

## Creating\_Functions\_part\_2

May 8, 2023

```
[17]: import numpy
import glob
import matplotlib
import matplotlib.pyplot
```

```
[21]: def visualize(filename):

    data = numpy.loadtxt(fname = filename, delimiter = ',')

    fig = matplotlib.pyplot.figure(figsize=(10.0, 3.0))

    axes1 = fig.add_subplot(1, 3, 1)
    axes2 = fig.add_subplot(1, 3, 2)
    axes3 = fig.add_subplot(1, 3, 3)

    axes1.set_ylabel('average')
    axes1.plot(numpy.mean(data, axis=0))

    axes2.set_ylabel('max')
    axes2.plot(numpy.amax(data, axis=0))

    axes3.set_ylabel('min')
    axes3.plot(numpy.amin(data, axis=0))

    fig.tight_layout()
    matplotlib.pyplot.show()
```

```
[22]: def detect_problems(filename):

    data = numpy.loadtxt(fname = filename, delimiter = ',')

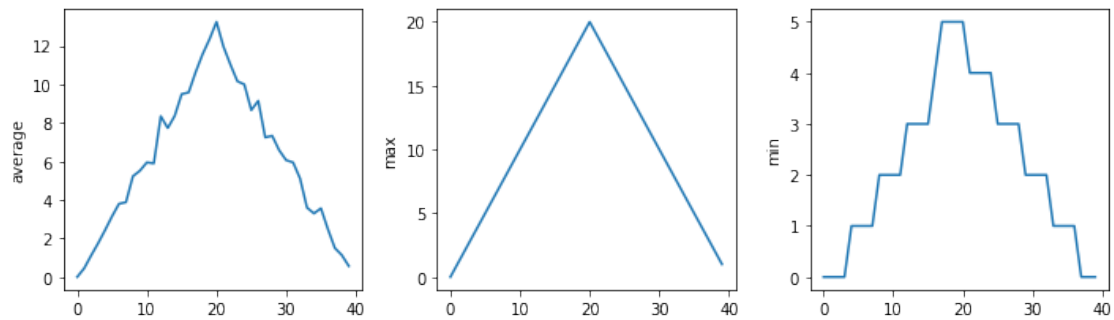
    if numpy.amax(data, axis = 0)[0] == 0 and numpy.amax(data, axis = 0)[20] == 0:
        print('Suspicious looking maxima!')
    elif numpy.sum(numpy.amin(data, axis = 0)) == 0:
        print('Minima add up to zero!')
    else:
```

```
print('seems ok!')
```

```
[25]: filenames = sorted(glob.glob('inflammation*.csv'))

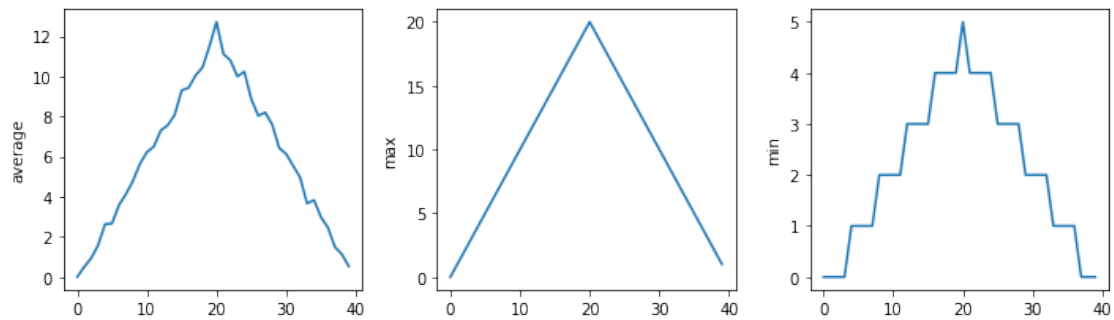
for filename in filenames:
    print(filename)
    visualize(filename)
    detect_problems(filename)
```

