# Dictionaries and strings (part 2)

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- Quiz
- Exercise 6.7
- String manipulation

#### Question A

```
d = {-2:-1, -1:0, 0:1, 1:2, 2:-2}
print(d[0])
# What is printed out?
```

#### Question B

```
d = {-2:-1, -1:0, 0:1, 1:2, 2:-2}
print(d[d[0]])
# What is printed out?
```

#### Question C

```
d = {-2:-1, -1:0, 0:1, 1:2, 2:-2}
print(d[-2]*d[2])
# What is printed out?
```

#### Question A

table = {'age':[35,20], 'name':['Anna','Peter']}
for key in table:
 print('%s: %s' % (key,table[key]))
# What is printed out?

#### Question B

```
table = {'age':[35,20], 'name':['Anna','Peter']}
vals = list(table.values())
print(vals)
print(vals[0])
print(vals[0][0])
# What is printed out?
```

#### Question C

```
table = {'age':[35,20], 'name':['Anna','Peter']}
print(table['name'][1], table['age'][1])
# What is printed out?
```

### Question A

d = {3:5, 6:7}
e = {4:6, 7:8}
d.update(e)
# What is the content of dictionary d now?

### Question B

### Question C

```
d = {6:100}
e = {6:6, 7:8}
d.update(e)
# What is the content of dictionary d now?
```

#### The file 'teledata.txt' gives information about mobile customers:

Age 45	Income 720k	Gender Female	Monthly calls 46	ID A001
27	440k	Male	3	A002
17	0	Male Fomalo	52 19	A006
24		remare		A014

- How could you store the data using five lists?
- How could you store the data using one list?
- How could you store the data in a dictionary (what information would be key and what datatype would you use for the values)?

### Make a nested dictionary from a file

The file human\_evolution.txt holds information about various human species and their height, weight, and brain volume. Make a program that reads this file and stores the tabular data in a nested dictionary humans. The keys in humans correspond to the species name (e.g., H. erectus), and the values are dictionaries with keys 'period', 'height', 'weight', 'volume'. For example,

humans['H. habilis']['weight']

should equal '55 - 70'. Let the program print to screen the humans dictionary in a nice tabular form similar to that in the file.

Filename: humans

## Step 1: reading the file

### We first download the file and inspect it visually:

Species	Lived when (mill. yrs)	Adult height (m)	Adult mass (kg)	Brain volume (cm**3)
<ul> <li>H. habilis</li> <li>H. erectus</li> <li>H. ergaster</li> <li>H. heidelbergensis</li> <li>H. neanderthalensis</li> <li>H. sapiens sapiens</li> <li>H. floresiensis</li> </ul>	2.2 - 1.6 1.4 - 0.2 1.9 - 1.4 0.6 - 0.35 0.35 - 0.03 0.2 - present 0.10 - 0.012	1.0 - 1.5 1.8 1.9 1.8 1.6 1.4 - 1.9 1.0	$\begin{array}{r} 33 & - 55 \\ 60 \\ 55 & - 70 \\ 50 & - 100 \\ 25 \end{array}$	660 850 (early) - 1100 (late) 700 - 850 1100 - 1400 1200 - 1700 1000 - 1850 400

Source: http://en.wikipedia.org/wiki/Human\_evolution

To read the table, we need to skip some lines at the top and bottom. How do we determine where the data start and stop?

- Solution 1: we see that the data span lines 4-10.
- Solution 2: data lines always start with 'H. '.
- Solution 3: data occur between the lines with hyphens.

All would work, but here we go for the third solution.

```
# Read all lines into a list
infile = open('human_evolution.txt', 'r')
lines = infile.readlines()
# Find first line with data
k = 0
while lines[k][0] != '-': # When no hyphen
 k = k + 1
                  # ... we continue the search
first = k + 1
                         # First line after hyphen
# Find last line with data
k = first
                         # Start point for search
while lines[k][0] != '-': # When no hyphen
   k = k + 1
                      # ... we continue the search
                       # Last line before hyphen
last = k - 1
# Now we are ready to process the data
for i in range(first, last+1):
    # Do something with lines[i]
```

