Lab 8
Matrices

Sup’Biotech 3
Python

Pierre Parutto

November 23, 2016
Preamble

Document Property

<table>
<thead>
<tr>
<th>Authors</th>
<th>Pierre Parutto</th>
</tr>
</thead>
<tbody>
<tr>
<td>Version</td>
<td>1.0</td>
</tr>
<tr>
<td>Number of pages</td>
<td>5</td>
</tr>
</tbody>
</table>

Contact

Contact the assistant team at: supbiotech-bioinfo-bt3@googlegroups.com

Copyright

The use of this document is strictly reserved to the students from the Sup’Biotech school. This document must have been downloaded from www.intranet.supbiotech.fr, if this is not the case please contact the author(s) at the address given above.

© Assistants Sup’Biotech 2016.
## Contents

1 **Introduction** ........................................... 3

2 **Characteristics of Matrices** .......................... 3
    2.1 Only Negative Values? ............................... 3
        Example ............................................. 3
    2.2 Has negative value? .................................. 3
        Example ............................................. 3
    2.3 Maximum Value ...................................... 3
        Example ............................................. 3
    2.4 Common Values ...................................... 3
        Example ............................................. 4
    2.5 Positions Of Maximum ................................ 4
        Example ............................................. 4

3 **Computations on Matrices** ............................ 4
    3.1 Matrix Transposition ................................. 4
        Example ............................................. 4
    3.2 Matrix Multiplication ................................ 4
        Example ............................................. 5

4 **Creating Matrices** ..................................... 5
    4.1 Count Of Successive Bases ......................... 5
        Example ............................................. 5
1 Introduction

In this lab we will manipulate matrices using the library numpy.

2 Characteristics of Matrices

2.1 Only Negative Values?

Write a function `all_neg(m: array) -> bool` that returns `True` if the matrix `m` contains only strictly negative values and `False` otherwise. If the matrix is empty, the function must return `True`.

Example

```python
>>> from numpy import array
>>> all_neg(array([[-1,-2,-3], [-4,-5,6]]))
False
>>> all_neg(array([[-1,-2,-3], [-3,-2,-1]]))
True
```

2.2 Has negative value?

Write a function `has_neg(m: array) -> bool` that returns `True` if the matrix `m` possesses at least one strictly negative value and `False` otherwise. If `m` is empty the function must return `False`.

Example

```python
>>> from numpy import array
>>> has_neg(array([[1,2,3], [4,5,5]]))
False
>>> has_neg(array([[-1,3], [4,5]]))
True
```

2.3 Maximum Value

Write a function `max_mat(M: array) -> int` that returns the greatest value in the matrix `M`. If `M` is empty your function must return `None`.

Example

```python
>>> from numpy import array
>>> max_mat(array([[1,9], [2,3]]))
9
>>> max_mat(array([[1,2,5], [12,-2,0]]))
12
```

2.4 Common Values

Write a function `common_values(A: array, B: array) -> list` that returns the list of the common values between the two matrices `A` and `B`. The matrices `A` and `B` have the same size.
Example

```python
>>> from numpy import array
>>> common_values(array([[3,-5],[4,8]]), array([[1,2], [3,8]]))
[3,8]
>>> common_values(array([[1,2,3],[-1,4,9]]), array([[3,1,4], [9,8,2]]))
[1,2,3,4,9]
```

2.5 Positions Of Maximum

Write a function `pos_max(M: array) -> list` that returns the positions of the maximum value in the matrix M. The returned value is a list of tuples `[(i,j), ...]` where i is the line number and j is a column number.

Example

```python
>>> from numpy import array
>>> pos_max(array([[1,3],[4,6]]))
[(1,1)]
>>> pos_max(array([[1,2,3],[3,-4,3]]))
[(0,2), (1,0), (1,2)]
```

3 Computations on Matrices

3.1 Matrix Transposition

Transposition is the operation that transforms lines into columns and vice versa. For a matrix $M = m_{i,j}$ of size $m \times n$, its transpose is written as: $M^t$, has size $n \times m$ and follows the formula:

$$
M^t_{i,j} = m_{j,i}
$$

Write a function `transpose(M: array) -> array` that returns the transpose of the matrix M.

Example

```python
>>> from numpy import array
>>> transpose(array([[1,2],[3,4],[5,6]]))
array([[1,3,5],
      [2,4,6]])
>>> transpose(array([[1,2,3],[4,5,6]]))
array([[1,4],
      [2,5],
      [3,6]])
```

3.2 Matrix Multiplication

The multiplication between two compatible matrices $A = a_{i,j}$ of size $m \times n$ and $B = b_{i,j}$ of size $n \times k$ produces a matrix $C = c_{i,j}$ of size $m \times k$ such that $\forall 0 < i < n$ and $0 < j < k$:

$$
c_{i,j} = \sum_{i=0}^{n} a_{i,k} b_{k,j}
$$

Write a function `mat_mult(A: array, B: array) -> array` that returns the multiplication between the two matrices A and B.
Example

```python
>>> from numpy import array
>>> mat_mult(array([[1,2],[4,5]]), array([[1,0],[0,1]]))
array([[1,2],[4,5]])
>>> mat_mult(array([[-1,-2],[-4,-5],[-3,-6]]), array([[1,2,3],[4,5,6]]))
array([[-9,-12,-15],[-24,-33,-42],[-27,-36,-45]])
```

4 Creating Matrices

4.1 Count Of Successive Bases

Consider a DNA sequence, we want to count at which a base pair $b_1$ is followed by a base pair $b_2$. We will create a matrix where the lines correspond to the base $b_1$ and the columns to the base $b_2$. As there are 4 bases, A, T, G, C, the matrix will be of size $4 \times 4$, where we consider the mapping:

- A $\rightarrow$ 0
- T $\rightarrow$ 1
- G $\rightarrow$ 2
- C $\rightarrow$ 3

Write a function `count_succ(s: str) -> array` that returns the matrix presented above given the string $s$.

Example

```python
>>> count_succ("ATTGTGACT")
array([[0,1,0,1],[0,1,2,0],[1,1,0,0],[0,1,0,0]])
>>> count_succ("AAAAATGAGTA")
array([[4,1,1,0],[1,0,1,0],[1,1,0,0],[0,0,0,0]])
```