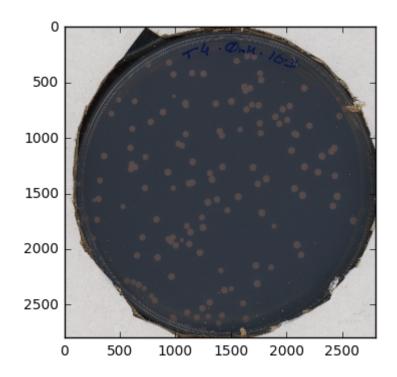
```
In [1]: # first: find regions that are plates
In [2]:
        # import settings and packages
        %matplotlib inline
        from skimage import io
        import matplotlib.pyplot as plt
        import matplotlib.patches as mpatches
        from skimage.filters import threshold otsu
        from skimage.segmentation import clear border
        from skimage.measure import label, regionprops
        from skimage.morphology import closing, square
        from skimage.color import label2rgb
        from scipy import ndimage as ndi
        import glob
        # create new class of object called cultureplate:
In [3]:
        class cultureplate:
            def init (self, name, boundaries, colonies, image, label,
        pos, scanner):
                self.name = name
                self.colonies = dict()
                self.boundaries = tuple()
                self.image = ()
                self.label = ()
                self.pos = ()
                self.scanner=()
            def add colony(self, colonies, key, value):
                self.colonies[key]=value
            def rename(self, newname):
                self.name = newname
            def add image(self, newimage):
                self.image = newimage
        # import final scanned image
In [5]:
        # after having imported it into the local folder
        # and separate out blue channel (could alternatively use green)
        plates1 = io.imread('/home/cmarx/Documents/Jessica/images/deathat4mM_
        170917/scanner2/scan 2017-09-24 12:02:02.tiff')
In [6]: cultureplates_manual = list()
In [7]: plateA = plates1[300:3100, 3400:6200,:]
```

```
In [8]: plt.imshow(plateA)
```

Out[8]: <matplotlib.image.AxesImage at 0x7fc56d61b470>

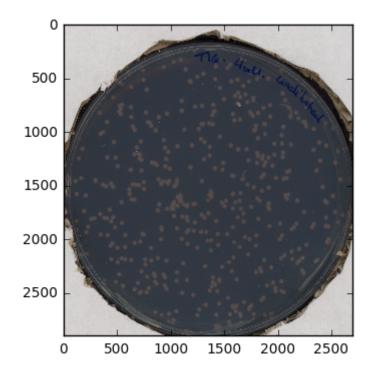


Out[9]: 1

```
In [10]: cultureplates_manual[0].add_image(plates1[300:3100, 3400:6200,:])
    cultureplates_manual[0].boundaries=(300,3400,3100,6200)
    cultureplates_manual[0].pos='topright'
    cultureplates_manual[0].scanner='2'
```

```
In [11]: plateB = plates1[3200:6100, 3600:6300,:]
    plt.imshow(plateB)
```

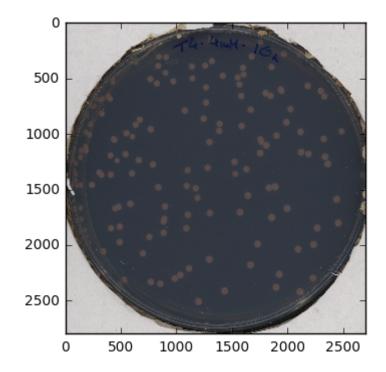
Out[11]: <matplotlib.image.AxesImage at 0x7fc5688e69b0>



Out[12]: 2

```
In [13]: plateC= plates1[300:3100, 400:3100,:]
    plt.imshow(plateC)
```

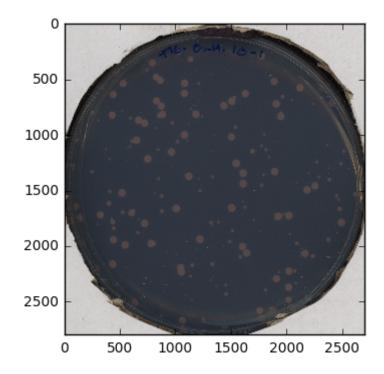
Out[13]: <matplotlib.image.AxesImage at 0x7fc566a77438>



Out[14]: 3

```
In [15]: plateD= plates1[3400:6200, 500:3200,:]
    plt.imshow(plateD)
```