## Step 2: splitting a line into columns

Species	Lived when	Adult	Adult	Brain volume
	(mill. yrs)	height (m)	mass (kg)	(cm**3)
H. habilis H. erectus H. ergaster H. heidelbergensis H. neanderthalensis H. sapiens sapiens H. floresiensis	2.2 - 1.6 1.4 - 0.2 1.9 - 1.4 0.6 - 0.35 0.35 - 0.03 0.2 - present 0.10 - 0.012	1.0 - 1.5 1.8 1.9 1.8 1.6 1.4 - 1.9 1.0	33 - 55 60 55 - 70 50 - 100 25	660 850 (early) - 1100 (late) 700 - 850 1100 - 1400 1200 - 1700 1000 - 1850 400

Source: http://en.wikipedia.org/wiki/Human\_evolution

Want to split each data line into columns, for example:

```
words[0] : 'H. habilis'
words[1] : '2.2 - 1.6'
words[2] : '1.0 - 1.5'
...
```

Possible solutions:

- Split on whitespace but how to go from there?
- Find position of each column from the header

Here we go for the second solution.

```
# Read all lines into a list
infile = open('human_evolution.txt', 'r')
lines = infile.readlines()
# Find column positions from second line in file
s = lines[1]
start = [0, s.index('(mill. yrs)'),
               s.index('height (m)'),
               s.index('mass (kg)'),
               s.index('(cm**3)')]
stop = start[1:len(start)] + [80]
# start: [ 0, 21, 37, 50, 62]
# stop: [21, 37, 50, 62, 80]
# The k'th column in the i'th line is now easy to find:
# words[0] = lines[i][start[0]:stop[0]]
# words[1] = lines[i][start[1]:stop[1]]
# ...etc
```

## Putting step 1 and 2 together

```
infile = open('human_evolution.txt', 'r')
lines = infile.readlines()
s = lines[1]
start = [0, s.index('(mill. yrs)'), s.index('height (m)'), ...]
stop = start[1:len(start)] + [80]
\mathbf{k} = 0
while lines[k][0] != '-':
k = k + 1
first = k + 1
k = first
while lines[k][0] != '-':
 k = k + 1
last = k - 1
humans = \{\}
for i in range(first, last+1):
    species = lines[i][start[0]:stop[0]]
    period = lines[i][start[1]:stop[1]]
    height = lines[i][start[2]:stop[2]]
    weight = lines[i][start[3]:stop[3]]
    volume = lines[i][start[4]:stop[4]]
    # Store the data in a dictionary
```

Consider the last step in the algorithm above:

```
for i in range(first, last+1):
    species = lines[i][start[0]:stop[0]].strip()
    period = lines[i][start[1]:stop[1]].strip()
    height = lines[i][start[2]:stop[2]].strip()
    weight = lines[i][start[3]:stop[3]].strip()
    volume = lines[i][start[4]:stop[4]].strip()
    # Store the data in a dictionary
```

The variables represent one line of data from the file. We want to store it in the dictionary humans as one (key,value) pair.

We want the key to be species and the value to be another dictionary. We can achieve this as follows:

## Putting step 1, 2 and 3 together

```
infile = open('human_evolution.txt', 'r')
lines = infile.readlines()
s = lines[1]
start = [0, s.index('(mill. yrs)'), s.index('height (m)'), ...]
stop = start[1:len(start)] + [80]
\mathbf{k} = 0
while lines[k][0] != '-':
k = k + 1
first = k + 1
k = first
while lines[k][0] != '-':
 k = k + 1
last = k - 1
for i in range(first, last+1):
    species = lines[i][start[0]:stop[0]].strip()
    period = lines[i][start[1]:stop[1]].strip()
    height = lines[i][start[2]:stop[2]].strip()
    weight = lines[i][start[3]:stop[3]].strip()
    volume = lines[i][start[4]:stop[4]].strip()
    humans[species] = {'period': period, 'height': height,
         'weight': weight, 'volume': volume}
```

```
# Print a title
s = '%-23s %-13s %-13s %-13s %-25s' %
        ('species', 'period', 'height', 'weight', 'volume')
print(s)
# Print table contents
for sp in humans:
    d = humans[sp]
    period = d['period']
    height = d['height']
    weight = d['weight']
    volume = d['volume']
    s = '%-23s %-13s %-13s %-25s' % \
        (sp, period, height, weight, volume)
    print(s)
```

