Introduction to Python

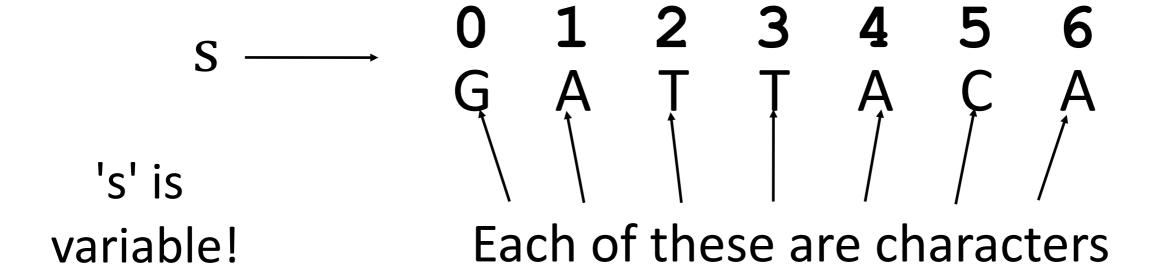
Strings

Components of Python

- 1) Variables (Me hold data)
- 2) Data-type (integers, float,)
- 3) Data-structures (list, tuple, dicitonary)
- 4) Loop (for, while)
- 5) Branch (if, elif, else)
- 6) Methods
- 7) Functions

Computers store text as strings

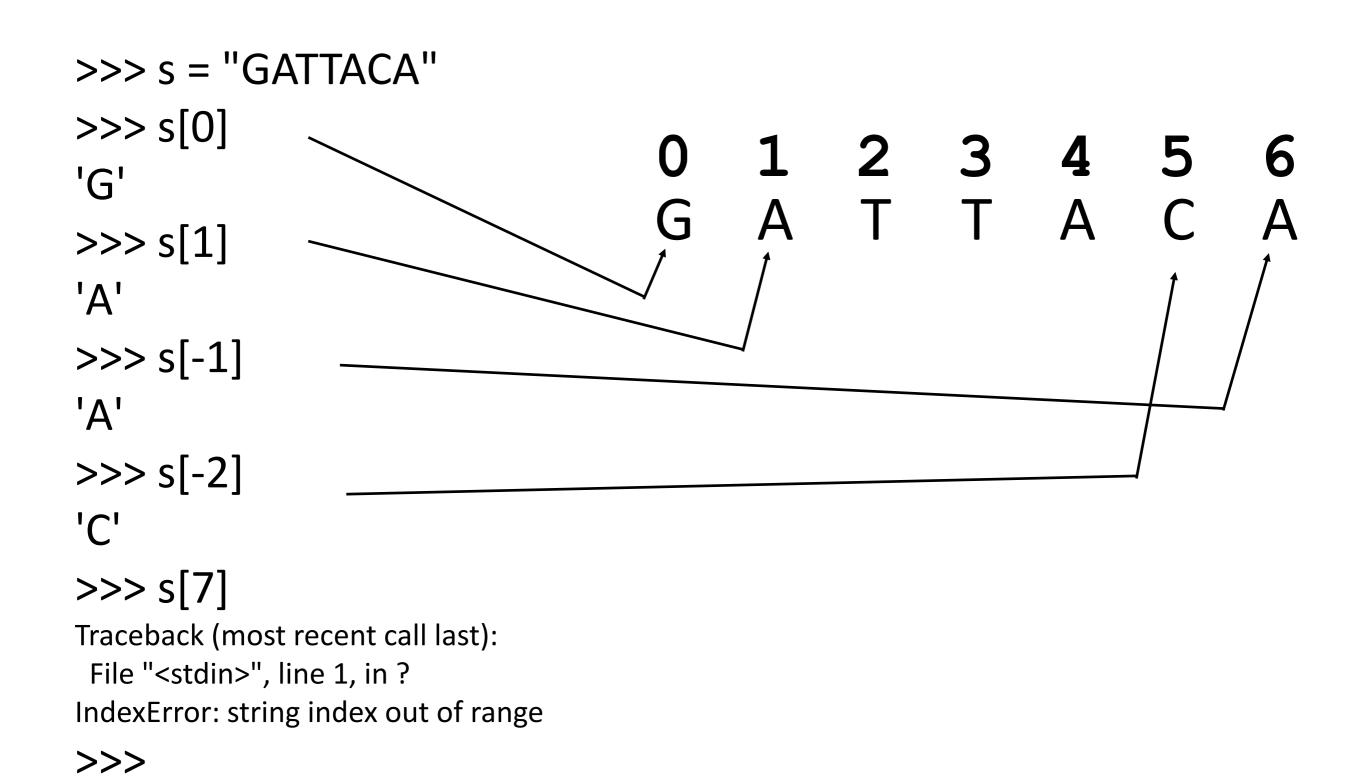
$$>>>$$
 s = "GATTACA"