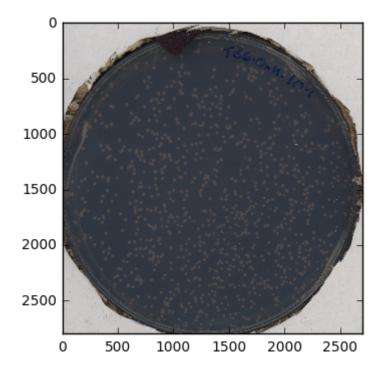
Out[15]: <matplotlib.image.AxesImage at 0x7fc566a5dd30>



Out[16]: 4

```
In [17]: plateE= plates1[6500:9300, 400:3100,:]
    plt.imshow(plateE)
```

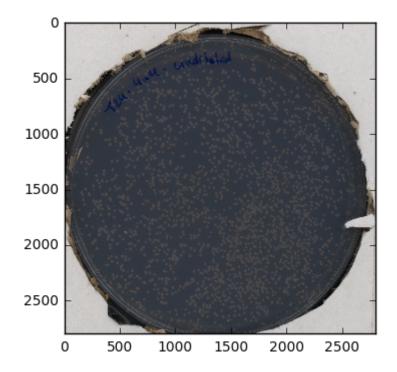
Out[17]: <matplotlib.image.AxesImage at 0x7fc5669b9cf8>



Out[18]: 5

```
In [19]: plateF= plates1[6400:9200, 3500:6300,:]
    plt.imshow(plateF)
```

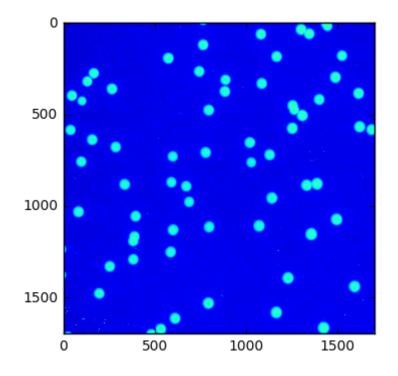
Out[19]: <matplotlib.image.AxesImage at 0x7fc5669a3518>



Out[20]: 6

In [23]: # now find thresholds
 testplate = plateC[600:2300, 500:2200, 0]
 plt.imshow(testplate)

Out[23]: <matplotlib.image.AxesImage at 0x7fc564a3db70>



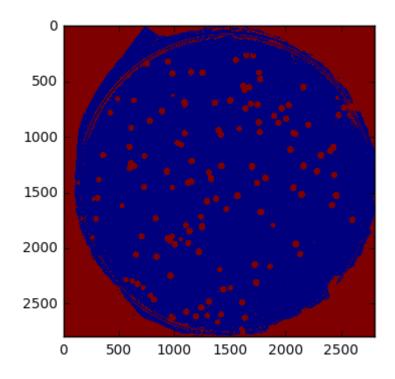
In [24]: thresh = threshold_otsu(testplate)

In [25]: thresh

Out[25]: 68

```
In [26]: bw_plate1 = closing(plateA[:,:,0] > thresh, square(3))
plt.imshow(bw_plate1)
```

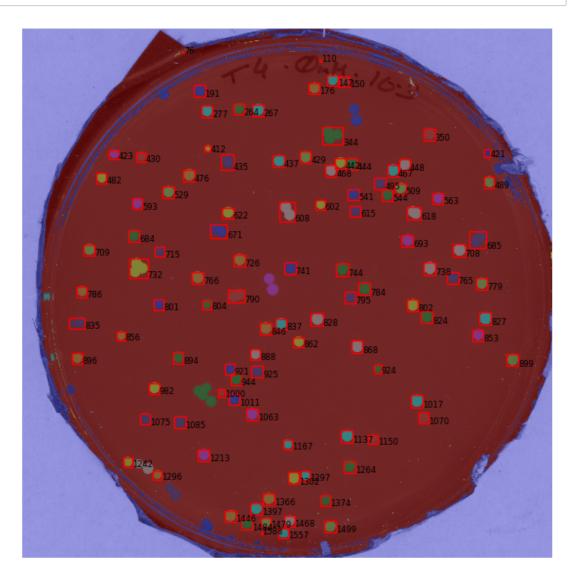
Out[26]: <matplotlib.image.AxesImage at 0x7fc564a18e10>



