

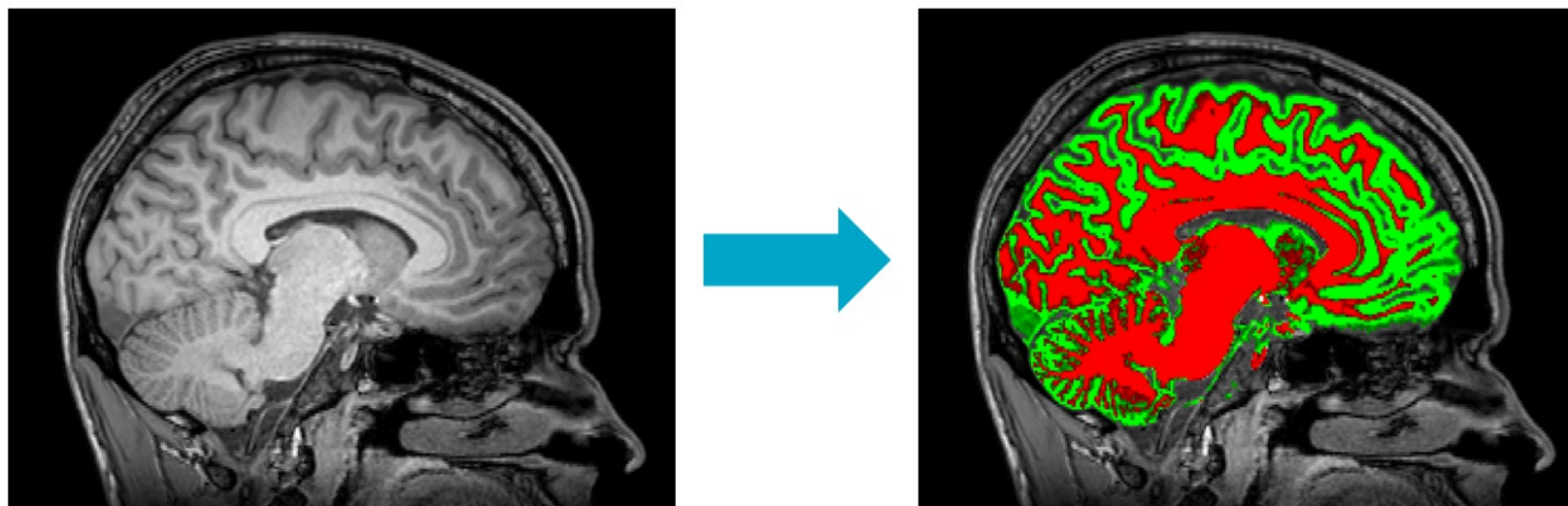
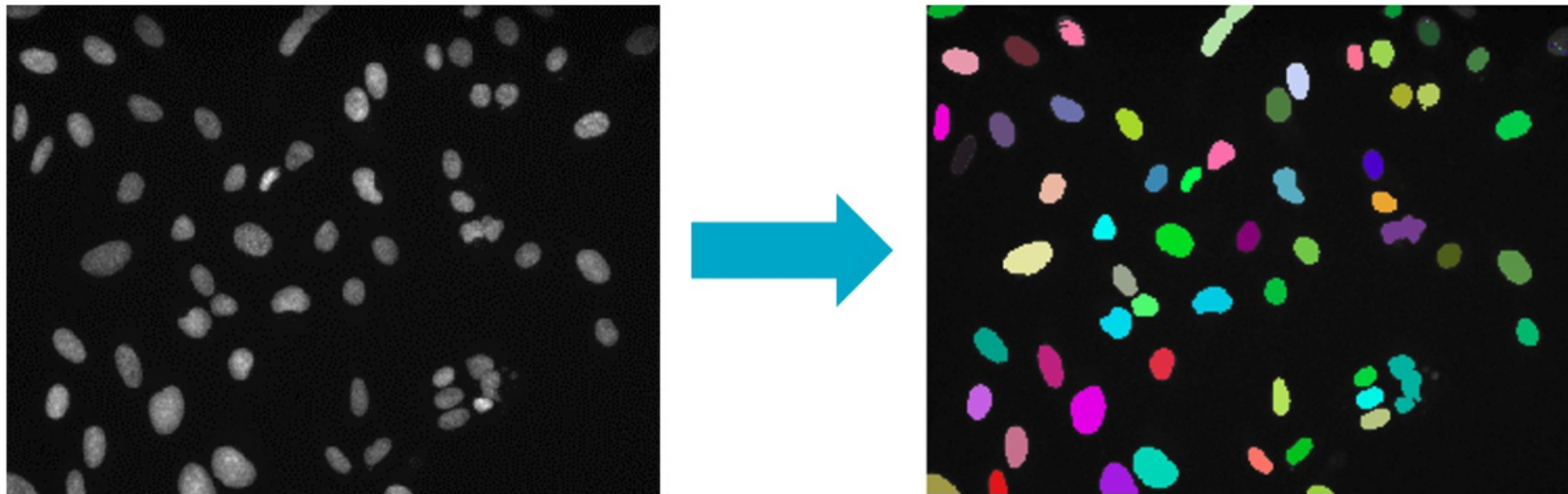


