Objects and Labels

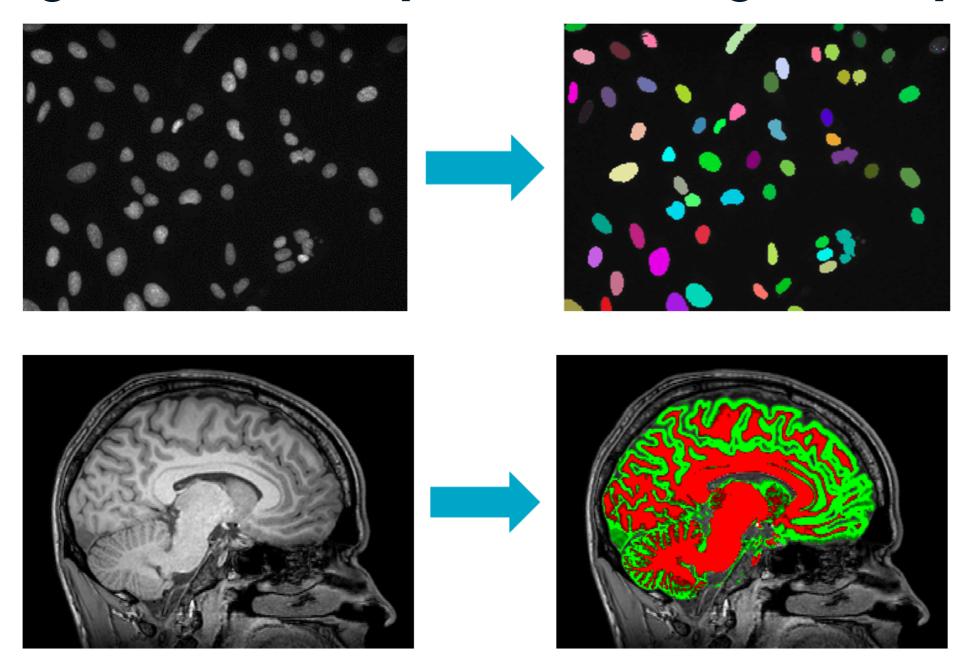
BIOMEDICAL IMAGE ANALYSIS IN PYTHON



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Instructor

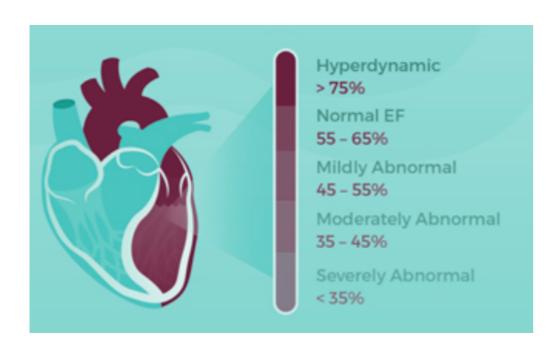


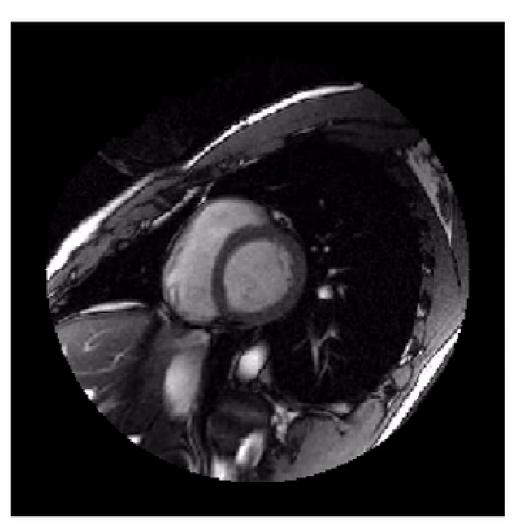
Segmentation splits an image into parts



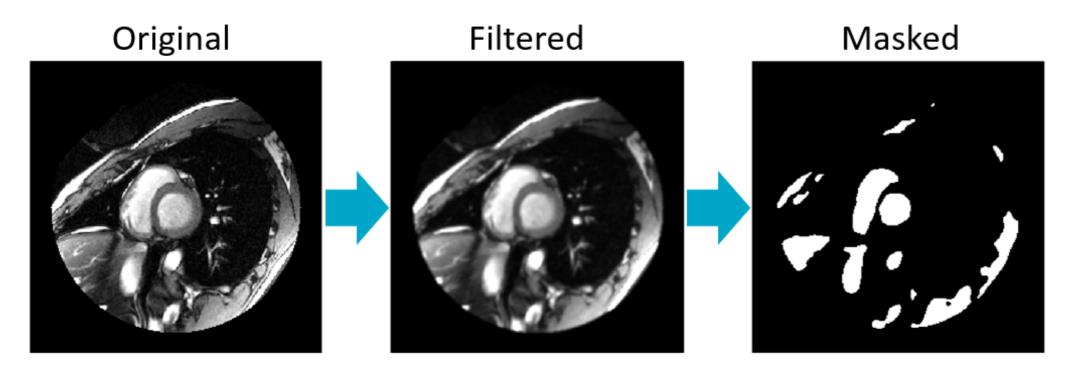
Sunnybrook Cardiac Database

Ejection fraction: the proportion of blood pumped out of the heart's left ventricle (LV).





Labeling image components

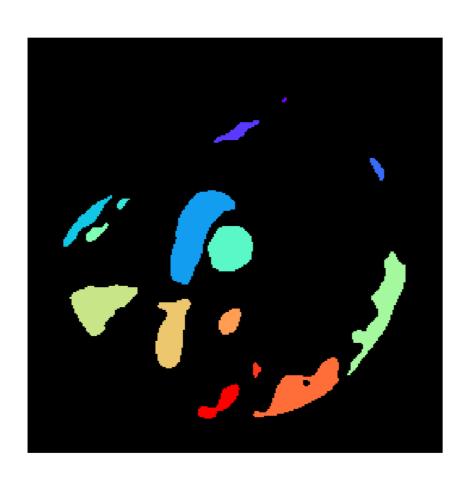


Labeling image components

nlabels

14

```
plt.imshow(labels, cmap='rainbow')
plt.axis('off')
