

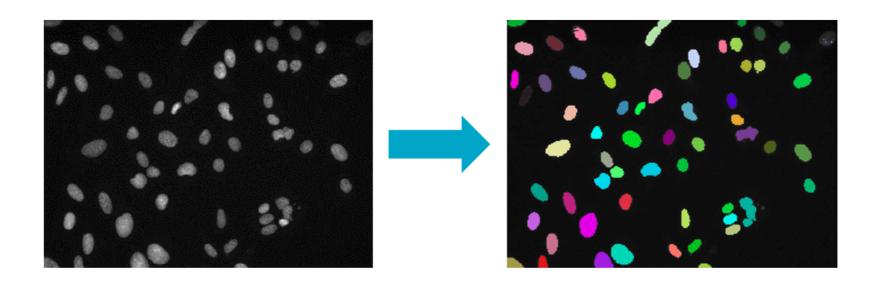


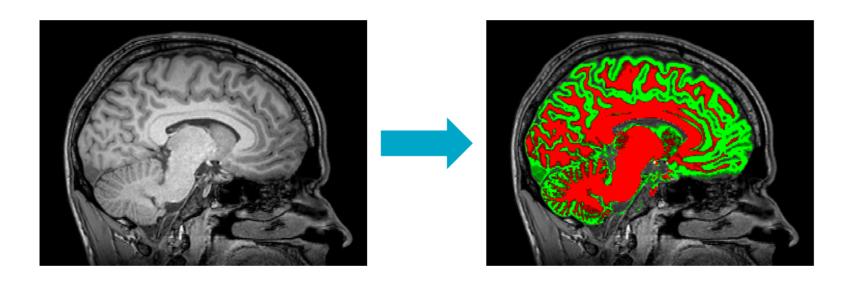
#### BIOMEDICAL IMAGE ANALYSIS IN PYTHON

