

```
In [1]: import sys
from sklearn.neighbors import KernelDensity

sys.path.insert(0, "gudhi-devel-multi/build/src/python/gudhi/")
import gudhi as gd
import numpy as np
import matplotlib.pyplot as plt
```

# From scratch

Definition of a simplextree

```
In [2]: simplextree = gd.SimplexTreeMulti(num_parameters=2)
simplextree.insert([0,1,2], [0,0])          # we add the triangle 0-1-2 to the complex.

#    0---2
#    | */
#    | */
#    |/
#    1
```

Out[2]: True

```
In [3]: simplextree.num_parameters
```

Out[3]: 2

Filtration assignment

```
In [4]: #simplextree.assign_filtration(simplex, bifiltration value)
simplextree.assign_filtration([0], np.array([1,2], dtype=int)) # The vertices appear
simplextree.assign_filtration([1], np.array([1,2], dtype=np.float128))
simplextree.assign_filtration([2], [1,2])
simplextree.assign_filtration([0,1], [1,2]) # The edges (and the cycle) appear at [1,
simplextree.assign_filtration([0,2], [1,2])
simplextree.assign_filtration([1,2], [1,2])
simplextree.assign_filtration([0,1,2], [3,3]) # The triangle appears at [3,3], and ki
```

```
In [5]: simplextree2 = gd.SimplexTreeMulti(num_parameters = 2)
simplextree2.insert([0,1,2], [0,0])
for s,f in simplextree2.get_skeleton(1):
    simplextree2.assign_filtration(s, [1,2])
simplextree2.assign_filtration([0,1,2], [3,3])
print(simplextree == simplextree2)
simplextree2.assign_filtration([0], [1,1])
print(simplextree == simplextree2)
```

True

False

Multi-critical filtrations

```
In [6]: simplextree.filtration([0])
```

Out[6]: [1.0, 2.0]

```
In [7]: simplextree.insert([0], [1,3]) # (1,3) >= (1,2), simplex already appeared
```

Out[7]: False

```
In [8]: # (2, 1) is incomparable to previous filtrations, it can be added as a new birth  
simplextree.insert([0], [2,1])
```

```
Out[8]: True
```

```
In [9]: # Vector on which all filtration values of this simplex are stacked.
```

```
f = simplextree.filtration([0])
```

```
np.array_split(f, len(f) // simplextree.num_parameters) # They can be unstack using a
```

```
Out[9]: [array([1., 2.]), array([2., 1.])]
```

```
In [10]: ## Prevents assigning filtration values
```

```
# that don't have the same shape as self.num_parameter
```

```
try:
```

```
    simplextree.assign_filtration([0],[1,1,1])
```

```
except AssertionError:
```

```
    print("len([1,1,1]) % 2 != 0")
```

```
    # I don't know how to make the assert message print with cython
```

```
len([1,1,1]) % 2 != 0
```

## Example of (useful) bifiltration : Rips + Density

```
In [11]: def noisy_annulus(r1:float=1, r2:float=2, n1:int=1000,n2:int=200, dim:int=2, center:n  
        """Generates a noisy annulus dataset.
```

```
Parameters
```

```
-----
```

```
r1 : float.
```

```
    Lower radius of the annulus.
```

```
r2 : float.
```

```
    Upper radius of the annulus.
```

```
n1 : int
```

```
    Number of points in the annulus.
```

```
n2 : int
```

```
    Number of points in the square.
```

```
dim : int
```

```
    Dimension of the annulus.
```

```
center: list or array
```

```
    center of the annulus.
```

```
Returns
```

```
-----
```

```
numpy array
```

```
    Dataset. size : (n1+n2) x dim
```

```
"""
```

```
from numpy.random import uniform
```

```
from numpy.linalg import norm
```

```
set =[]
```

```
while len(set)<n1:
```

```
    draw=uniform(low=-r2, high=r2, size=dim)
```

```
    if norm(draw) > r1 and norm(draw) < r2:  
        set.append(draw)
```

```
annulus = np.array(set) if center == None else np.array(set) + np.array(cente
```

```
diffuse_noise = uniform(size=(n2,dim), low=-1.1*r2,high=1.1*r2)
```

```
if center is not None: diffuse_noise += np.array(center)
```

```
return np.vstack([annulus, diffuse_noise])
```

```
In [12]: npts = 200
```