plt.show()
```



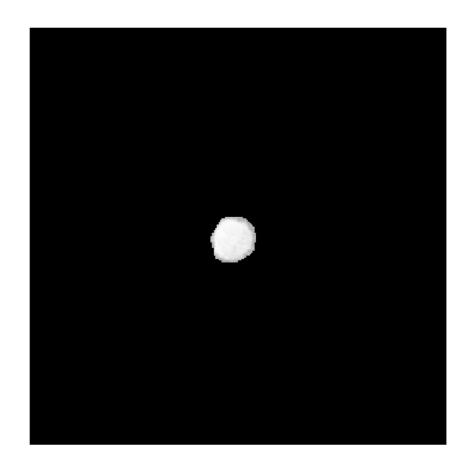


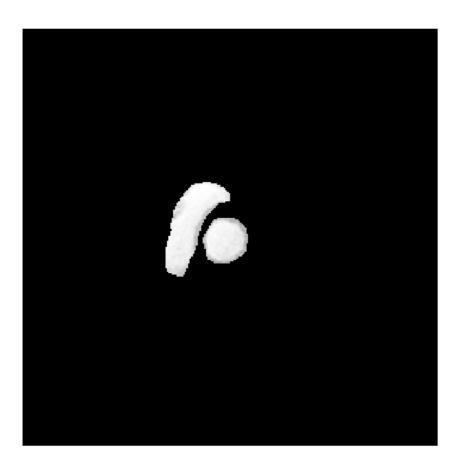
Label selection

Select a single label within image:

Select many labels within image:

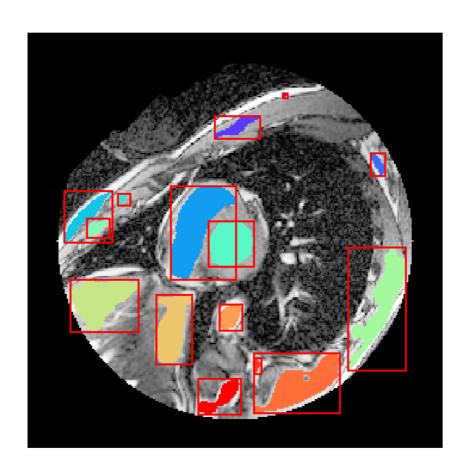
np.where(labels == 1, im, 0)





Object extraction

- Bounding box: range of pixels that completely encloses an object
- ndi.find_objects()
 returns a list of bounding
 box coordinates



Object extraction

```
labels, nlabels = ndi.label(mask)
boxes = ndi.find_objects(labels)
boxes[0]
```

(slice(116,139), slice(120, 141))

im[boxes[0]] im[boxes[1]] im[boxes[2]]



Let's practice!