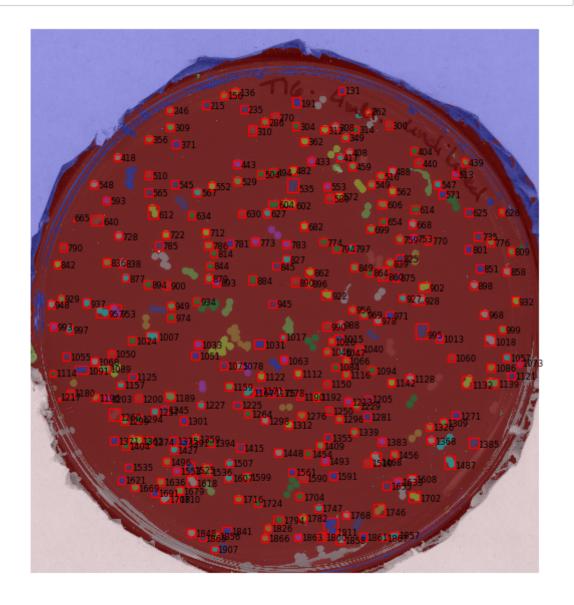
```
In [27]: del plateA
    del plateB
    del plateC
    del plateD
    del plateE
    del plateF
```

```
In [28]:
         for platenum in range(0,6):
              bw plate = closing(cultureplates manual[platenum].image[:,:,0] >
         thresh, square(3))
             labeled plate = label(bw plate)
              overlaid plate = label2rgb(labeled plate, image=cultureplates man
         ual[platenum].image)
              cultureplates manual[platenum].label = labeled plate # add the in
         fo about labeling
             cultureplates manual[platenum].add image(overlaid plate) # add th
         is overlay image to the object
              for region in regionprops(labeled plate):
                  if 40 < region.area < 10000:</pre>
                      minr, minc, maxr, maxc = region.bbox
                      if 0.6<(maxr-minr)/(maxc-minc)<1.4:</pre>
                          if region.area>0.6*((maxr-minr)*(maxc-minc)):
                              cultureplates manual[platenum].colonies[region.la
         bel] = region.coords # add only the good colonies to the dictionary
```

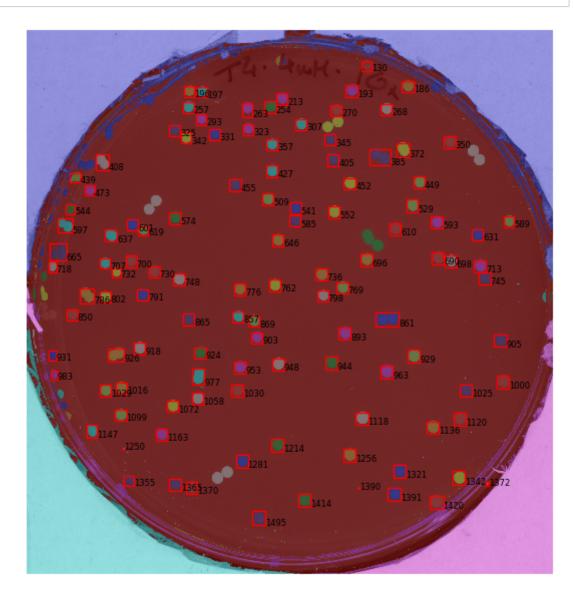
```
In [30]:
         # find the good colonies
         fig, ax = plt.subplots(figsize=(10, 6))
         ax.imshow(cultureplates manual[0].image)
         for region in regionprops(cultureplates manual[0].label):
             if 40 < region.area < 10000:
                 minr, minc, maxr, maxc = region.bbox
                 if 0.6<(maxr-minr)/(maxc-minc)<1.4:</pre>
                          if region.area>0.6*((maxr-minr)*(maxc-minc)):
                              rect = mpatches.Rectangle((minc, minr), maxc - mi
         nc, maxr - minr,
                                                fill=False, edgecolor='red', li
         newidth=1)
                              ax.add patch(rect)
                              ax.text(maxc, maxr, str(region.label),
         fontsize=6)
         ax.set axis off()
         plt.tight layout()
         plt.show()
```