sp	ecies	period	height	weight	volume
н.	neanderthalensis	0.35 - 0.03	1.6	55 - 70	1200 - 1700
н.	sapiens sapiens	0.2 - present	1.4 - 1.9	50 - 100	1000 - 1850
н.	heidelbergensis	0.6 - 0.35	1.8	60	1100 - 1400
н.	erectus	1.4 - 0.2	1.8	60	850 (early) - 1100
н.	floresiensis	0.10 - 0.012	1.0	25	400
н.	ergaster	1.9 - 1.4	1.9		700 - 850
н.	habilis	2.2 - 1.6	1.0 - 1.5	33 - 55	660

- We have seen that Python is well suited for mathematical calculations and visualizations.
- Python is also an efficient tool for processing of text strings. \* Applications involving text processing are very common.
- Many advanced applications of text processing (e.g. web search and DNA analysis) involve mathematical and statistical computations.

## Example: web search

Google and other web search tools do advanced text processing. Crawlers browse WWW for files and analyse their content.



## Example: DNA analysis

DNA sequences are very long strings with known and undiscovered patterns. Algorithms to find and compare such patterns are very important in modern biology and medicine.



```
s = 'This is a string, ok?'
# To split a string into individual words:
s.split() # ['This', 'is', 'a', 'string,', 'ok?']
# To split a string with another delimiter
s.split(',') # ['This is a string', ' ok?']
s.split('a string') # ['This is ', ', ok?']
# To find the location of a substring:
s.index('is') # 2
# To check if a string contains a substring:
'This' in s # True
'this' in s # False
# To select a particular character in a string:
s[0] # 'T'
s[1] # 'h'
s[2] # 'i'
s[3] # 's'
```

### Extracting substrings

```
s = 'This is a string, ok?'
# Remove the first character
s[1:] # 'his is a string, ok?'
# Remove the first and the last character
s[1:-1] # 'his is a string, ok'
# Remove the two first and two last characters
s[2:-2] # 'is is a string, o'
# The characters with index 2,3,4
s[2:5] # 'is '
# Select everything starting from a substring
s[s.index('is a'):] # 'is a string, ok?'
# Remove trailing blanks
\mathbf{s} = \mathbf{A} \mathbf{B} \mathbf{C}
```

```
a = ['I', 'am', 'happy']
# Join list elements
''.join(a) # 'Iamhappy'
# Join list elements with space between them
' '.join(a) # 'I am happy'
# Join list elements with '%%' between them
'%%'.join(a) # 'I%%am%%happy'
```

```
s = 'This is a string, ok?'
# Replace every blank by 'X'
s.replace(' ', 'X') # 'ThisXisXaXstring,Xok?'
# Replace one word by another
s.replace('string', 'text') # 'This is a text, ok?'
# Replace the text before the comma by 'Fine'
s.replace(s[:s.index(',')], 'Fine') # 'Fine, ok?'
# Replace the text from the comma by ' dummy'
s.replace(s[s.index(','):], ' dummy') # 'This is a string dummy'
```

Lines are separated by different control characters on different platforms.

```
# Concatenate with Unix/Linux/Mac type line break
s1 = '\n'.join(['Line A', 'Line B', 'Line C'])
# Concatenate with Windows type line break
s2 = '\r\n'.join(['Line A', 'Line B', 'Line C'])
# Platform dependent line splitting:
s1.split('\n') # Works: ['Line Ă', 'Line B', 'Line C']
s1.split('\r\n') # FAILS: ['Line A\nLine B\nLine C']
s2.split('\n') # FAILS: ['Line A\r', 'Line B\r', 'Line C']
s2.split('\r\n') # Works: ['Line A', 'Line B', 'Line C']
# Better line splitting (platform independent):
s1.splitlines() # Works: ['Line A', 'Line B', 'Line C']
s2.splitlines() # Works: ['Line A', 'Line B', 'Line C']
```

```
# Check if a string only contains digits
s = '314'
s.isdigit() # True
s = ' 314'
s.isdigit() # False
s = '3.14'
s.isdigit() # False
# Change to lower-case or upper-case
s = 'ABC def'
s.lower() # 'abc def'
s.upper() # 'ABC DEF'
# Starts with and ends with substring
s = 'This is a string'
s.startswith('This is') # True
s.endswith('This is') # False
```

Suppose we want to read pairs of numbers (x,y) from a file.