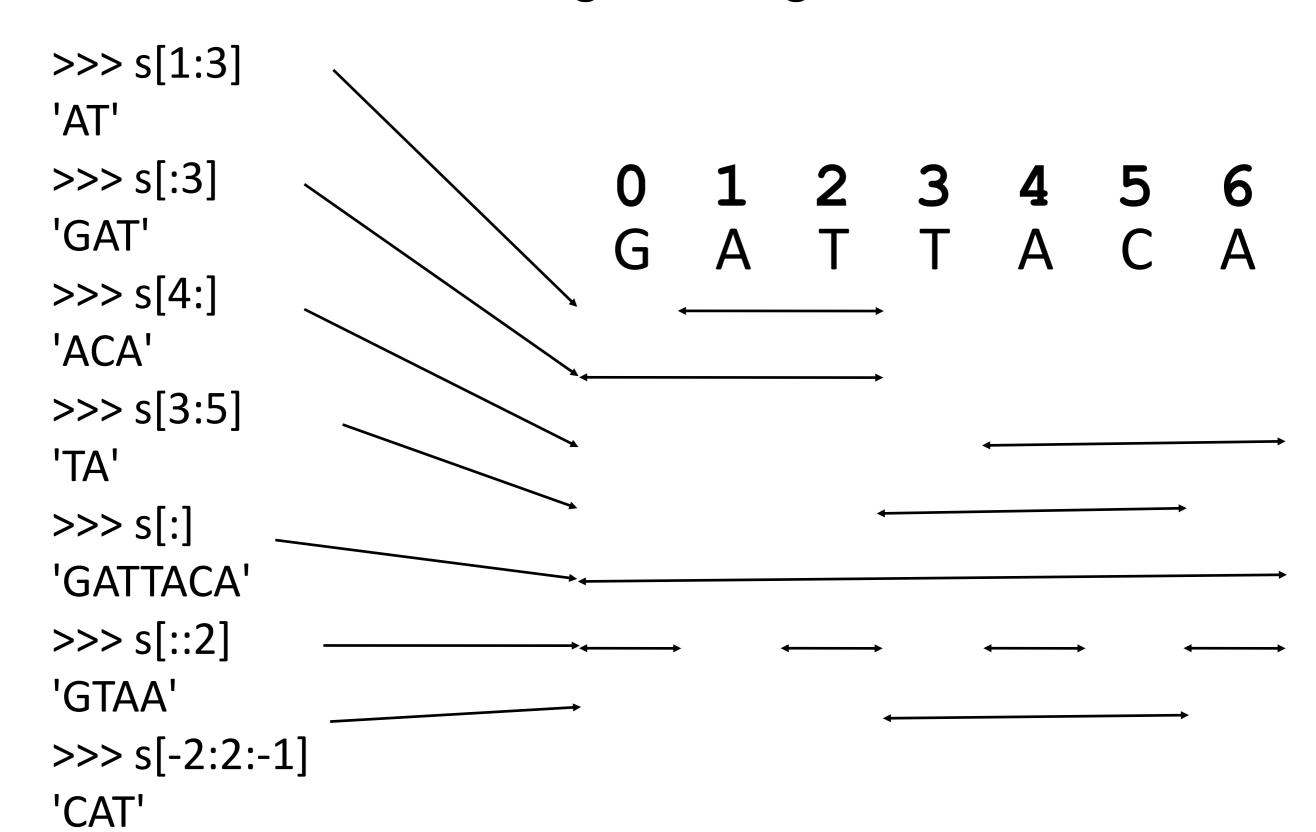
Why are strings important?