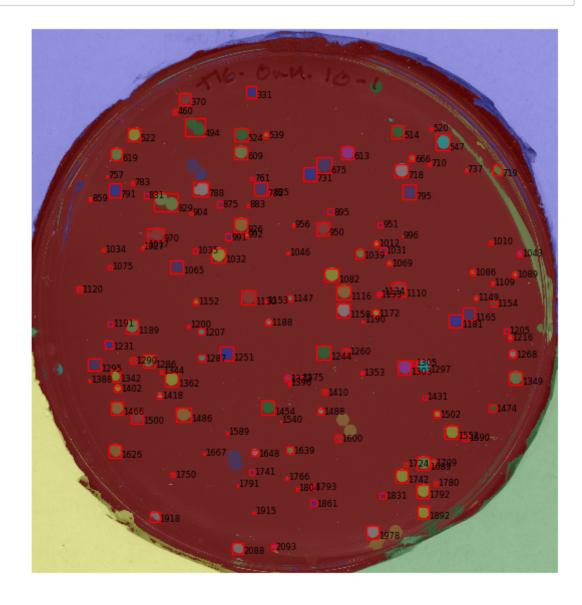
```
In [31]:
         fig, ax = plt.subplots(figsize=(10, 6))
         ax.imshow(cultureplates manual[1].image)
         for region in regionprops(cultureplates_manual[1].label):
             if 40 < region.area < 10000:
                 minr, minc, maxr, maxc = region.bbox
                  if 0.6<(maxr-minr)/(maxc-minc)<1.4:</pre>
                          if region.area>0.6*((maxr-minr)*(maxc-minc)):
                              rect = mpatches.Rectangle((minc, minr), maxc - mi
         nc, maxr - minr,
                                                fill=False, edgecolor='red', li
         newidth=1)
                              ax.add_patch(rect)
                              ax.text(maxc, maxr, str(region.label),
         fontsize=6)
         ax.set axis off()
         plt.tight layout()
         plt.show()
```



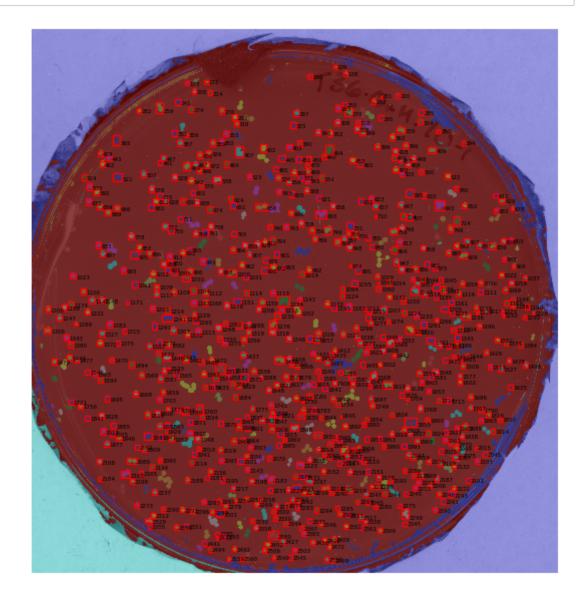
```
In [32]:
         fig, ax = plt.subplots(figsize=(10, 6))
         ax.imshow(cultureplates manual[2].image)
         for region in regionprops(cultureplates_manual[2].label):
             if 40 < region.area < 10000:
                 minr, minc, maxr, maxc = region.bbox
                  if 0.6<(maxr-minr)/(maxc-minc)<1.4:</pre>
                          if region.area>0.6*((maxr-minr)*(maxc-minc)):
                              rect = mpatches.Rectangle((minc, minr), maxc - mi
         nc, maxr - minr,
                                                fill=False, edgecolor='red', li
         newidth=1)
                              ax.add patch(rect)
                              ax.text(maxc, maxr, str(region.label),
         fontsize=6)
         ax.set axis off()
         plt.tight layout()
         plt.show()
```