Sample file:			
(1.3,0) (0,1) (0,-0.01)	(-1,2) (1,0) (10.5,-1)	(3,-1.5) (1,1) (2.5,-2.5)	

### Algorithm:

- Read one line at a time
- 2 For each line, split line into words
- For each word, strip off parentheses and split the rest on comma

```
infile = open('pairs.dat', 'r')
pairs = []  # Create a list to hold all the pairs
for line in infile:
    words = line.split()
    for w in words:
        w = w[1:-1]  # Remove parentheses
        numbers = w.split(',')
        pair = (float(numbers[0]), float(numbers[1]))
        pairs.append(pair)
```

$$\begin{matrix} [(1.3, 0.0), \\ (-1.0, 2.0), \\ (3.0, -1.5), \\ (0.0, 1.0), \\ (1.0, 0.0), \\ (1.0, 1.0), \\ (0.0, -0.01), \\ (10.5, -1.0), \\ (2.5, -2.5) \end{matrix}$$

```
Suppose the file format
```

```
(1.3, 0) (-1, 2) (3, -1.5)
```

was slightly different:

```
[(1.3, 0), (-1, 2), (3, -1.5),
...]
```

Running eval on the perturbed format produces the desired list!

```
text = open('read_pairs2.dat', 'r').read()
text = '[' + text.replace(')', '),') + ']'
pairs = eval(text)
```

The text is a mix of HTML commands and the text displayed in the browser:

```
<html>
<body bgcolor="orange">
<h1>A Very Simple Web Page</h1> <!-- headline -->
Ordinary text is written as ordinary text, but when we
need headlines, lists,
<111>
<em>emphasized words</em>, or
<b>boldfaced words</b>,
</11]>
we need to embed the text inside HTML tags. We can also
insert GIF or PNG images, taken from other Internet sites,
if desired.
<hr> <!-- horizontal line -->
<img src="http://www.simula.no/simula_logo.gif">
</body>
</html>
```

- A program can download a web page, as an HTML file, and extract data by interpreting the text in the file (using string operations).
- Example: climate data from the UK

Download oxforddata.txt to a local file Oxford.txt:

```
import urllib
baseurl = 'http://www.metoffice.gov.uk/climate/uk/stationdata'
filename = 'oxforddata.txt'
url = baseurl + '/' + filename
urllib.urlretrieve(url, filename='Oxford.txt')
```

Oxford								
Location: 4509E 2072N, 63 metres amsl								
Estimated data is marked with a $*$ after the value.								
Missing	data	(more t	han 2 day:	s missir	ng <mark>in</mark> mo	nth) is	marked by	
Sunshine	data	taken	from an a	utomatic	:			
уууу	mm	tmax	tmin	af	rain	sun		
		degC	degC	days	mm	hours		
1853	1	8.4	2.7	4	62.8			
1853	2	3.2	-1.8	19	29.3			
1853	3	7.7	-0.6	20	25.9			
1853	4	12.6	4.5	0	60.1			
1853	5	16.8	6.1	0	59.5			
2010	5	17.6	7.3	0	28.6	207.4		
2010	6	23.0	11.1	0	34.5	230.5		
2010	7	23.3*	14.1*	0*	24.4*	184.4*	Provisional	
2010	10	14.6	7.4	2	43.5	128.8	Provisional	

## Reading the climate data

### Algorithm:

- Read the place and location in the file header
- 2 Skip the next 5 (for us uninteresting) lines
- Sead the column data and store in dictionary
- Test for numbers with special annotation, "provisional" column, etc.

#### Program, part 1:

#### Program, part 2:

```
data['data'] ={}
for line in infile:
    columns = line.split()
    year = int(columns[0])
    month = int(columns[1])
    if columns[-1] == 'Provisional':
        del columns[-1]
    for i in range(2, len(columns)):
        if columns[i] == '---':
            columns[i] = None
        elif columns[i][-1] == '*' or columns[i][-1] == '#':
            # Strip off trailing character
            columns[i] = float(columns[i][:-1])
        else:
            columns[i] = float(columns[i])
```

#### Program, part 3