BIOMEDICAL IMAGE ANALYSIS IN PYTHON

# Objects and Labels

Stephen Bailey  
Instructor

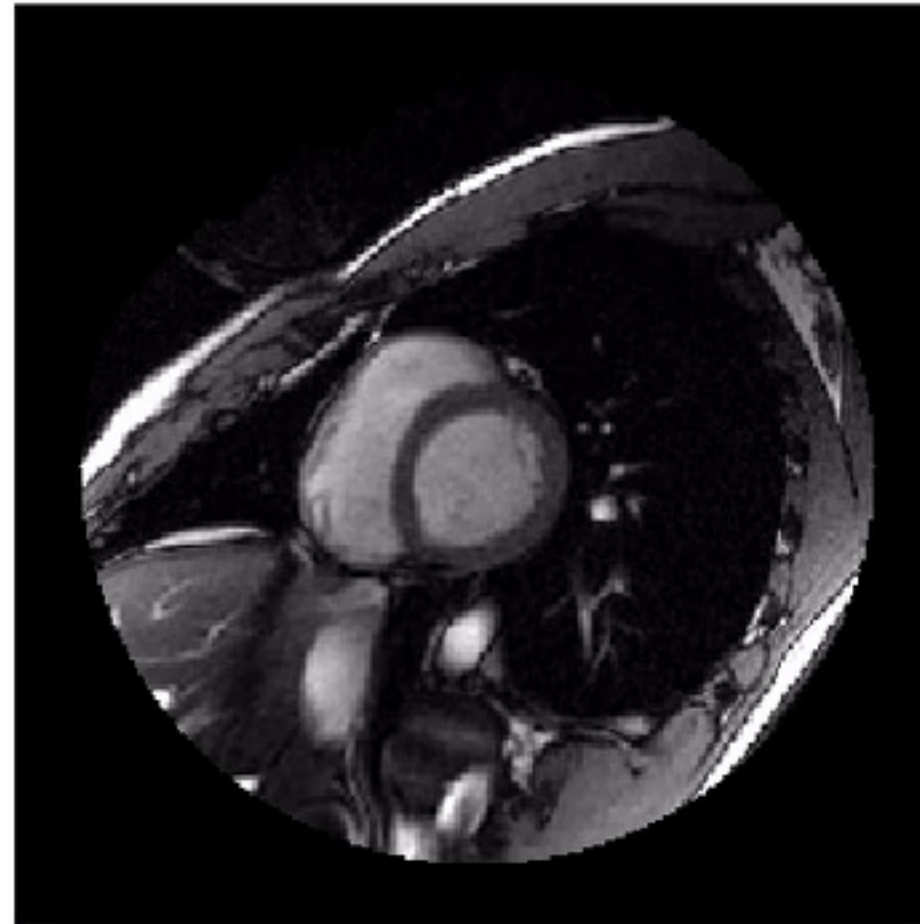
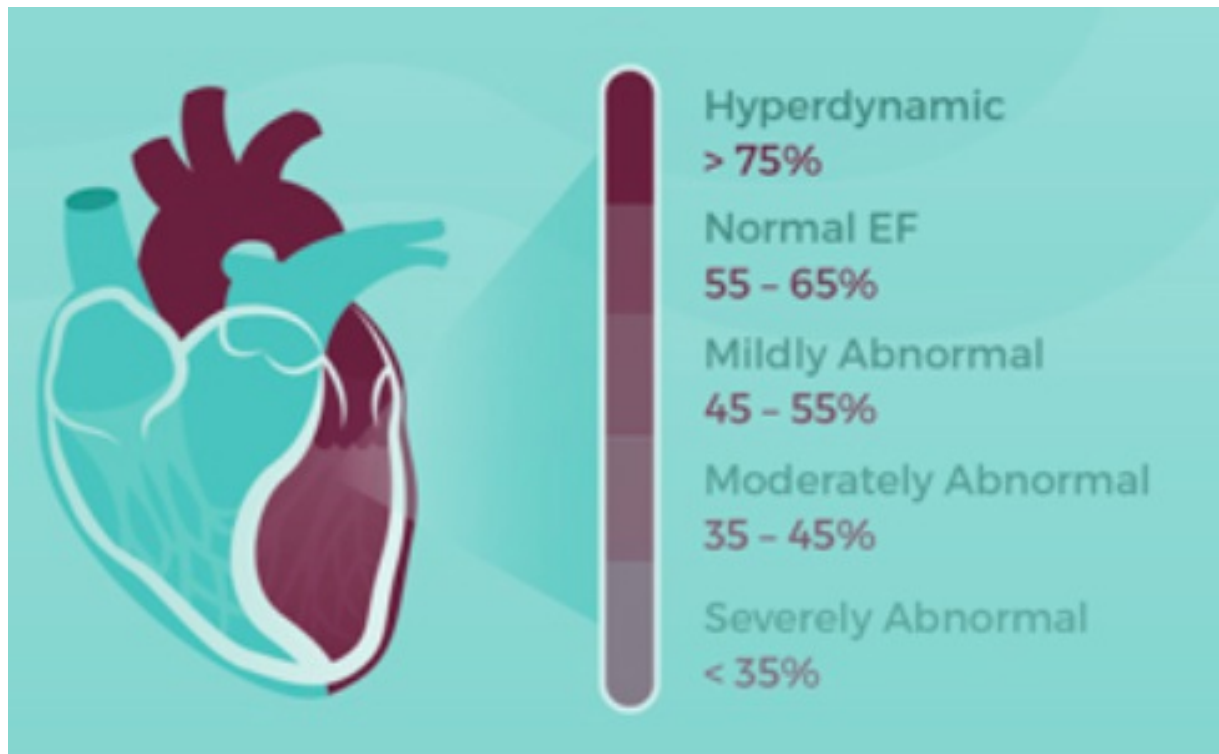
# Segmentation splits an image into parts