- Sequences are strings
 - ..catgaaggaa ccacagccca gagcaccaag ggctatccat..
- Database records contain strings
 - LOCUS AC005138
 - DEFINITION Homo sapiens chromosome 17, clone hRPK.261_A_13, complete sequence
 - AUTHORS Birren, B., Fasman, K., Linton, L., Nusbaum, C. and Lander, E.
- HTML is one (big) string

Getting Characters



Getting substrings



Creating strings

Strings start and end with a single or double quote characters (they must be the same)

"This is a string"
"This is another string"

"Strings can be in double quotes"
'Or in single quotes.'
'There's no difference.'
'Okay, there\'s a small one.'

Special Characters and Escape Sequences

Backslashes (\) are used to introduce special characters

>>> s = 'Okay, there\'s a small one.'

The \ "escapes" the following single quote

```
>>> print s
Okay, there's a small one.
```

Some special characters

Esca	pe	Seq	uer	ıce
	L			

Meaning

\\ Backslash (keep a \)

Single quote (keeps the ')

\" Double quote (keeps the ")

\n Newline

\t Tab

Working with strings

```
length
>>> len("GATTACA")
>>> "GAT" + "TACA"
                               concatenation
'GATTACA'
>>> "A" * 10
                               repeat
'AAAAAAAAAA'
>>> "G" in "GATTACA"
                               substring test
True
>>> "GAT" in "GATTACA"
True
>>> "AGT" in "GATTACA"
False
                               substring location
>>> "GATTACA".find("ATT")
                               substring count
>>> "GATTACA".count("T")
>>>
```

Converting from/to strings

```
>>> "38" + 5
Traceback (most recent call last):
  File "<stdin>", line 1, in ?
TypeError: cannot concatenate 'str' and 'int' objects
>>> int("38") + 5
43
>>> "38" + str(5)
13851
>>> int("38"), str(5)
(38, '5')
>>> int("2.71828")
Traceback (most recent call last):
  File "<stdin>", line 1, in ?
ValueError: invalid literal for int(): 2.71828
>>> float("2.71828")
2.71828
>>>
```

Change a string?

Strings cannot be modified They are immutable Instead, create a new one

```
>>> s = "GATTACA"
>>> s[3] = "C"
Traceback (most recent call last):
   File "<stdin>", line 1, in ?
TypeError: object doesn't support item
assignment
>>> s = s[:3] + "C" + s[4:]
>>> s
'GATCACA'
>>>
```

Some more methods

```
>>> "GATTACA".lower()
'gattaca'
>>> "gattaca".upper()
'GATTACA'
>>> "GATTACA".replace("G", "U")
'UATTACA'
>>> "GATTACA".replace("C", "U")
'GATTAUA'
>>> "GATTACA".replace("AT", "**")
'G**TACA'
>>> "GATTACA".startswith("G")
True
>>> "GATTACA".startswith("g")
False
>>>
```

Ask for a string

The Python function "raw_input" asks the user (that's you!) for a string

```
>>> seq = raw_input("Enter a DNA sequence: ")
Enter a DNA sequence: ATGTATTGCATATCGT
>>> seq.count("A")
4
>>> print "There are", seq.count("T"), "thymines"
There are 7 thymines
>>> "ATA" in seq
True
>>> substr = raw_input("Enter a subsequence to find: ")
Enter a subsequence to find: GCA
>>> substr in seq
True
>>>
```

Ask the user for a sequence then print its length

```
Enter a sequence: ATTAC It is 5 bases long
```

Modify the program so it also prints the number of A, T, C, and G characters in the sequence

```
Enter a sequence: ATTAC
```

It is 5 bases long

adenine: 2

thymine: 2

cytosine: 1

guanine: 0

Modify the program to allow both lowercase and upper-case characters in the sequence

```
Enter a sequence: ATTgtc It is 6 bases long adenine: 1 thymine: 3 cytosine: 1 quanine: 1
```

Modify the program to print the number of unknown characters in the sequence

```
Enter a sequence: ATTU*gtc
It is 8 bases long
adenine: 1
thymine: 3
cytosine: 1
guanine: 1
unknown: 2
```