inflammation-01.csv



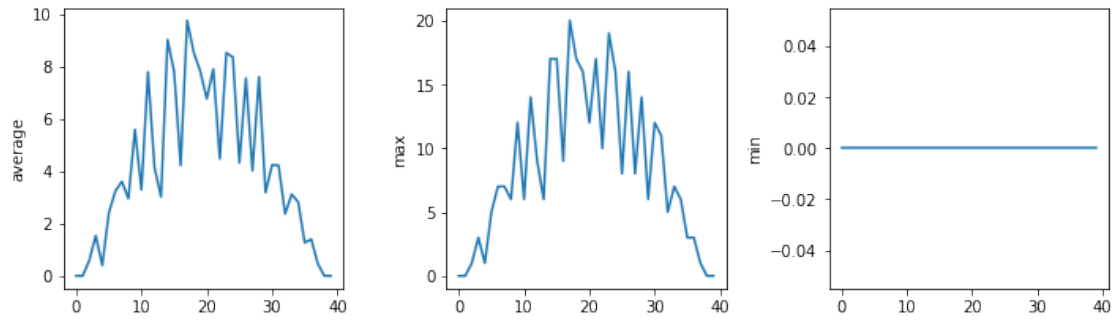
Suspicious looking maxima!

inflammation-02.csv

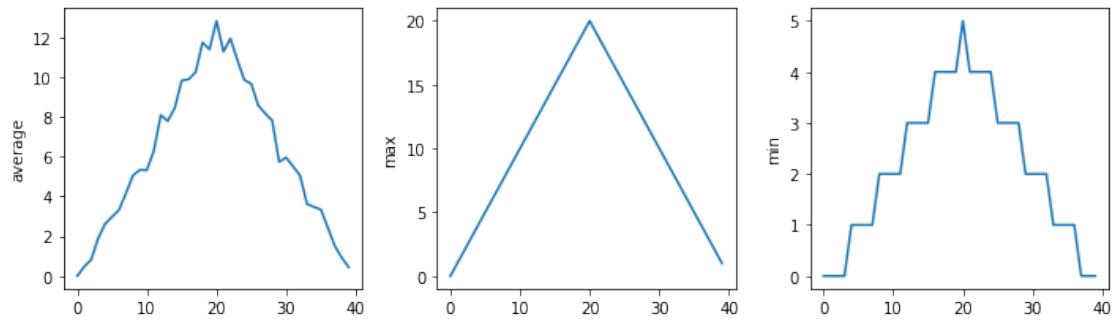


Suspicious looking maxima!

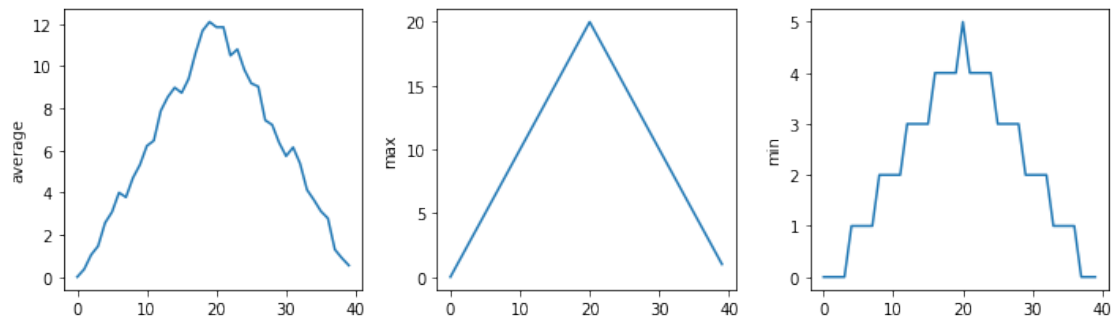
inflammation-03.csv



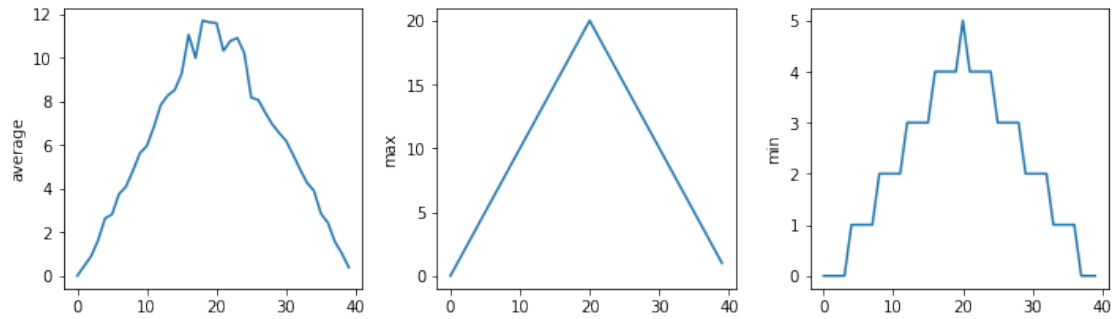
Minima add up to zero!  
inflammation-04.csv



