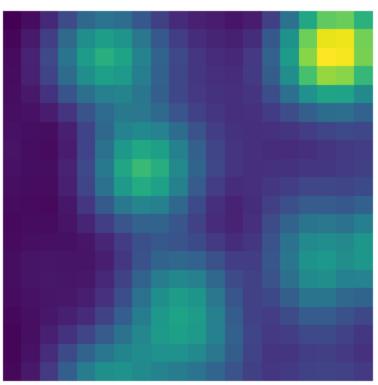
Persistent Homology (dim=0)

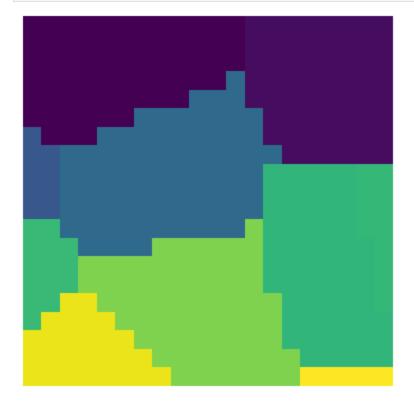
```
input: image I with shape (H,W)
           output: dgm array with birth and death values
           1. M = arg-maxpool2d(I)
           2. pbirth = unique(M), M = reindex(M)
           3. B = maxpool(M) != -maxpool(-M)
           4. B = distillation(B)
           5. B = sort(B)
              For x in B:
                if x is border of 2 partitions P:
                   x is added in pdeath
                   C[x] = min(P)
           6. dgm = (I[pbirth], I[pdeath])
        import numpy as np
In [1]:
        from astropy.io import fits
        import matplotlib.pyplot as plt
        from utils import maxpool2d, neighbors, my saddle
In [2]: img = fits.getdata("")
        img = img[0:20,0:20].astype(np.float32)
        img -= img.min()
        img /= img.max()
In [3]:
        plt.imshow(img)
        plt.axis('off')
        plt.show()
```



1. Identification of the concave components

The initial step involves identifying the concave elements within the image I. To accomplish this, we compute, for each pixel, the index of the pixel with the highest value among its eight neighboring pixels, using the arg-maxpool2d(x) operation. The generated index array is recursively processed until a labeled image M is achieved.

```
In [5]: plt.imshow(m.reshape(H,W))
   plt.axis('off')
   plt.show()
```



2. Locate birth points (Re-Indexing of components)

Currently, the detected elements are labeled with the index of their neighbor relative maximum. We gather all the indices of relative maxima into an array pbirth. Before moving on, we assign sequential numbers to the zones, starting from the one with the smallest relative maximum to the one with the largest one.

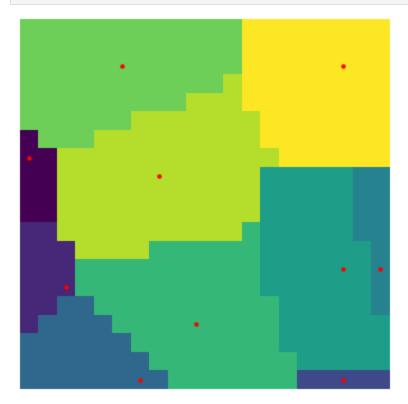
```
In [6]: pbirth = np.unique(m)
  birth = img[pbirth]
  idxs = np.argsort(birth)
  pbirth = pbirth[idxs]
```

```
In [7]: dict_replace = {pbirth[i]: i+1 for i in range(birth.size)}
p_idxs = np.sort(idxs + 1)
```

```
def replace(element):
    return dict_replace.get(element, element)
vreplace = np.vectorize(replace)
new_m = vreplace(m)

del birth, idxs, dict_replace, vreplace
```

```
In [8]: plt.imshow(new_m.reshape(H,W))
plt.scatter(pbirth % W, pbirth // W, 5, 'r')
plt.axis('off')
plt.show()
```



3. Find the edges

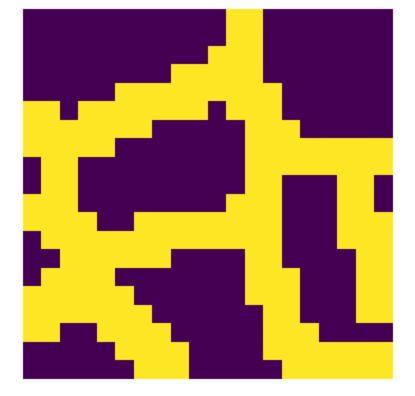
After have partitioned the image into distinct components, we detect the edges, and within the found edges we later locate the dead points pdeath.