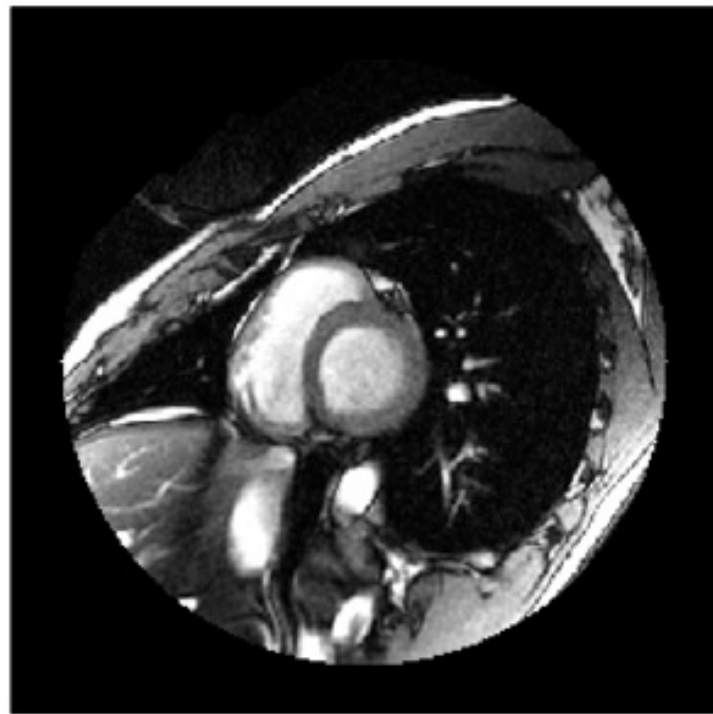
# Sunnybrook Cardiac Database

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

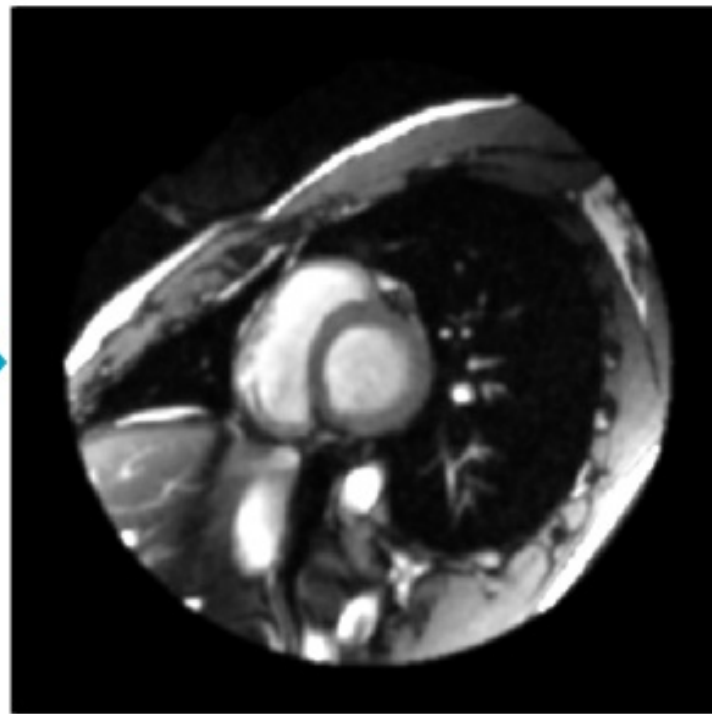


# Labeling image components

Original



Filtered



Masked



# Labeling image components

```
import scipy.ndimage as ndi

im=imageio.imread('SCD4201-2d.dcm')
filt=ndi.gaussian_filter(im,
                        sigma=2)

mask = filt > 150

labels, nlabels = ndi.label(mask)

nlabels
14

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