# **Objects and Labels**

Stephen Bailey Instructor

# Segmentation splits an image into parts

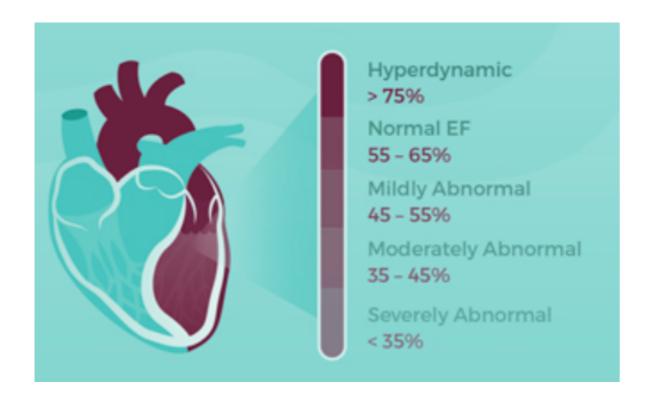


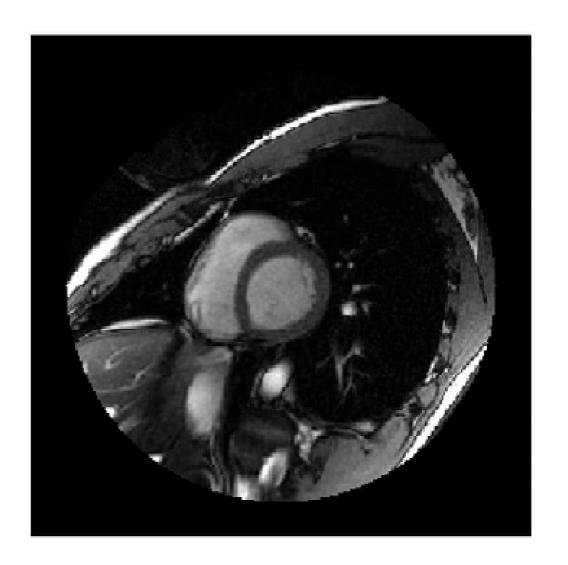




### Sunnybrook Cardiac Database

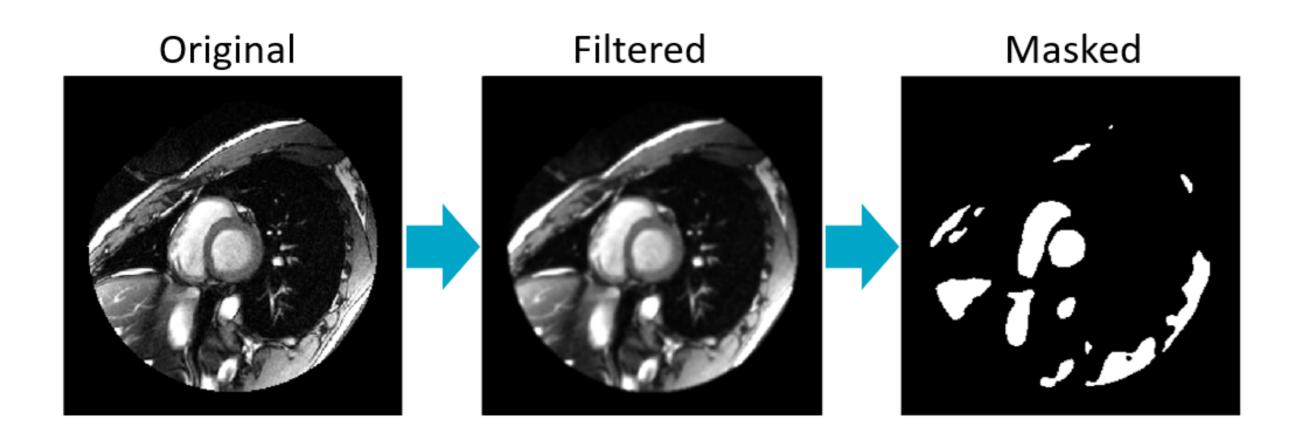
**Ejection fraction**: the proportion of blood pumped out of the heart's left ventricle (LV).







# Labeling image components





### Labeling image components

```
nlabels
14
```

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





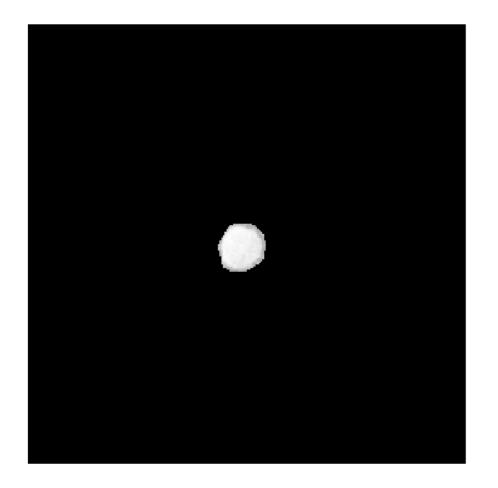
#### Label selection

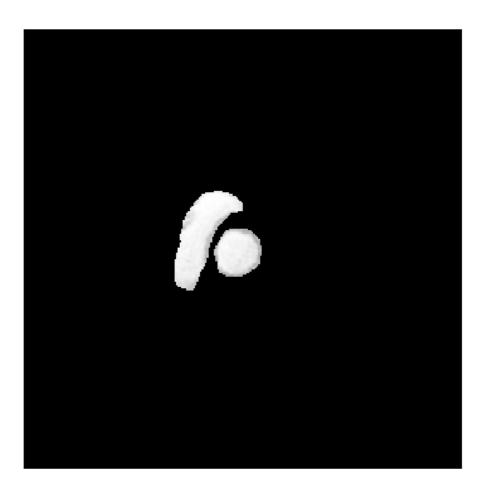
Select a single label within image:

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

Select many labels within image:

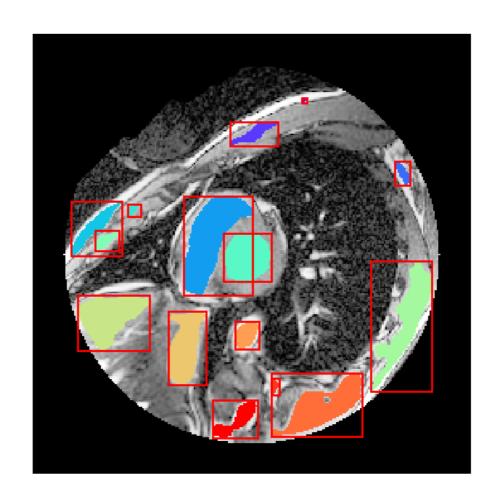
np.where(labels < 3, im, 0)</pre>





### Object extraction

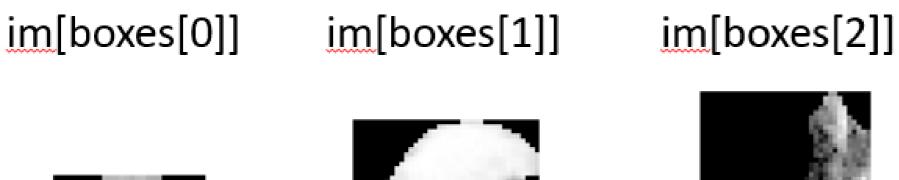
- Bounding box: range of pixels that completely encloses an object
- ndi.find\_objects() returns a list of bounding box coordinates