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Measuring Intensity

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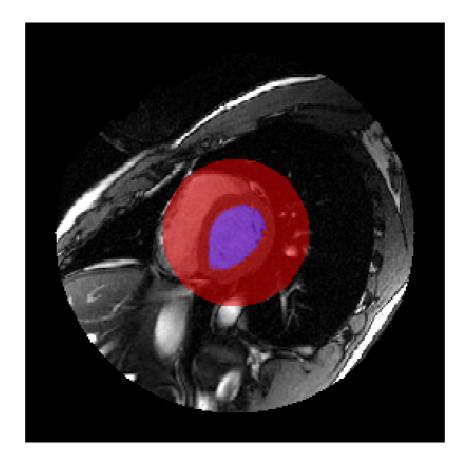
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Measuring intensity

We have the following labels for a single volume of the cardiac time series:

- 1. Left ventricle
- 2. Central portion



Functions

scipy.ndimage.measurements ndi.mean() ndi.median() ndi.sum() ndi.maximum() ndi.standard_deviation() ndi.variance()

Functions applied over all dimensions, optionally at specific labels.

Custom functions:

ndi.labeled_comprehension()

Calling measurement functions

```
import imageio
import scipy.ndimage as ndi
vol=imageio.volread('SCD-3d.npz')
label=imageio.volread('labels.npz')
# All pixels
ndi.mean(vol)
```

3.7892

```
# Labeled pixels
ndi.mean(vol, label)
```

89.2342

```
# Label 1
ndi.mean(vol, label, index=1)
```

163.2930

```
# Labels 1 and 2
ndi.mean(vol, label, index=[1,2])
```

[163.2930, 60.2847]

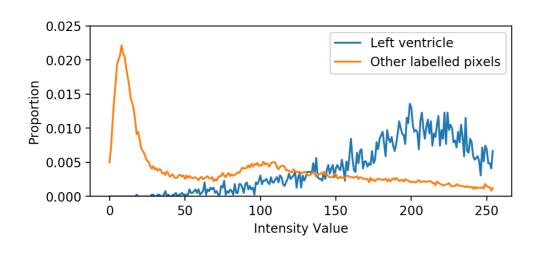
Object histograms

2



Object histograms

```
plt.plot(obj_hists[0],
    label='Left ventricle')
plt.plot(obj_hists[1],
    label='Other labelled pixels')
plt.legend()
plt.show()
```



- Histograms containing multiple tissue types will have several peaks
- Histograms for wellsegmented tissue often resemble a normal distribution

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Measuring morphology

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Morphology

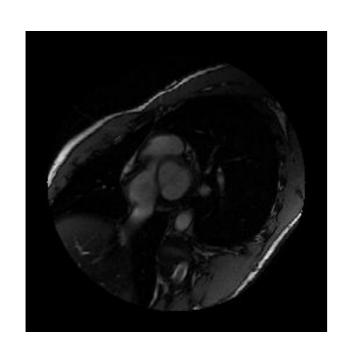




Spatial extent

Spatial extent is the product of:

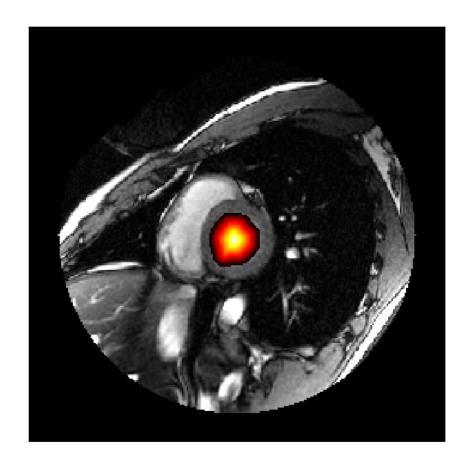
- Space occupied by each element
- 2. Number of array elements



```
# Calculate volume per voxel
d0, d1, d2 = vol.meta['sampling']
dvoxel = d0 * d1 * d2
# Count label voxels
nvoxels=ndi.sum(1, label, index=1)
# Calculate volume of label
volume = nvoxels * dvoxel
volume
```

1249023

Distance transformation



Euclidean Distance

```
# Create a left ventricle mask
mask=np.where(labels == 1, 1, 0)
# In terms of voxels
d=ndi.distance_transform_edt(mask)
d.max()
```