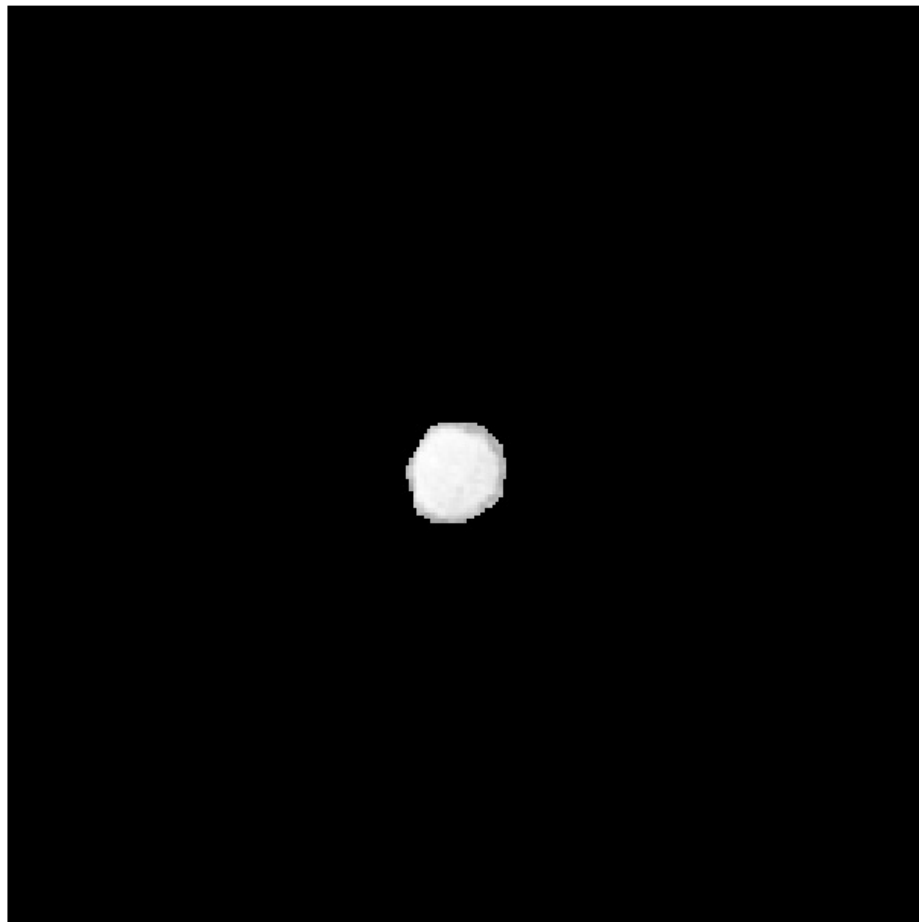




# Label selection

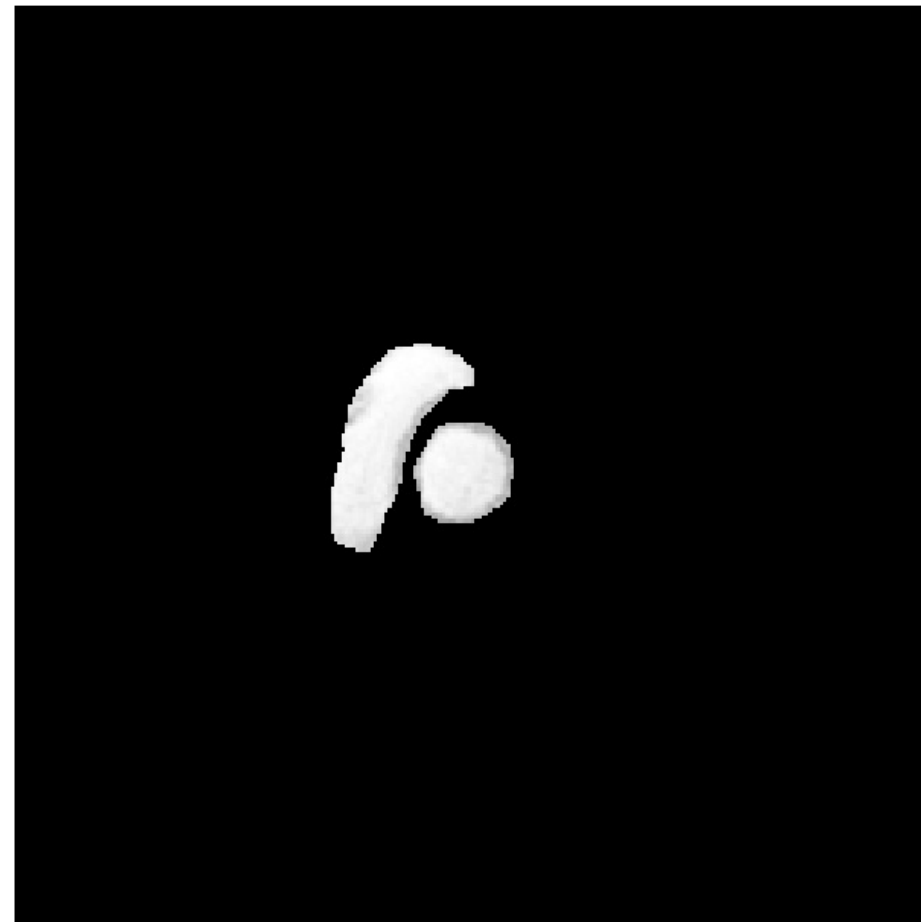
Select a single label within image:

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



Select many labels within image:

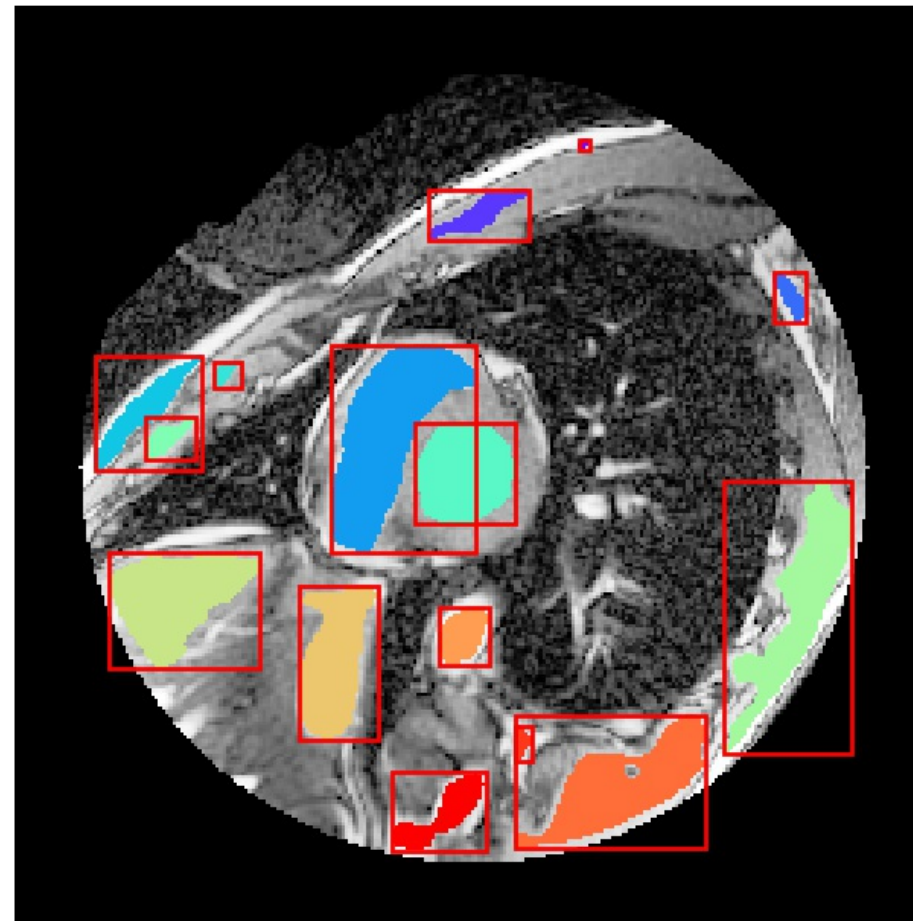
```
np.where(labels < 3, im, 0)
```





# Object extraction

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



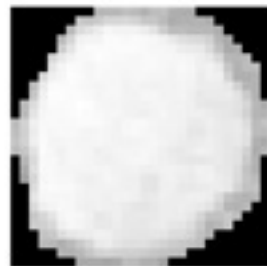


# Object extraction

```
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]]