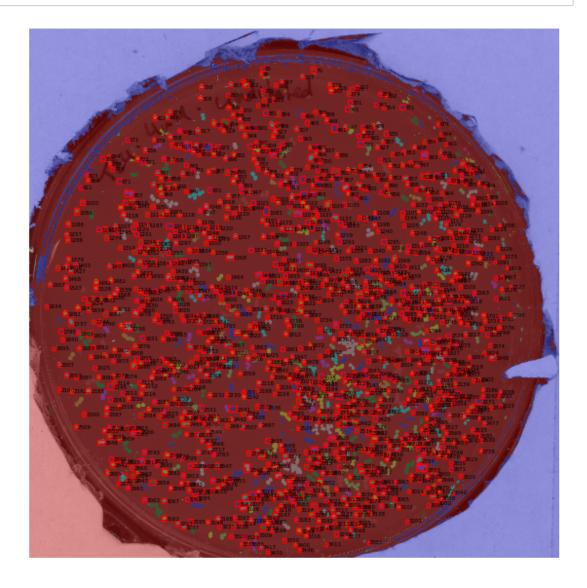
```
In [33]:
         fig, ax = plt.subplots(figsize=(10, 6))
         ax.imshow(cultureplates manual[3].image)
         for region in regionprops(cultureplates_manual[3].label):
             if 40 < region.area < 10000:
                 minr, minc, maxr, maxc = region.bbox
                  if 0.6<(maxr-minr)/(maxc-minc)<1.4:</pre>
                          if region.area>0.6*((maxr-minr)*(maxc-minc)):
                              rect = mpatches.Rectangle((minc, minr), maxc - mi
         nc, maxr - minr,
                                                fill=False, edgecolor='red', li
         newidth=1)
                              ax.add patch(rect)
                              ax.text(maxc, maxr, str(region.label),
         fontsize=6)
         ax.set axis off()
         plt.tight layout()
         plt.show()
```



```
In [35]:
         fig, ax = plt.subplots(figsize=(10, 6))
         ax.imshow(cultureplates manual[4].image)
         for region in regionprops(cultureplates_manual[4].label):
             if 40 < region.area < 10000:
                 minr, minc, maxr, maxc = region.bbox
                  if 0.6<(maxr-minr)/(maxc-minc)<1.4:</pre>
                         if region.area>0.6*((maxr-minr)*(maxc-minc)):
                              rect = mpatches.Rectangle((minc, minr), maxc - mi
         nc, maxr - minr,
                                                fill=False, edgecolor='red', li
         newidth=1)
                              ax.add_patch(rect)
                              ax.text(maxc, maxr, str(region.label),
         fontsize=4)
         ax.set axis off()
         plt.tight layout()
         plt.show()
```



```
In [37]:
         fig, ax = plt.subplots(figsize=(10, 6))
         ax.imshow(cultureplates manual[5].image)
         for region in regionprops(cultureplates_manual[5].label):
             if 40 < region.area < 10000:
                 minr, minc, maxr, maxc = region.bbox
                  if 0.6<(maxr-minr)/(maxc-minc)<1.4:</pre>
                         if region.area>0.6*((maxr-minr)*(maxc-minc)):
                              rect = mpatches.Rectangle((minc, minr), maxc - mi
         nc, maxr - minr,
                                                fill=False, edgecolor='red', li
         newidth=1)
                              ax.add_patch(rect)
                              ax.text(maxc, maxr, str(region.label),
         fontsize=4)
         ax.set axis off()
         plt.tight layout()
         plt.show()
```



```
In [ ]: # now, loop through all the plates to get the timecourse data
         # if you're absolutely certain the experiment is over,
         # sequester the last 3 timepoints in another folder
         # to prevent anomalous flatlines at the end of growth
In [38]:
         out = open('/home/cmarx/Documents/Jessica/imageprocessing/imageproces
         sing 4mM 170924/scanner2 plate IDs 170924.csv', 'w')
         for eachplate in cultureplates manual:
             out.write(str(eachplate.name)+','+str(eachplate.pos)+','+str(each
         plate.scanner)+'\n')
         out.close()
In [39]:
         path = "/home/cmarx/Documents/Jessica/images/deathat4mM 170917/scanne
         r2/scan *.tiff"
         out = open('/home/cmarx/Documents/Jessica/imageprocessing/imageproces
         sing 4mM 170924/scanner2 colony trajectories 170924.csv', 'w')
         for eachplate in cultureplates manual:
             topbound = eachplate.boundaries[0]
             bottombound = eachplate.boundaries[2]
             leftbound = eachplate.boundaries[1]
             rightbound = eachplate.boundaries[3]
             for filename in glob.glob(path):
                 image1 = io.imread(filename)[topbound:bottombound,
         leftbound:rightbound,01
                 image thresh = closing(image1 > thresh, square(3))
                 colony pixel counts = dict()
                 for colony in eachplate.colonies.keys():
                     whitepix = 0
                     for pixel in eachplate.colonies[colony]:
                         x coord=pixel[0]
                         y coord=pixel[1]
                         if image thresh[x coord, y coord]==True:
                             whitepix = whitepix+1
                     colony_pixel_counts[colony] = (whitepix)
                 for colony2 in colony_pixel_counts.keys():
                     out.write(str(eachplate.name)+','
                               +str(colony2)+','
                               +str(colony pixel counts[colony2])+','
                               +str(filename[-24:-5])+'\n')
         out.close()
```

In []: