

```

## Importing necessary modules

import numpy as np
import matplotlib.pyplot as plt
import cv2 as cv
import os
from math import floor

## This is a helper function for plotting the images and feature maps
## for detailed representation

def plot_images(images, title, cmap=None):
    plt.figure(figsize=(12, 7), dpi=150)
    plt.suptitle(title)
    for idx, image in enumerate(images):
        plt.subplot(int(f"24{idx+1}"))

        if cmap=='gray':
            plt.imshow(image, cmap=cmap)
        else:
            plt.imshow(image)
        plt.axis("off")

## This part of the code is used to load the dataset in
## different color spectrums such as RGB, HSV and Gray
## and resize the images into a shape of (1000, 1000)

path = "/content/Dataset"

def load_images(path, gray=False, color=None):
    images = []
    for img in os.listdir(path):
        img = cv.imread(os.path.join(path, img))

        if color==None:
            img = cv.cvtColor(img, cv.COLOR_BGR2RGB)

        elif color=='HSV':
            img = cv.cvtColor(img, cv.COLOR_BGR2HSV)

        if gray:
            img = cv.cvtColor(img, cv.COLOR_BGR2GRAY)

        img = cv.resize(img, (1000,1000))
        images.append(img)

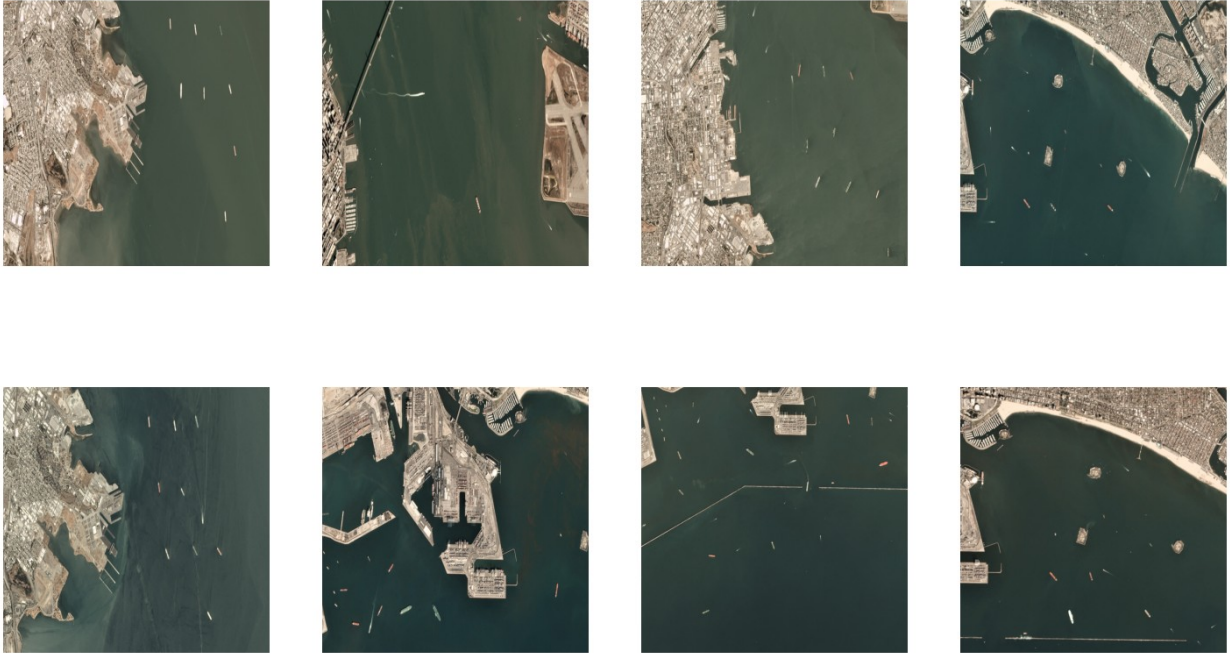
    return images

## Loading the dataset an viewing it with help of the
## helper functions mentioned above

```

```
images = load_images(path)
plot_images(images, "Dataset")
```

Dataset



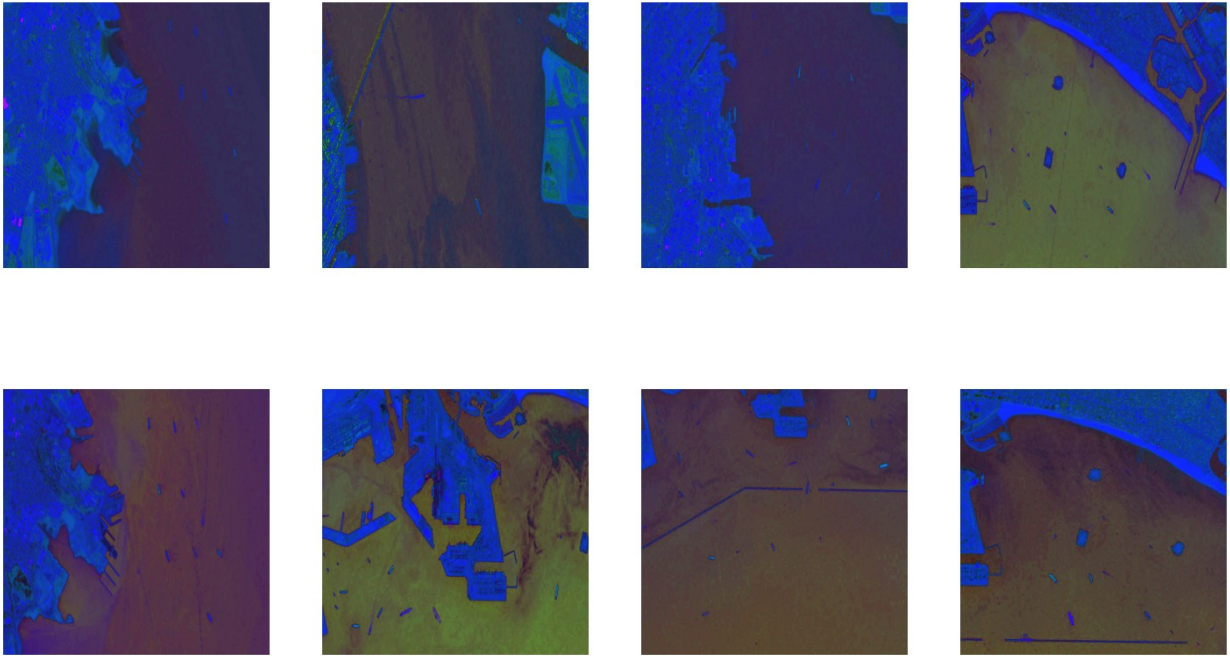
Applying Canny Edge Detection

Using HSV Spectrum

```
## Loading the images in HSV format and viewing it for comparing  
## them with regular RGB images, to see if there is any difference.  
## HSV images helped us a lot in differentiating the water bodies  
## from other objects in the images.
```

```
hsv_images = load_images(path, color='HSV', gray=False)  
plot_images(hsv_images, title="HSV Images")
```