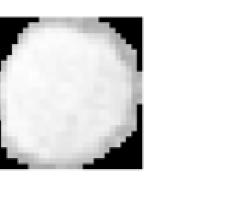


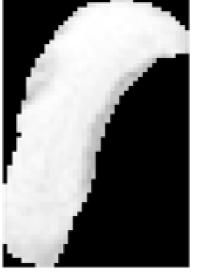
### Object extraction

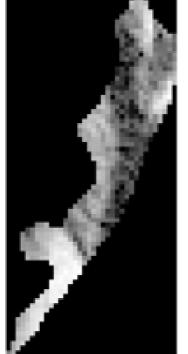
```
labels, nlabels = ndi.label(mask)
boxes = ndi.find_objects(labels)

boxes[0]
   (slice(116,139), slice(120, 141))
```













# Let's practice!





BIOMEDICAL IMAGE ANALYSIS IN PYTHON

# **Measuring Intensity**

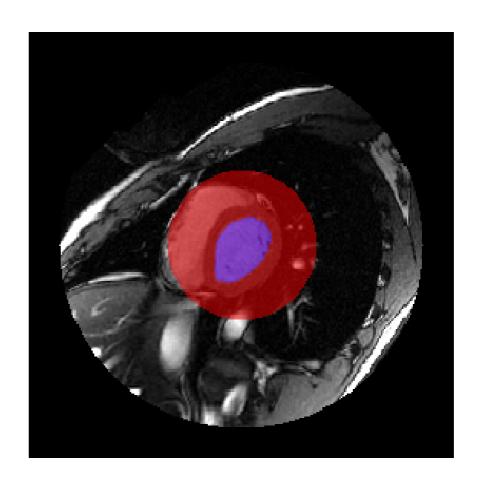
Stephen Bailey Instructor



# 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

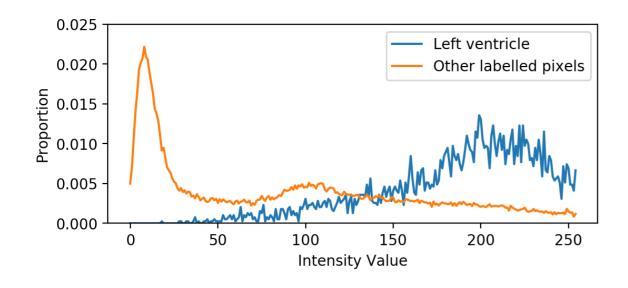
```
import imageio
import scipy.ndimage as ndi
vol=imageio.volread('SCD-3d.npz')
label=imageio.volread('labels.npz')
# All pixels
ndi.mean(vol)
    3.7892
# Labeled pixels
ndi.mean(vol, label)
    89.2342
# Label 1
ndi.mean(vol, label, index=1)
    163.2930
# Labels 1 and 2
ndi.mean(vol, label, index=[1,2])
    [163.2930, 60.2847]
```