To detect the edges, we calculate the maxpool2d of M and -M. The area of our interest will be the region B where these two arrays differ.

```
In [9]: p1 = maxpool2d(new_m.reshape(H, W), kernel_size=3, stride=1, padding=1, return_indices=F
    p2 = -maxpool2d(-new_m.reshape(H, W), kernel_size=3, stride=1, padding=1, return_indices
    borders_idxs = set(np.nonzero(p1 != p2)[0].tolist())
```

```
In [10]: mask = np.zeros((H,W)).flatten()
    mask[list(borders_idxs)] = 1
    mask = mask.reshape(H,W)

plt.imshow(mask)
    plt.axis('off')
    plt.show()
```



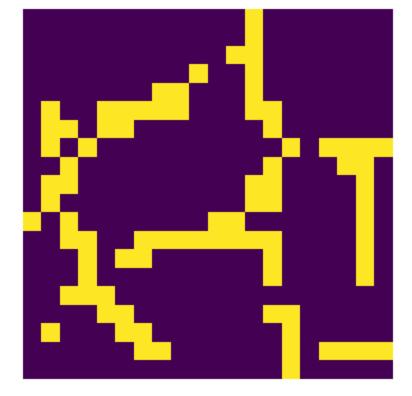
4. Distillation

By definition, dead points are located along border regions. Moreover, to connect two adjacent areas, they must be either relative minimum points or saddle points. Any point failing to meet these criteria is subsequently removed from the previous array B.

```
img pad = np.pad(img.reshape(H,W), ((1, 1), (1, 1)), 'constant', constant values=0)
In [11]:
         for x in borders idxs.copy():
            w = x // W
            h = x % W
            if not my saddle(img pad[w:(w+3), h:(h+3)]):
                 borders idxs.discard(x)
         for x in np.flip(pbirth):
             for y in neighbors (x, H, W):
                 if new m[y] != new m[x]:
                    borders idxs.add(y)
            borders idxs.discard(x)
         del img pad
        borders idxs = np.array(list(borders idxs))
         sort = np.flip(np.argsort(img[borders idxs]))
        borders idxs = borders idxs[sort]
```

```
In [12]: mask2 = np.zeros((H,W)).flatten()
   mask2[borders_idxs] = 1
   mask2 = mask2.reshape(H,W)

plt.imshow(mask2)
  plt.axis('off')
  plt.show()
```

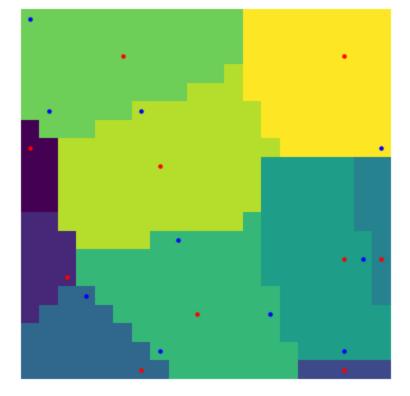


5. Locate dead points

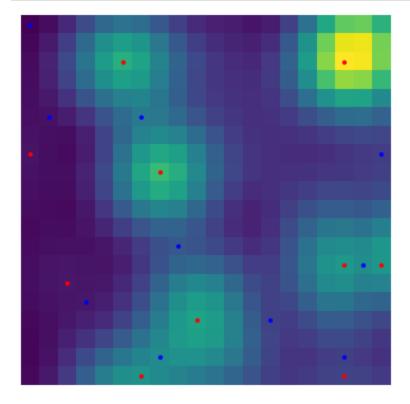
The potential death points are arranged in descending order to preserve the sequence of merging partitions in M. For each point x in set B, if it borders two partitions, these partitions are merged, and x is designated as the pdeath point for the partition with the smallest index. The history of these unions is tracked in vector C, which holds the new index in relation to the initial partition index. To enhance the algorithm's efficiency, modifications are restricted to the eight pixels close to the point x, rather than applying changes directly to M

```
pdeath = []
In [13]:
         death idxs = set()
         changer = p idxs.copy()
         for x in borders idxs:
             check = np.unique(changer[new m[np.array(neighbors(x, H, W, mode=8))] -1])
             if len(check) >= 2:
                 check = tuple(check[:2])
                 if check not in death idxs:
                     death idxs.add(check)
                     changer[np.where(changer == check[0])[0]] = check[1]
                     pdeath.append([check[0] -1, x])
         del changer
         pdeath = np.array(pdeath)
         pdeath = pdeath[np.argsort(pdeath[:,0])][:,1]
         pdeath = np.append(pdeath, np.argmin(img))
```

```
In [14]: plt.imshow(new_m.reshape(H,W))
  plt.scatter(pbirth % W, pbirth // W, 5, 'r')
  plt.scatter(pdeath % W, pdeath // W, 5, 'b')
  plt.axis('off')
  plt.show()
```



```
In [15]: plt.imshow(img.reshape(H,W))
  plt.scatter(pbirth % W, pbirth // W, 5, 'r')
  plt.scatter(pdeath % W, pdeath // W, 5, 'b')
  plt.axis('off')
  plt.show()
```