## BIOMEDICAL IMAGE ANALYSIS IN PYTHON

**Let's practice!**



BIOMEDICAL IMAGE ANALYSIS IN PYTHON

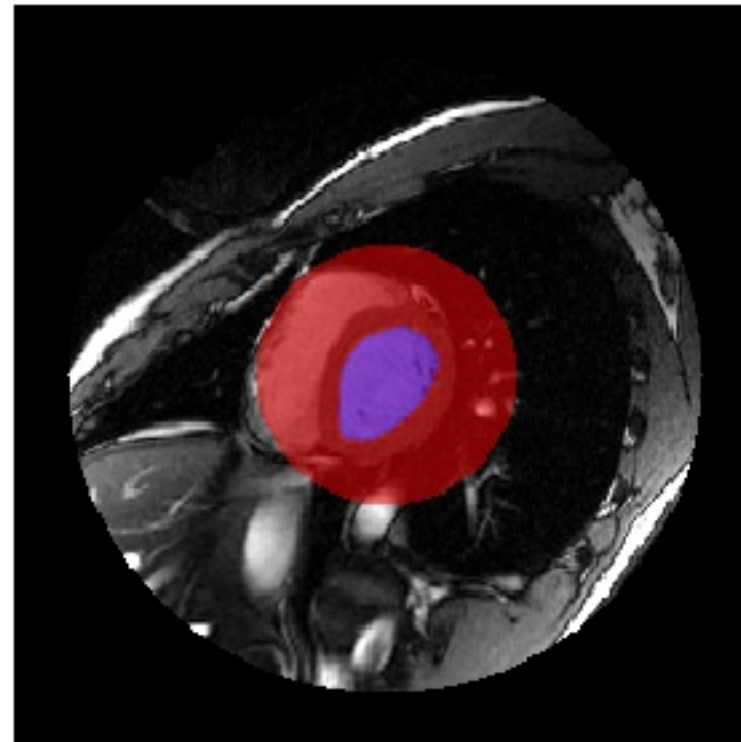
# Measuring Intensity

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

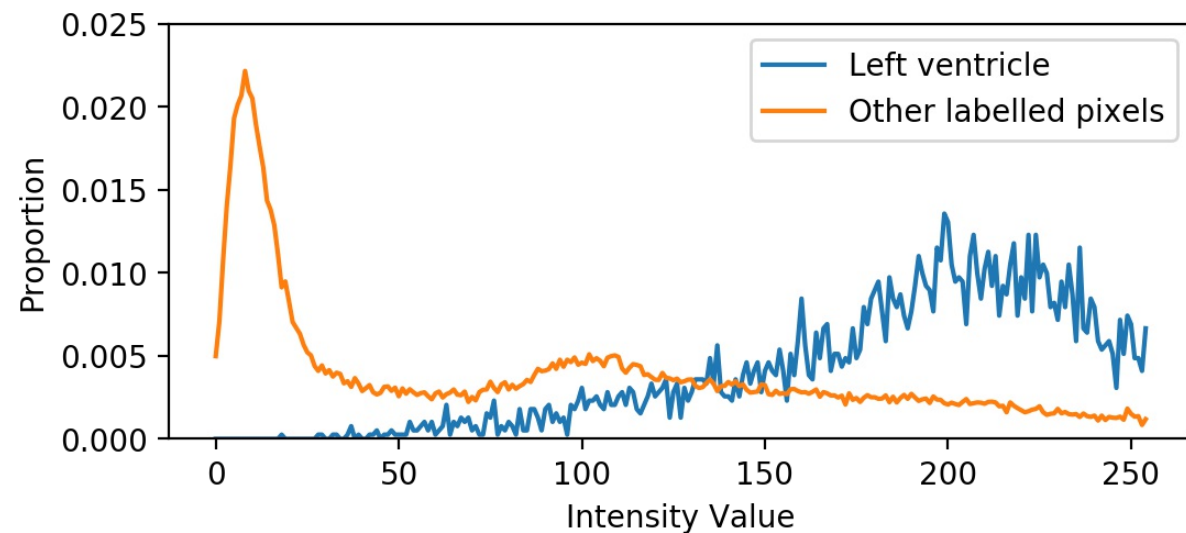
```
hist=ndi.histogram(vol, min=0, max=255, bins=256)

obj_hists=ndi.histogram(vol, 0, 255, 256,
                        labels, index=[1, 2])

len(obj_hists)
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 well-segmented tissue often resemble a normal distribution