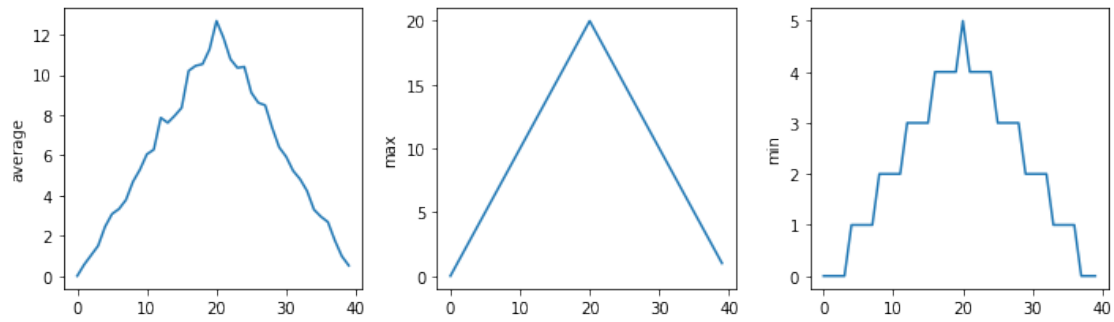
Suspicious looking maxima!  
inflammation-05.csv



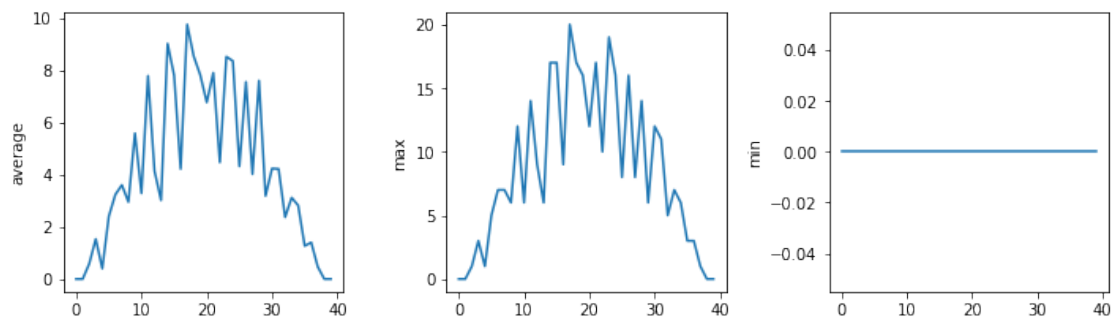
Suspicious looking maxima!  
inflammation-06.csv



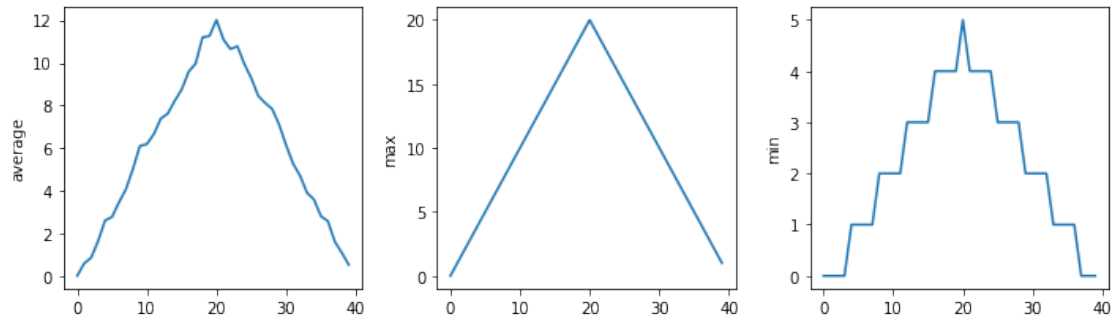
Suspicious looking maxima!  
inflammation-07.csv