# Object histograms

### 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 well-segmented tissue often resemble a normal distribution





# Let's practice!





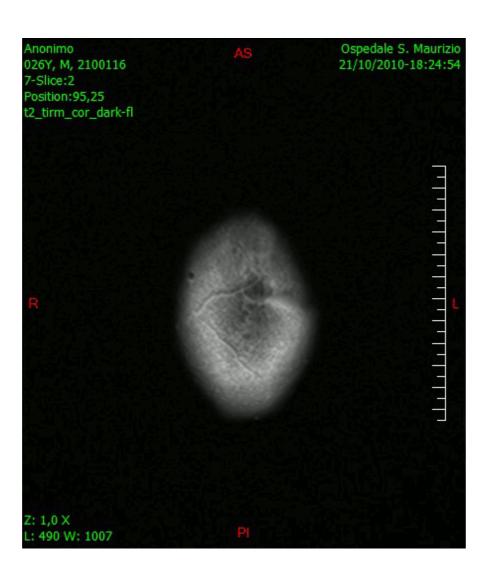
BIOMEDICAL IMAGE ANALYSIS IN PYTHON

# Measuring Morphology

Stephen Bailey Instructor



# Morphology





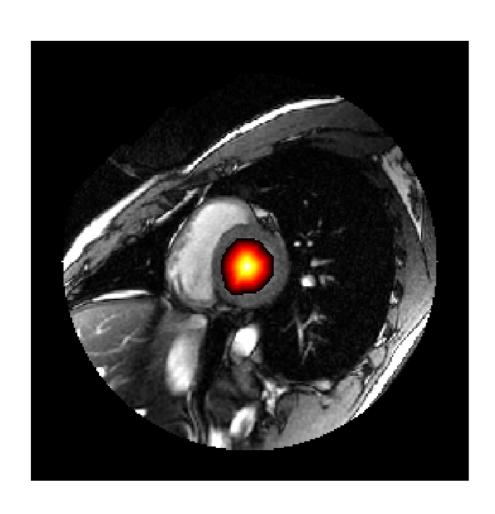
## Spatial extent

#### **Spatial extent** is the product of:

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



#### Distance transformation



#### **Euclidean Distance**

```
# Create a left ventricle mask
mask=np.where(labels == 1, 1, 0)

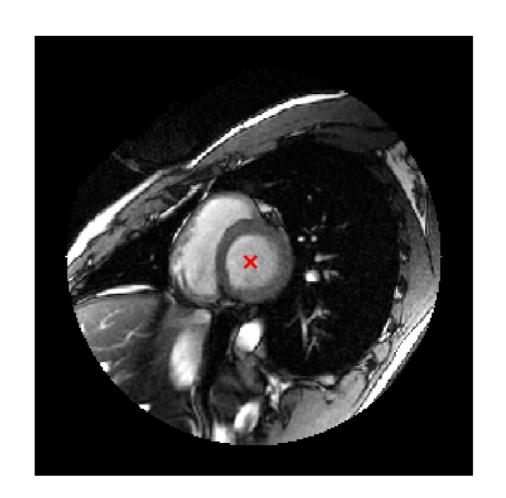
# In terms of voxels
d=ndi.distance_transform_edt(mask)

d.max()
    12.3847
```



#### Center of mass

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







# Let's practice!





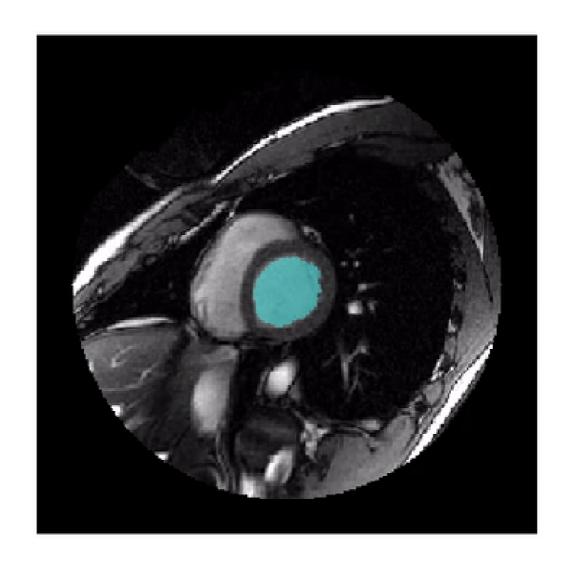
#### BIOMEDICAL IMAGE ANALYSIS IN PYTHON

# **Measuring in Time**

Stephen Bailey Instructor

# Ejection fraction

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





## Ejection fraction

#### **Procedure**

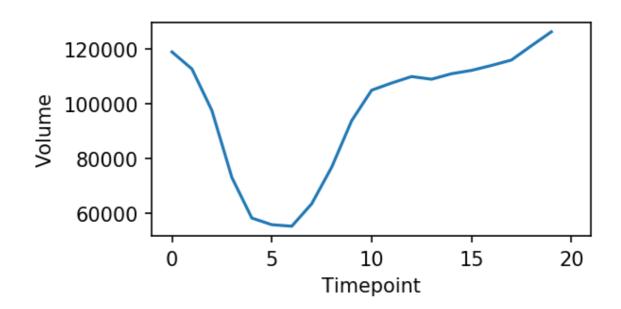
- 1. Segment left ventricle
- 2. For each 3D volume in the time series, calculate volume
- 3. Select minimum and maximum
- 4. Calculate ejection fraction



## Calculate volume for each time point

```
# Stored in (t,z,x,y) format
vol_ts.shape
      (20, 12, 256, 256)
labels.shape
      (20, 12, 256, 256)
```

```
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!