## BIOMEDICAL IMAGE ANALYSIS IN PYTHON

**Let's practice!**

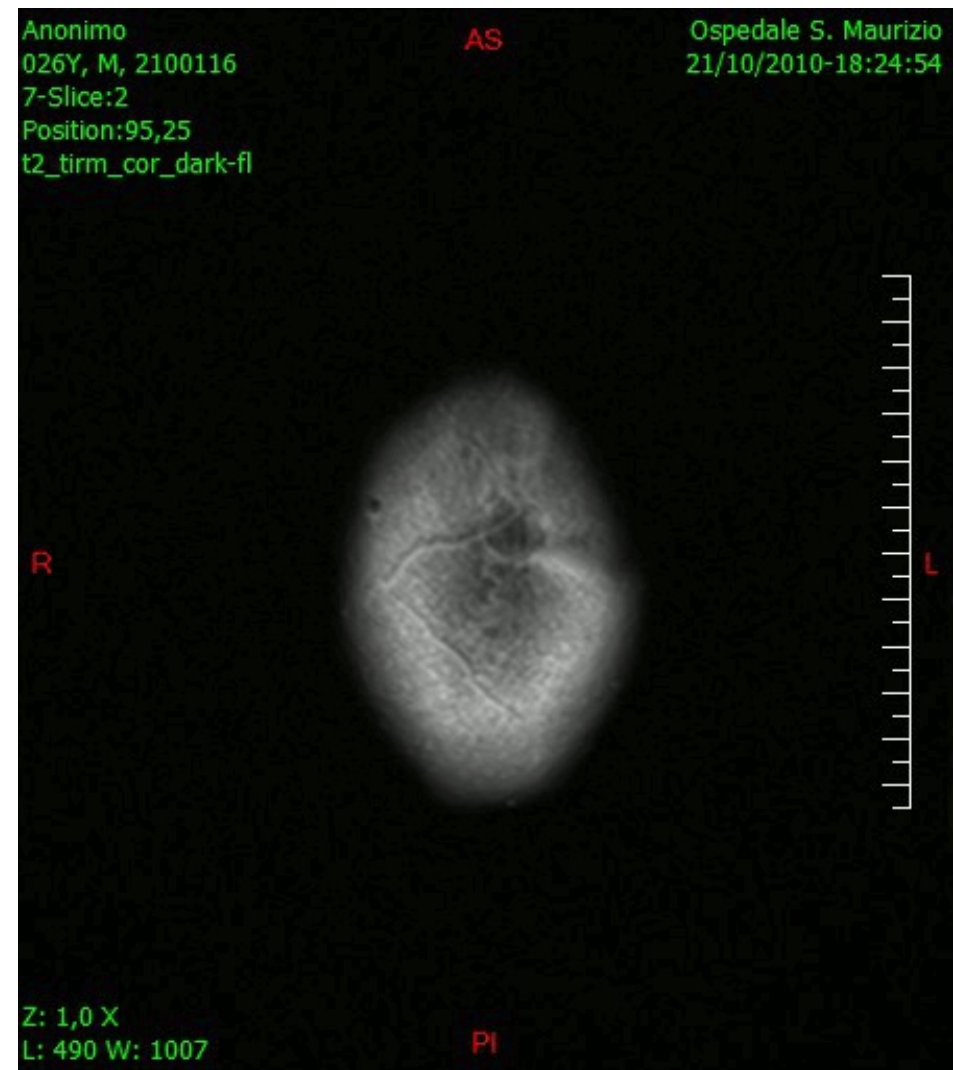


BIOMEDICAL IMAGE ANALYSIS IN PYTHON

# Measuring Morphology

Stephen Bailey  
Instructor

# Morphology

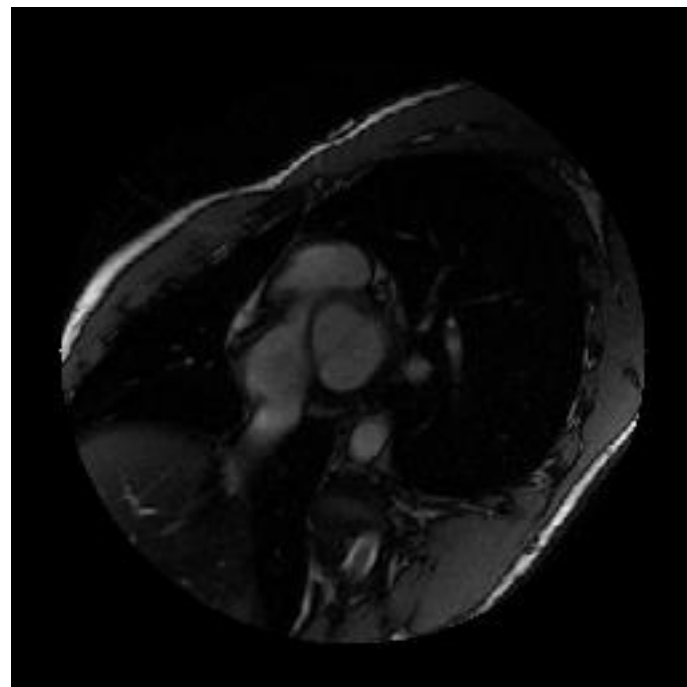




# Spatial extent

**Spatial extent** is the product of:

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



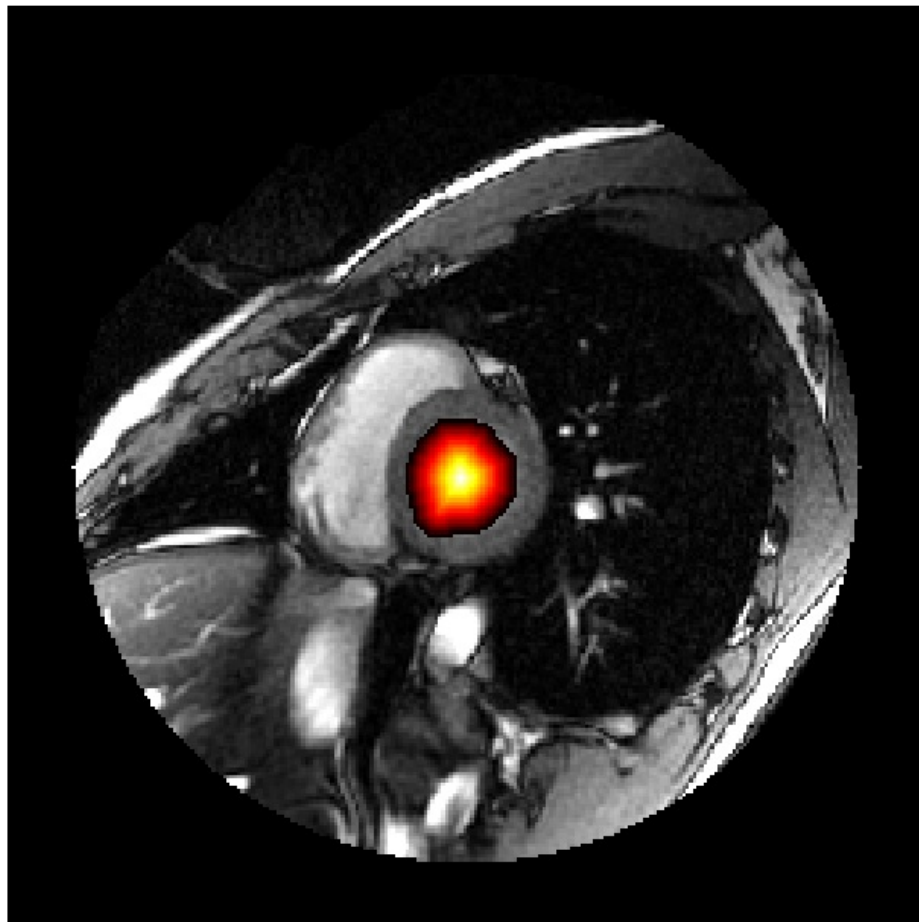
```
# Calculate volume per voxel
d0, d1, d2 = vol.meta['sampling']
dvoxel = d0 * d1 * d2