6. Persistence Diagram (DGM)

In the final step, a persistence diagram is created, linking each birth point with the corresponding relative death point within the same partition. The birth and death values for the pbirth and pdeath points are then extracted from the image I and aggregated into a DGM matrix.

```
In [16]:
                         img[pdeath]], axis=1)
         dam
         array([[0.05277771, 0.04943267],
Out[16]:
                [0.0656876, 0.06317271],
                [0.20702817, 0.18857712],
                [0.51076263, 0.4814553],
                [0.5293757, 0.49486426],
                [0.5322356, 0.19706908],
                [0.58093137, 0.2759763],
                [0.6282401, 0.40609908],
                [0.6722756, 0.18125333],
                            , 0.
                                        ]], dtype=float32)
In [17]: plt.figure(figsize=(5,5))
         plt.scatter(dgm[:,1], dgm[:,0])
         plt.plot([0, 1], [0, 1], 'k-', lw=1)
         plt.xlim([0,1])
         plt.ylim([0,1])
         plt.show()
          1.0
          0.8
          0.6
          0.4
          0.2
          0.0
                       0.2
                                 0.4
            0.0
                                           0.6
                                                      0.8
                                                                1.0
In [17]:
In [17]:
         f, arr = plt.subplots(2,4, figsize=(40,20))
In [18]:
         arr[0,0].imshow(img.reshape(H,W))
         arr[0,0].axis('off')
         arr[0,0].set title('a', y=-0.1, fontsize=36)
         arr[0,1].imshow(new m.reshape(H,W))
         arr[0,1].axis('off')
         arr[0,1].set_title('b', y=-0.1, fontsize=36)
```

dgm = np.stack([img[pbirth],

arr[0,2].imshow(new m.reshape(H,W))

```
arr[0,2].scatter(pbirth % W, pbirth // W, 25, 'r')
arr[0,2].axis('off')
arr[0,2].set title('c', y=-0.1, fontsize=36)
arr[0,3].imshow(mask.reshape(H,W))
arr[0,3].axis('off')
arr[0,3].set title('d', y=-0.1, fontsize=36)
arr[1,0].imshow(mask2.reshape(H,W))
arr[1,0].axis('off')
arr[1,0].set title('e', y=-0.1, fontsize=36)
arr[1,1].imshow(new m.reshape(H,W))
arr[1,1].scatter(pbirth % W, pbirth // W, 25, 'r')
arr[1,1].scatter(pdeath % W, pdeath // W, 25, 'b')
arr[1,1].axis('off')
arr[1,1].set title('f', y=-0.1, fontsize=36)
arr[1,2].imshow(img.reshape(H,W))
arr[1,2].scatter(pbirth % W, pbirth // W, 25, 'r')
arr[1,2].scatter(pdeath % W, pdeath // W, 25, 'b')
arr[1,2].axis('off')
arr[1,2].set title('g', y=-0.1, fontsize=36)
arr[1,3].scatter(dgm[:,1], dgm[:,0])
arr[1,3].plot([0, 1], [0, 1], 'k-', lw=1)
arr[1,3].set xlim([0,1])
arr[1,3].set ylim([0,1])
arr[1,3].set title('h', y=-0.1, fontsize=36)
plt.show()
                                b
        а
                                                        c
                                                        g
```