `m` (i.e., `dna[m]`)?

If not, describe what it is. Yes; `m` is the first character in the slice.

b) Is the value at the end of the resulting string the same as the value at index `n` (i.e., `dna[n]`)?

If not, describe what it is. No; `n` is the index after the last character.

c) Explain what it means when only a single number is referenced when creating a slice, such as `[m:]` or `[:n]`.

The slice `[m:]` means “from the index `m` to the end”. The slice `[:n]` means “from the beginning to the index just before `n`” (i.e., the first `n` characters).

10. What is the simplest way to get the first three characters of `dna`? What is the simplest way to get the last three characters?

Based on the previous question, we know that `dna[:3]` gets the first three characters. To get the last three, we use `dna[-3:]`.

11. Write a Python expression that slices `'GACT'` from `dna` using positive indexes. Then write another expression that slices the same string using negative indexes.

```
dna[5:9]    dna[-5:-1]
```

12. Write a Python assignment statement that uses the `len` function to assign the last letter of `dna` to the variable `last`.

```
last = dna[len(dna) - 1]
```

13. Write a Python assignment statement that uses a negative index to assign the last letter of `dna` to the variable `last`.

```
last = dna[-1]
```

## Model 2 Working with Lists

Lists have *methods* (like built-in functions) that can be called using dot notation. For example, to add a new element to the end of a list, we can use the `append` method. See <https://docs.python.org/3/tutorial/datastructures.html#more-on-lists> for more details. The back of the handout also has a list of select functions with a novice friendly documentation.

Python code	Output
<code>rolls = [4, 6, 6, 2, 6]</code>	
<code>len(rolls)</code>	5
<code>print(rolls[5])</code>	IndexError: list index out of range
<code>rolls.append(1)</code>	
<code>print(rolls)</code>	[4, 6, 6, 2, 6, 1]
<code>print(rolls[5])</code>	1
<code>lucky.append(1)</code>	NameError: name 'lucky' is not defined
<code>lucky = []</code>	
<code>print(lucky[0])</code>	IndexError: list index out of range
<code>lucky.append(5)</code>	
<code>print(lucky)</code>	[5]
<code>print(lucky[0])</code>	5
<code>rolls.count(6)</code>	3
<code>rolls.remove(6)</code>	
<code>print(rolls)</code>	[4, 6, 2, 6, 1]
<code>help(rolls.remove)</code>	remove first occurrence of value
<code>help(rolls)</code>	Help on list object (multiple pages)

14. What is the result of calling the `append` method on a list?

The value gets added to the end of the list. Nothing is returned.

15. What must be defined prior to using a method like `append`?

The list itself; `lucky.append(5)` is an error if `lucky` is not defined.

16. Explain why two lines of code caused an `IndexError`.

In both cases, we asked for an index that was out of range. If the length of an index is  $n$ , the highest index is  $n - 1$ .

17. What is the result of calling the remove method on a list?

It removes the first occurrence of a value. The list changes as a result of this method.

18. Give one example of a list method that requires an argument and one that does not.

Methods that require arguments: append, count, extend, index, insert, remove. Methods that do not: clear, copy, pop, reverse, sort.

19. Describe the similarities and differences between using a list method like append and Python built-in functions like print.

Both use parentheses and take arguments. The list methods come after the dot operator, and the built-in functions surround the list itself.

## Model 3 Common String Methods

Like lists, strings have *methods* (built-in functions) that can be called using dot notation. See <https://docs.python.org/3/library/stdtypes.html#string-methods> for more details. The back of the handout also has a list of select functions.

Python code	Output
<code>dna = 'CTGACGACTT'</code>	
<code>dna.lower()</code>	<code>'ctgacgactt'</code>
<code>print(dna)</code>	<code>CTGACGACTT</code>
<code>lowercase = dna.lower()</code>	
<code>print(lowercase)</code>	<code>ctgacgactt</code>
<code>dnalist = list(dna)</code>	
<code>print(dnalist)</code>	<code>['C', 'T', 'G', 'A', 'C', 'G', 'A', 'C', 'T', 'T']</code>
<code>dnalist.reverse()</code>	
<code>print(dnalist)</code>	<code>['T', 'T', 'C', 'A', 'G', 'C', 'A', 'G', 'T', 'C']</code>
<code>type(dna)</code>	<code>&lt;class 'str'&gt;</code>
<code>dna = dna.split('A')</code>	
<code>print(dna)</code>	<code>['CTG', 'CG', 'CTT']</code>
<code>type(dna)</code>	<code>&lt;class 'list'&gt;</code>
<code>dna.replace('C', 'g')</code>	<code>AttributeError: 'list' object has no 'replace'</code>
<code>print(dna[0])</code>	<code>CTG</code>
<code>type(dna[0])</code>	<code>&lt;class 'str'&gt;</code>
<code>dna[0].replace('C', 'g')</code>	<code>'gTG'</code>
<code>print(dna)</code>	<code>['CTG', 'CG', 'CTT']</code>