12.3847

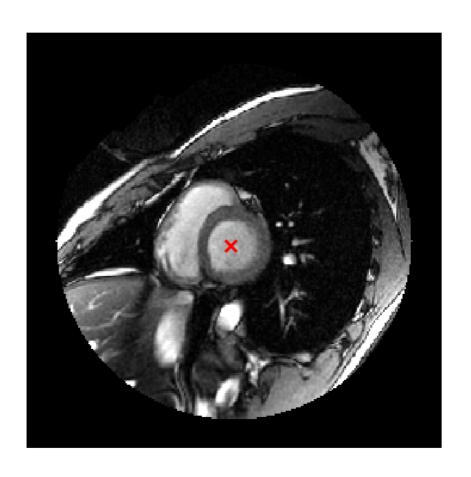
```
# In terms of space
d=ndi.distance_transform_edt(mask,
         sampling=vol.meta['sampling'])
d.max()
```

5.8038

Center of mass

```
(5.5235, 128.0590, 128.0993)
```

```
plt.imshow(vol[5], cmap='gray')
plt.scatter(com[2], com[1])
plt.show()
```



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Measuring in Time

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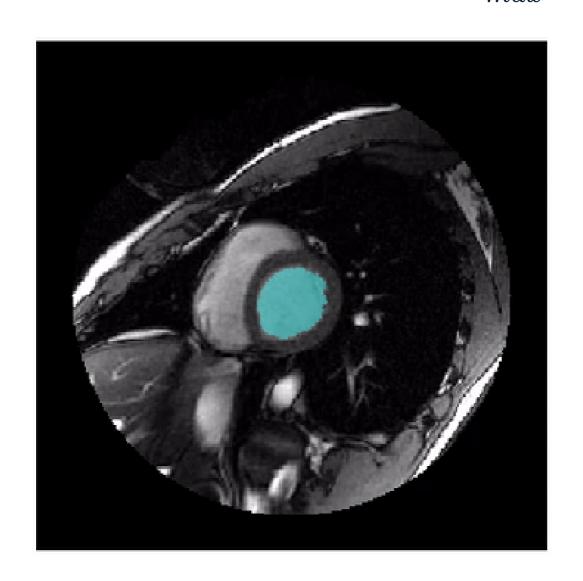


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Ejection fraction

$$Ejection \ Fraction = rac{LV_{max} - LV_{min}}{LV_{max}}$$



Ejection fraction

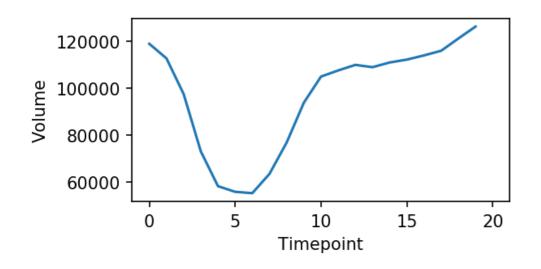
Procedure

- 1. Segment left ventricle
- 2. For each 3D volume in the time series, calculate volume
- 3. Select minimum and maximum
- 4. Calculate ejection fraction

Calculate volume for each time point

```
# Stored in (t,z,x,y) format
vol_ts.shape
labels.shape
```

```
(20, 12, 256, 256)
(20, 12, 256, 256)
```

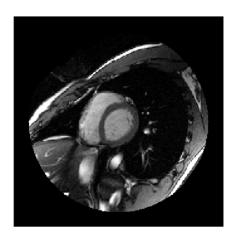


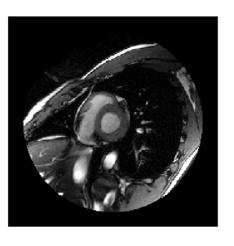
```
# Calculate voxel volume in mm^3
d0,d1,d2,d3=vol_ts.meta['sampling']
dvoxel = d1 * d2 * d3
# Instantiate empty list
ts = np.zeros(20)
# Loop through volume time series
for t in range(20):
    nvoxels=ndi.sum(1,
                    labels[t],
                    index=1)
    ts[t] = nvoxels * dvoxel
plt.plot(ts)
plt.show()
```

Calculate ejection fraction

```
min_vol = ts.min()
max_vol = ts.max()
ejec_frac = (max_vol - min_vol) / max_vol
ejec_frac
```

0.58672





Let's practice!

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