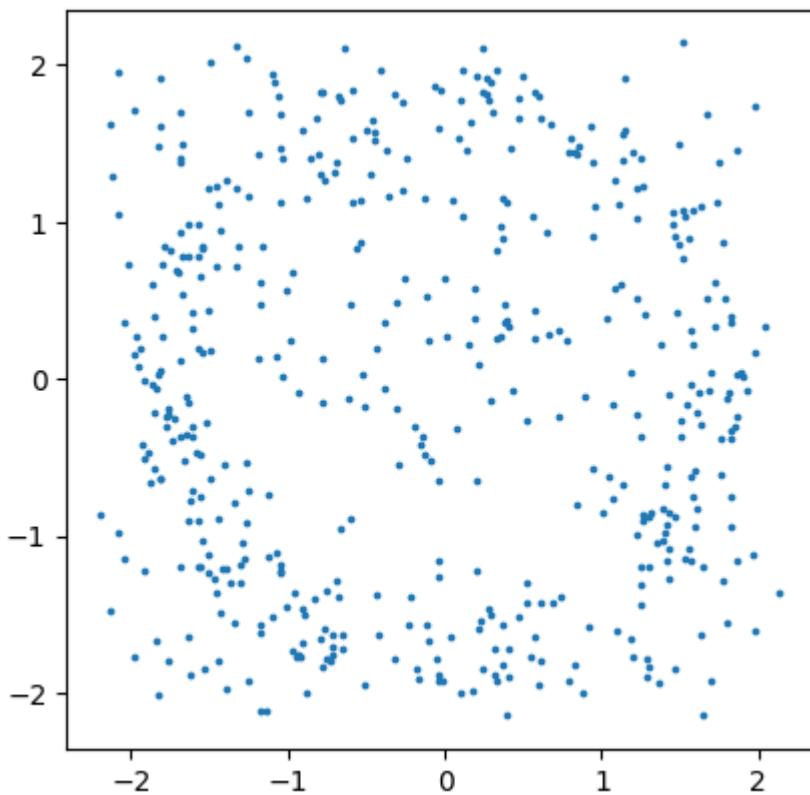
```
noutliers = (int)(npts * 0.4)
```

```
np.random.seed(100)
```

```

points = np.block([[np.array(noisy_annulus(1.5,2, npts))], [np.random.uniform(low=-2,
fig, ax = plt.subplots()
ax.set_aspect('equal')
plt.scatter(points[:,0], points[:,1], s=3)
plt.show()

```



```

In [13]: simplextree = gd.RipsComplex(points = points).create_simplex_tree(max_dimension=1)
## Creates a SimplexTreeMulti from a standard one. The first filtration is the same
simplextree = gd.SimplexTreeMulti(simplextree, num_parameters=2)
simplextree.num_simplices()

```

Out[13]: 115440

```

In [14]: kde = KernelDensity(bandwidth=0.25)
density = kde.fit(points).score_samples(points)
simplextree.fill_lowerstar(-density, parameter=1)

```

Out[14]: <gudhi.simplex\_tree\_multi.SimplexTreeMulti at 0x7f11ba922810>

```

In [15]: # Example of filtration
print(next(simplextree.get_simplices()))
([0, 1], [0.351698269996502, 2.6640137794676924])

```

```

In [16]: # Edge collapses from filtration_domination
simplextree.collapse_edges(strong=True, num=100, progress=1) # Takes time to get an e
simplextree.collapse_edges(strong=False, num=100, progress=1)
simplextree.expansion(2)
simplextree.num_simplices()

```

Removing edges: 13%	███████████	13/100 [00:02<00:17, 4.84it/s]
Removing edges: 11%	██████████	11/100 [00:01<00:14, 6.09it/s]

Out[16]: 7791

```

In [17]: ## The eulerchar of a list of points of a simplextree can be computed using this meth
# NOTE: Makes no sense on rips complexes, as they are cut in dimension.
# I have another implementation, relying on computing the rank invariant on a grid.
simplextree.euler_char([[0.1,2.5], [1,2.2]])

```

```
Out[17]: array([54,  6])
```