20. Does the `lower` method change the contents of the `dna` string? Justify your answer.

No, it does not. The next line of code prints `dna`, which is unchanged.

21. Describe the `list` function—what does `list(dna)` return?

It returns a list of the individual characters. Each element of the list is a string of length 1. (Note that Python does not have a character data type.)

22. Why is it possible to call the `replace` method on `dna[0]` but not `dna`?

The `list` data type does not include a `replace` method. However, strings allow you to “find and replace” any text.

23. Name several other string methods not shown in the provided code. (Read the documentation.)

There are dozens of string methods; the model only uses `lower`, `split`, and `replace`.

24. Consider the application of a method on a variable:

a) Does a string variable change after applying a method? Provide justification.

No it doesn't; neither `lower` nor `replace` modify the string.

b) Does a list variable change after applying a method? Provide justification.

It might; for example, the `reverse` method changes the list.

c) Identify the data type that is *immutable* (i.e., the value never changes). `String`

25. Write a single statement to change the final contents of `dna` to `['CTG', 'cc', 'CTT']`. Confirm that your code works in a Python Shell.

```
dna[1] = 'cc'
```

26. Why do you think Python has a `replace` method for strings but not for lists?

Answers may vary. One reason might be that lists are more complex than strings: they can store any type of data, not just characters. Another reason might be that there are fewer applications of replacing data in lists than patterns in text.

## List methods

- `append(item)` — Adds a new item to the end of a list
- `insert(position, item)` — Inserts a new item at the position given
- `extend(lst)` — Extend the list by appending all the items from lst
- `pop()` — Removes and returns the last item
- `pop(position)` — Removes and returns the item at position
- `sort()` — Modifies a list to be sorted
- `reverse()` — Modifies a list to be in reverse order
- `index(item)` — Returns the position of first occurrence of item
- `count(item)` — Returns the number of occurrences of item
- `remove(item)` — Removes the first occurrence of item
- `copy()` — Return a clone of the list
- `clear()` — Remove all items from the list

## String methods

- `upper()` — Returns a string in all uppercase
- `lower()` — Returns a string in all lowercase
- `capitalize()` — Returns a string with first character capitalized, the rest lower
- `strip()` — Returns a string with the leading and trailing whitespace removed
- `lstrip()` — Returns a string with the leading whitespace removed
- `rstrip()` — Returns a string with the trailing whitespace removed
- `count(item)` — Returns the number of occurrences of item
- `replace(old, new)` — Replaces all occurrences of old substring with new

- `center(width)` — Returns a string centered in a field of width spaces
- `ljust(width)` — Returns a string left justified in a field of width spaces
- `rjust(width)` — Returns a string right justified in a field of width spaces
- `find(item)` — Returns the leftmost index where the substring `item` is found, or -1 if not found
- `rfind(item)` — Returns the rightmost index where the substring `item` is found, or -1 if not found
- `index(item)` — Like `find` except causes a runtime error if `item` is not found
- `rindex(item)` — Like `rfind` except causes a runtime error if `item` is not found
- `split(separator)` — Return a list of the words in the string, using (optional) `separator` as the delimiter string
- `join(lst)` — Return a string which is the concatenation of the strings in `lst`
- `isalpha()` — Return True if all characters in the string are alphabetic and there is at least one character
- `isdigit()` — Return True if all characters in the string are decimal characters and there is at least one character
- `islower()` — Return True if all cased characters in the string are lowercase and there is at least one cased character
- `isspace()` — Return True if there are only whitespace characters in the string and there is at least one character
- `isupper()` — Return True if all cased characters in the string are uppercase and there is at least one cased character