# 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'])

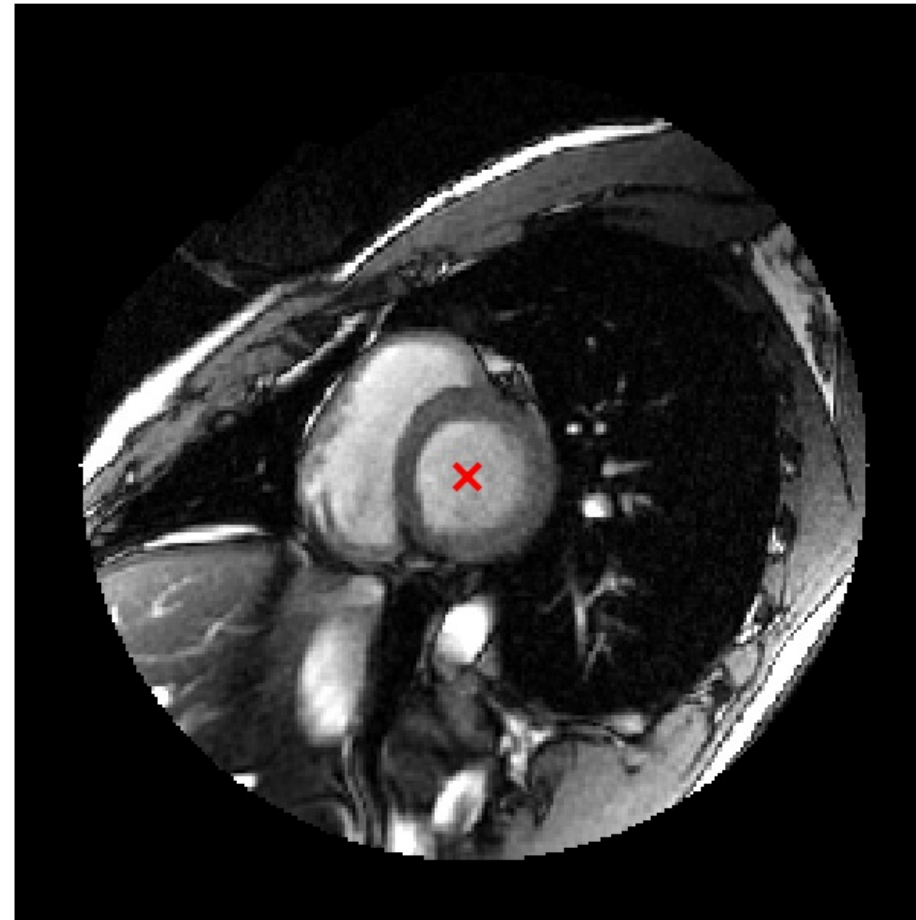
d.max()
5.8038
```

# Center of mass

```
com=ndi.center_of_mass(vol,  
                        labels,  
                        index=1)
```

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

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





## BIOMEDICAL IMAGE ANALYSIS IN PYTHON

**Let's practice!**





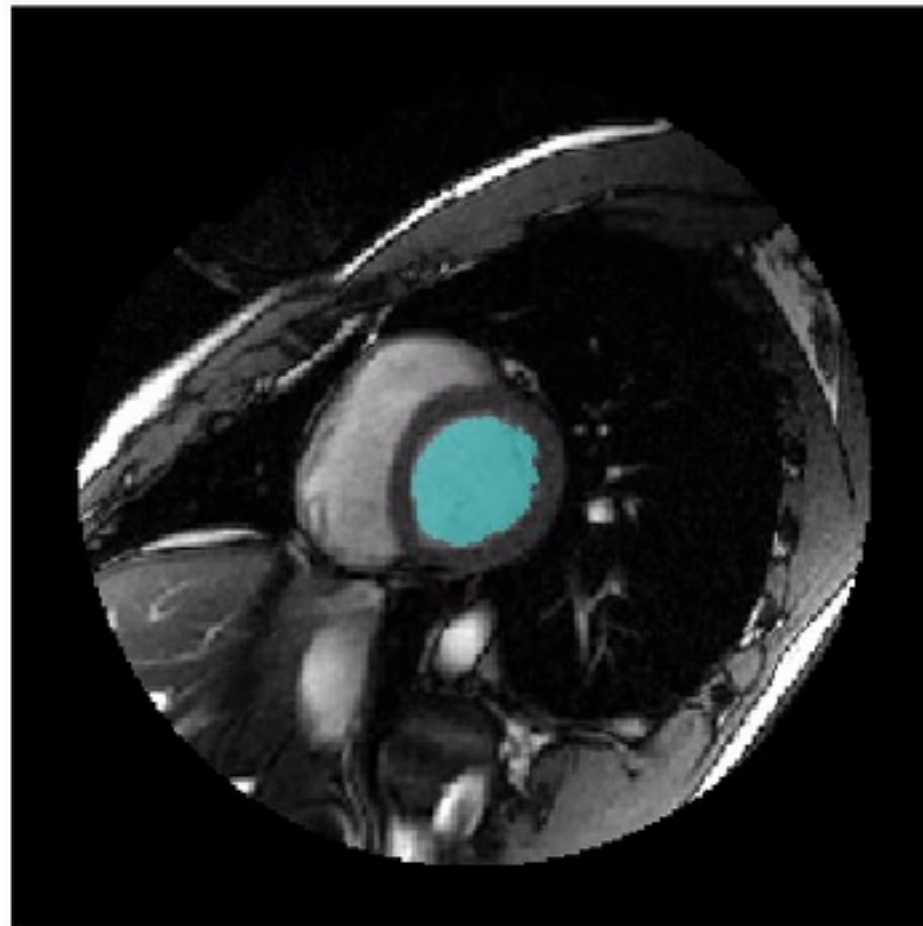
BIOMEDICAL IMAGE ANALYSIS IN PYTHON

# Measuring in Time

Stephen Bailey  
Instructor

# Ejection fraction

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





# 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)
```

```
# Calculate voxel volume in mm^3
d0,d1,d2,d3=vol_ts.meta['sampling']
dvoxel = d1 * d2 * d3
```

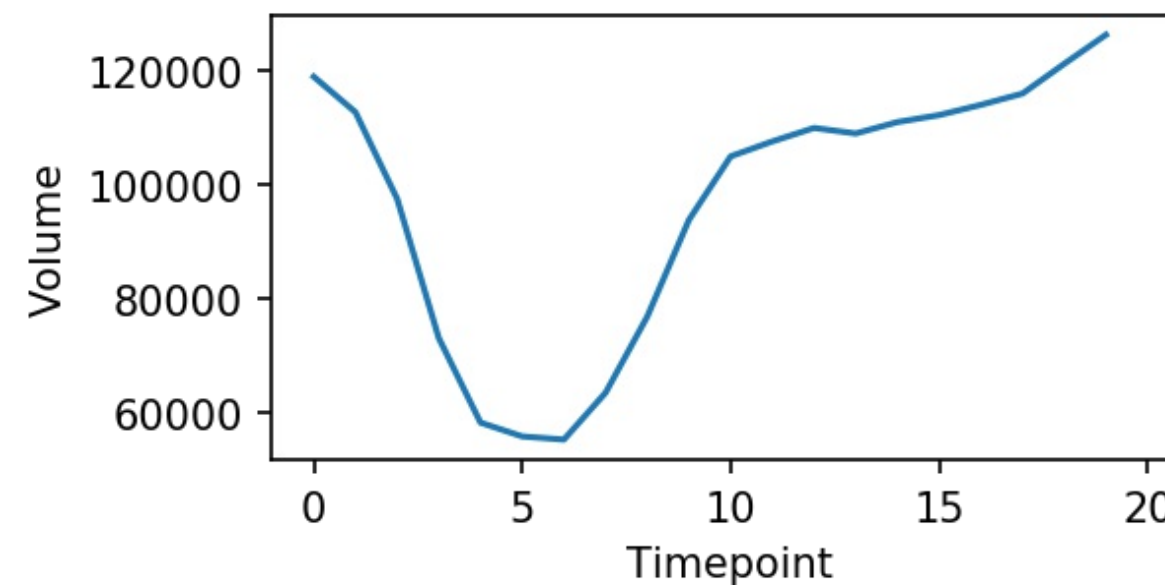
```
# Instantiate empty list
ts = np.zeros(20)
```

```
# 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()
```