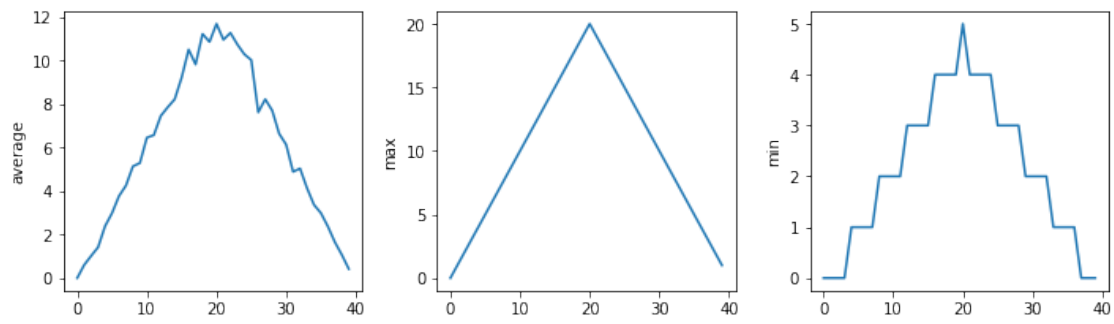
Suspicious looking maxima!  
inflammation-08.csv



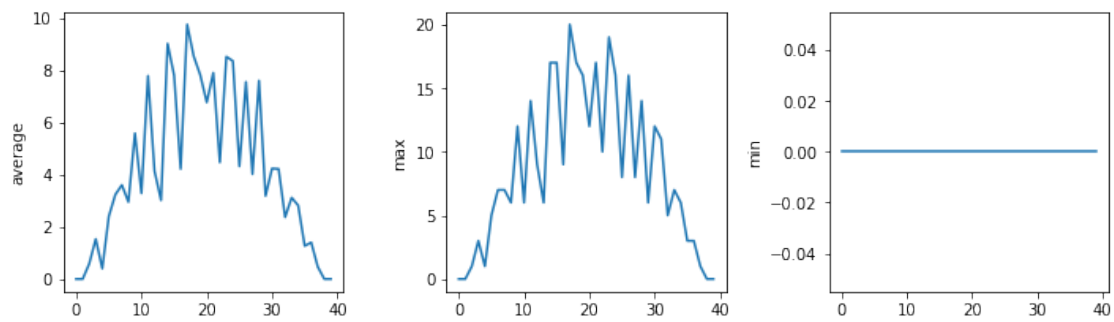
Minima add up to zero!  
inflammation-09.csv



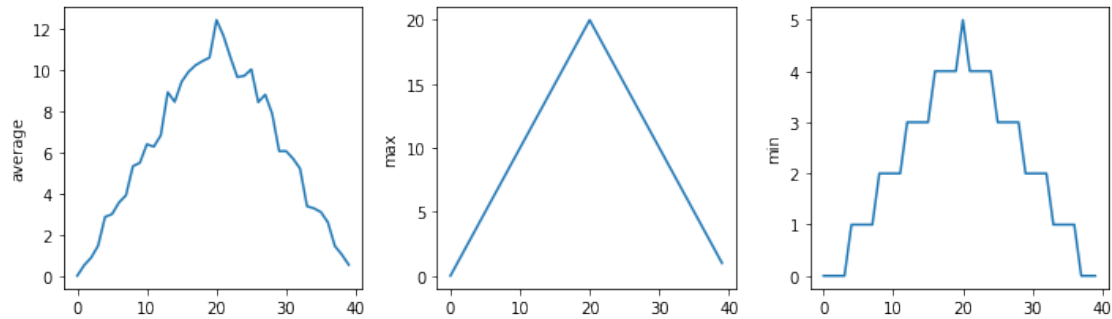
Suspicious looking maxima!  
inflammation-10.csv



Suspicious looking maxima!  
inflammation-11.csv



Minima add up to zero!  
inflammation-12.csv



Suspicious looking maxima!

```
[29]: def offset_mean(data, target_mean_value):
      return(data - numpy.mean(data)) + target_mean_value
```

```
[30]: z = numpy.zeros((2,2))
      print(offset_mean(z, 3))
```

```
[[3. 3.]
 [3. 3.]]
```

```
[32]: data = numpy.loadtxt(fname = 'inflammation-01.csv', delimiter = ',')

      print(offset_mean(data, 0))
```

```
[[ -6.14875  -6.14875  -5.14875  ...  -3.14875  -6.14875  -6.14875]
 [ -6.14875  -5.14875  -4.14875  ...  -5.14875  -6.14875  -5.14875]
 [ -6.14875  -5.14875  -5.14875  ...  -4.14875  -5.14875  -5.14875]
 ...
 [ -6.14875  -5.14875  -5.14875  ...  -5.14875  -5.14875  -5.14875]
 [ -6.14875  -6.14875  -6.14875  ...  -6.14875  -4.14875  -6.14875]
 [ -6.14875  -6.14875  -5.14875  ...  -5.14875  -5.14875  -6.14875]]
```

```
[33]: print('original mn, mean and max are:', numpy.amin(data), numpy.mean(data),
      ↪ numpy.amax(data))
      offset_data = offset_mean(data, 0)
      print('min, mean, and max of offset data are:',
            numpy.amin(offset_data),
            numpy.mean(offset_data),
            numpy.amax(offset_data))
```

```
original mn, mean and max are: 0.0 6.14875 20.0
```

```
min, mean, and max of offset data are: -6.14875 2.842170943040401e-16 13.85125
```

```
[34]: print('std dev before and after:', numpy.std(data), numpy.std(offset_data))
```