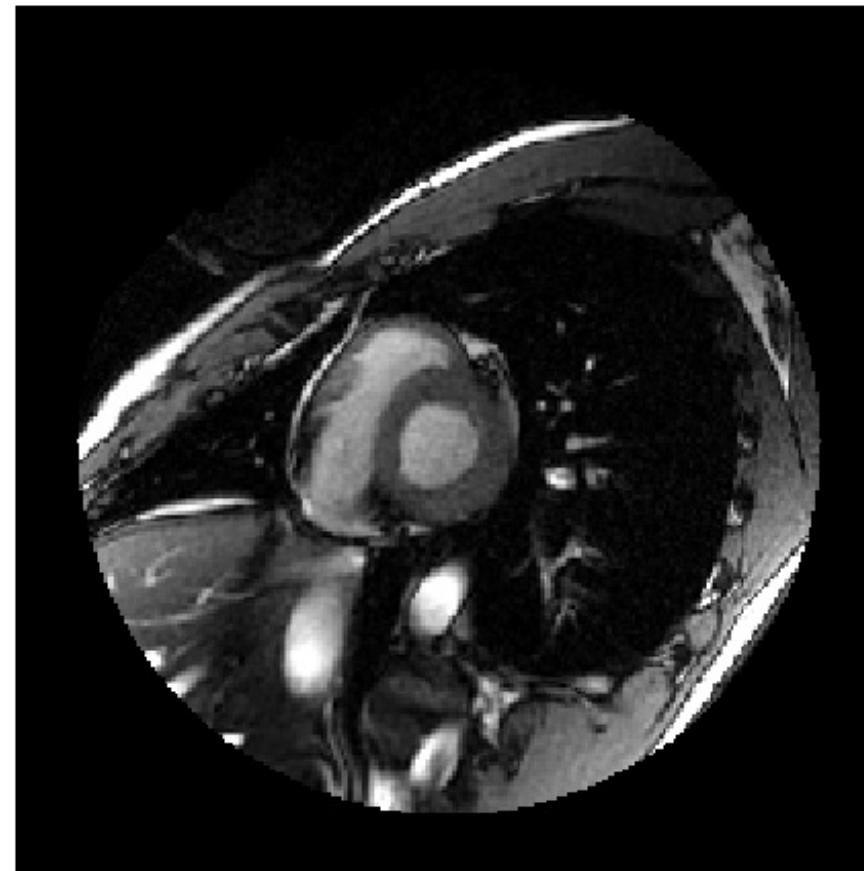
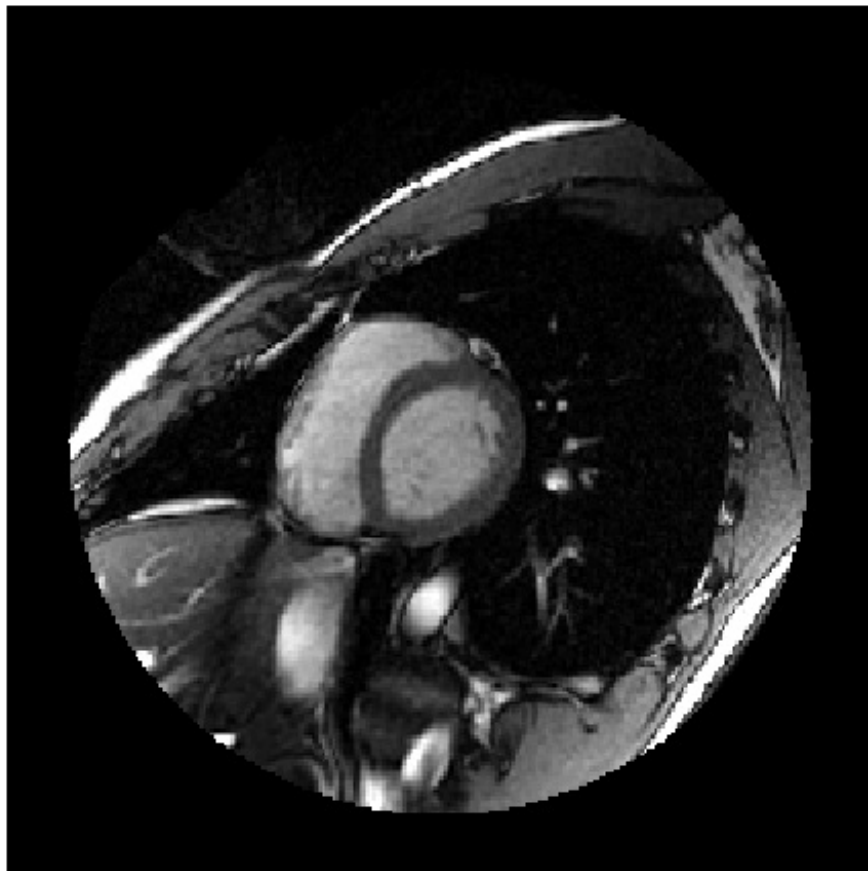


# Calculate ejection fraction

```
min_vol = ts.min()
max_vol = ts.max()

ejec_frac = (max_vol - min_vol) / max_vol

ejec_frac
0.58672
```





## BIOMEDICAL IMAGE ANALYSIS IN PYTHON

**Let's practice!**