std dev before and after: 4.613833197118566 4.613833197118566

```
[35]: print('difference in standard deviation before and after:',  
        numpy.std(data)- numpy.std(offset_data))
```

difference in standard deviation before and after: 0.0

```
[36]: # offset_mean(data, target_mean_value):  
# return a new array containing the original data with its mean offset to match  
# the desired value  
def offset_mean(data, target_mean_value):  
    return(data - numpy.mean(data)) + target_mean_value
```

```
[37]: def offset_mean(data, target_mean_value):  
    """Return a new array containing the original data with its mean offset to  
    match the desired value"""  
    return(data - numpy.mean(data)) + target_mean_value
```

```
[38]: help(offset_mean)
```

Help on function offset\_mean in module \_\_main\_\_:

offset\_mean(data, target\_mean\_value)

Return a new array containing the original data with its mean offset to match the desired value

```
[39]: def offset_mean(data, target_mean_value):  
    """Return a new array containing the original data with its mean offset to  
    match desired value.  
  
    Examples  
    -----  
  
    >>>offset_mean([1,2,3],0)  
    array([-1., 0., 1.])  
    """  
    return (data - numpy.mean(data)) + target_mean_value
```

```
[40]: help(offset_mean)
```

Help on function offset\_mean in module \_\_main\_\_:

offset\_mean(data, target\_mean\_value)

Return a new array containing the original data with its mean offset to match desired value.

Examples

-----

```
>>>offset_mean([1,2,3],0)
array([-1., 0., 1.]
```

```
[41]: numpy.loadtxt('inflammation-01.csv', delimiter = ',')
```

```
[41]: array([[0., 0., 1., ..., 3., 0., 0.],
            [0., 1., 2., ..., 1., 0., 1.],
            [0., 1., 1., ..., 2., 1., 1.],
            ...,
            [0., 1., 1., ..., 1., 1., 1.],
            [0., 0., 0., ..., 0., 2., 0.],
            [0., 0., 1., ..., 1., 1., 0.]])
```

```
[42]: def offset_mean(data, target_mean_value = 0.0):
      """Return a new array containing the original data with its mean offset to_
      ↪match desired value, (0 by default).
```

```
Examples
-----
```

```
>>>offset_mean([1,2,3])
array([-1., 0., 1.]
      """
      return (data - numpy.mean(data)) + target_mean_value
```

```
[43]: test_data = numpy.zeros((2,2))
      print(offset_mean(test_data, 3))
```

```
[[3. 3.]
 [3. 3.]]
```

```
[44]: print(offset_mean(test_data))
```

```
[[0. 0.]
 [0. 0.]]
```

```
[45]: def display(a=1, b=2, c=3):
      print('a:', a, 'b:', b, 'c:', c)

      print('no parameters:')
      display()
      print('one parameter:')
      display(55)
      print('two parameters')
      display(55,66)
```



```
no parameters:
a: 1 b: 2 c: 3
one parameter:
a: 55 b: 2 c: 3
two parameters
a: 55 b: 66 c: 3
```

```
[46]: print('only setting the value of c')
      display(c = 77)
```

```
only setting the value of c
a: 1 b: 2 c: 77
```

```
[47]: help(numpy.loadtxt)
```

Help on function loadtxt in module numpy:

```
loadtxt(fname, dtype=<class 'float'>, comments='#', delimiter=None,
converters=None, skiprows=0, usecols=None, unpack=False, ndmin=0,
encoding='bytes', max_rows=None)
    Load data from a text file.
```

Each row in the text file must have the same number of values.

Parameters

-----

**fname** : file, str, or pathlib.Path

File, filename, or generator to read. If the filename extension is ```.gz``` or ```.bz2```, the file is first decompressed. Note that generators should return byte strings for Python 3k.

**dtype** : data-type, optional

Data-type of the resulting array; default: float. If this is a structured data-type, the resulting array will be 1-dimensional, and each row will be interpreted as an element of the array. In this case, the number of columns used must match the number of fields in the data-type.

**comments** : str or sequence of str, optional

The characters or list of characters used to indicate the start of a comment. None implies no comments. For backwards compatibility, byte strings will be decoded as 'latin1'. The default is '#'.

**delimiter** : str, optional

The string used to separate values. For backwards compatibility, byte strings will be decoded as 'latin1'. The default is whitespace.

**converters** : dict, optional

A dictionary mapping column number to a function that will parse the column string into the desired value. E.g., if column 0 is a date string: ```converters = {0: datestr2num}```. Converters can also be used to provide a default value for missing data (but see also

```

    `genfromtxt`): ``converters = {3: lambda s: float(s.strip() or 0)}``.
    Default: None.
skiprows : int, optional
    Skip the first `skiprows` lines, including comments; default: 0.
usecols : int or sequence, optional
    Which columns to read, with 0 being the first. For example,
    ``usecols = (1,4,5)`` will extract the 2nd, 5th and 6th columns.
    The default, None, results in all columns being read.

.. versionchanged:: 1.11.0
    When a single column has to be read it is possible to use
    an integer instead of a tuple. E.g ``usecols = 3`` reads the
    fourth column the same way as ``usecols = (3,)`` would.
unpack : bool, optional
    If True, the returned array is transposed, so that arguments may be
    unpacked using ``x, y, z = loadtxt(...)``. When used with a structured
    data-type, arrays are returned for each field. Default is False.
ndmin : int, optional
    The returned array will have at least `ndmin` dimensions.
    Otherwise mono-dimensional axes will be squeezed.
    Legal values: 0 (default), 1 or 2.

.. versionadded:: 1.6.0
encoding : str, optional
    Encoding used to decode the inputfile. Does not apply to input streams.
    The special value 'bytes' enables backward compatibility workarounds
    that ensures you receive byte arrays as results if possible and passes
    'latin1' encoded strings to converters. Override this value to receive
    unicode arrays and pass strings as input to converters. If set to None
    the system default is used. The default value is 'bytes'.

.. versionadded:: 1.14.0
max_rows : int, optional
    Read `max_rows` lines of content after `skiprows` lines. The default
    is to read all the lines.

.. versionadded:: 1.16.0

Returns
-----
out : ndarray
    Data read from the text file.

See Also
-----
load, fromstring, fromregex
genfromtxt : Load data with missing values handled as specified.
scipy.io.loadmat : reads MATLAB data files

```

## Notes

-----

This function aims to be a fast reader for simply formatted files. The ``genfromtxt`` function provides more sophisticated handling of, e.g., lines with missing values.

.. versionadded:: 1.10.0

The strings produced by the Python `float.hex` method can be used as input for floats.

## Examples

-----

```
>>> from io import StringIO # StringIO behaves like a file object
>>> c = StringIO(u"0 1\n2 3")
>>> np.loadtxt(c)
array([[0., 1.],
       [2., 3.]])

>>> d = StringIO(u"M 21 72\nF 35 58")
>>> np.loadtxt(d, dtype={'names': ('gender', 'age', 'weight'),
...                          'formats': ('S1', 'i4', 'f4')})
array([(b'M', 21, 72.), (b'F', 35, 58.)],
      dtype=[('gender', 'S1'), ('age', '<i4'), ('weight', '<f4')])

>>> c = StringIO(u"1,0,2\n3,0,4")
>>> x, y = np.loadtxt(c, delimiter=',', usecols=(0, 2), unpack=True)
>>> x
array([1., 3.])
>>> y
array([2., 4.]])
```

```
[48]: numpy.loadtxt('inflammation-01.csv', delimiter = ',')
```

```
[48]: array([[0., 0., 1., ..., 3., 0., 0.],
            [0., 1., 2., ..., 1., 0., 1.],
            [0., 1., 1., ..., 2., 1., 1.],
            ...,
            [0., 1., 1., ..., 1., 1., 1.],
            [0., 0., 0., ..., 0., 2., 0.],
            [0., 0., 1., ..., 1., 1., 0.]])
```

```
[ ]:
```