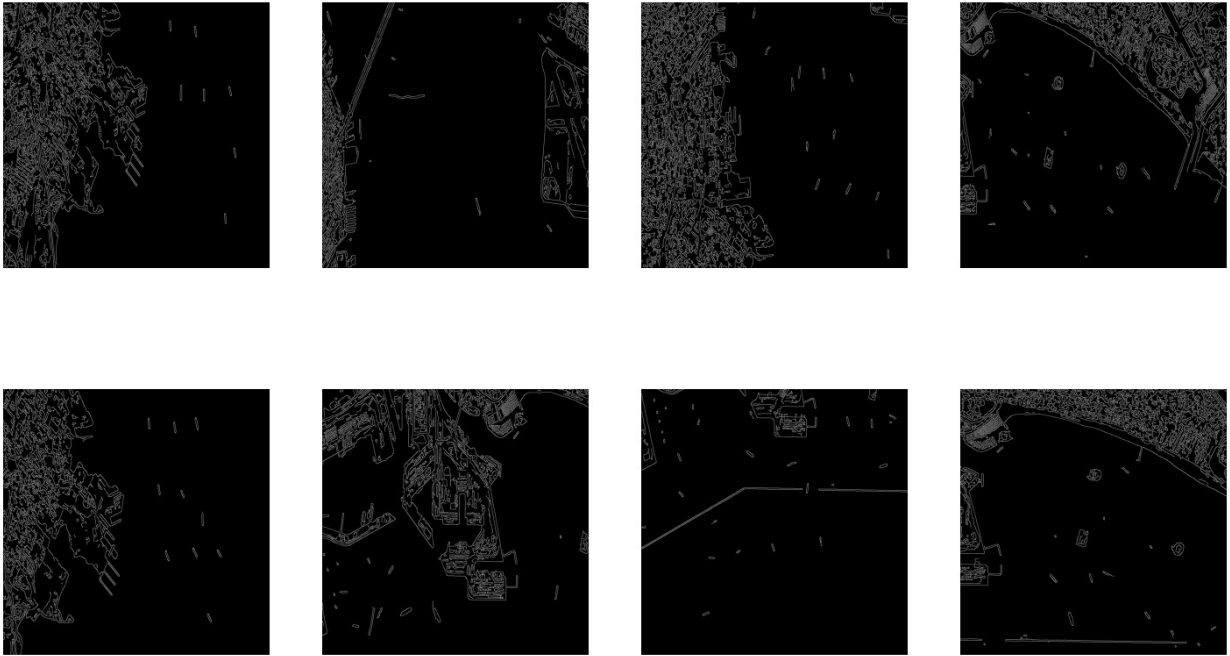
HSV Images



*## Applyin GaussinBlur and Canny Edge Detection on the HSV images
to recognize the edges and objects in the figure. This later helped
us in seperating water bodies with other objects.*

```
hsv_blur = [cv.GaussianBlur(image, (11,11), 0) for image in  
hsv_images]  
canny_hsv = [cv.Canny(image, 10, 100) for image in hsv_blur]  
plot_images(canny_hsv, title="HSV Canny", cmap="gray")
```

HSV Canny



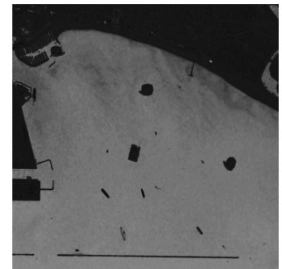
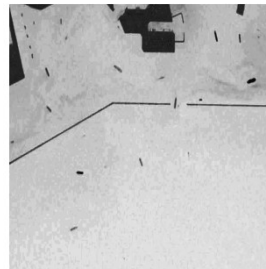
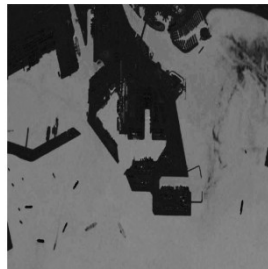
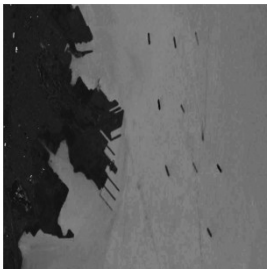
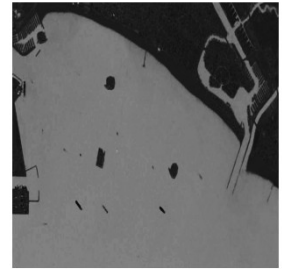
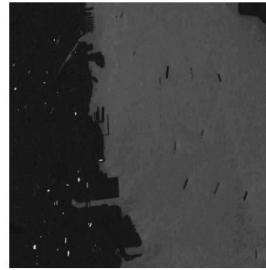
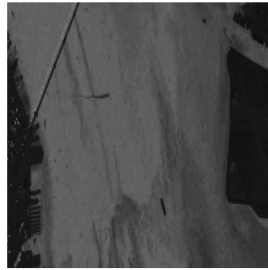
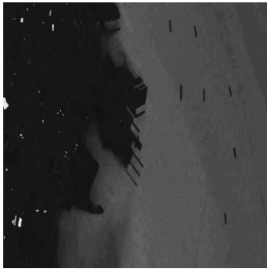
```
## median_smooth function is completely custom image smoothing  
## techniques that applies median smoothing on the images. We  
## are planning to implement this on the final feature maps or  
## predictions to remove any noise present in them.
```

```
def median_smooth(feature_map, kernel_size=3, pad=True, stride=1):  
    n = feature_map.shape[0]  
    padding = kernel_size - 2  
    op = floor((n + (2 * padding) - kernel_size) / stride) + 1  
    output = np.zeros((op, op), dtype=np.uint8)  
    feature_map = np.pad(feature_map, padding, 'reflect')  
    for row in range(op):  
        for col in range(kernel_size-1, op):  
            median = np.median(feature_map[row, col])  
            output[row, col] = median  
  
    return output
```

```
## Plotting the H cahnnel in HSV image to if we can gain any  
information.
```

```
plot_images([  
    image[:, :, 0]  
    for image in hsv_images  
], title="H in HSV Spectrum", cmap='gray')
```

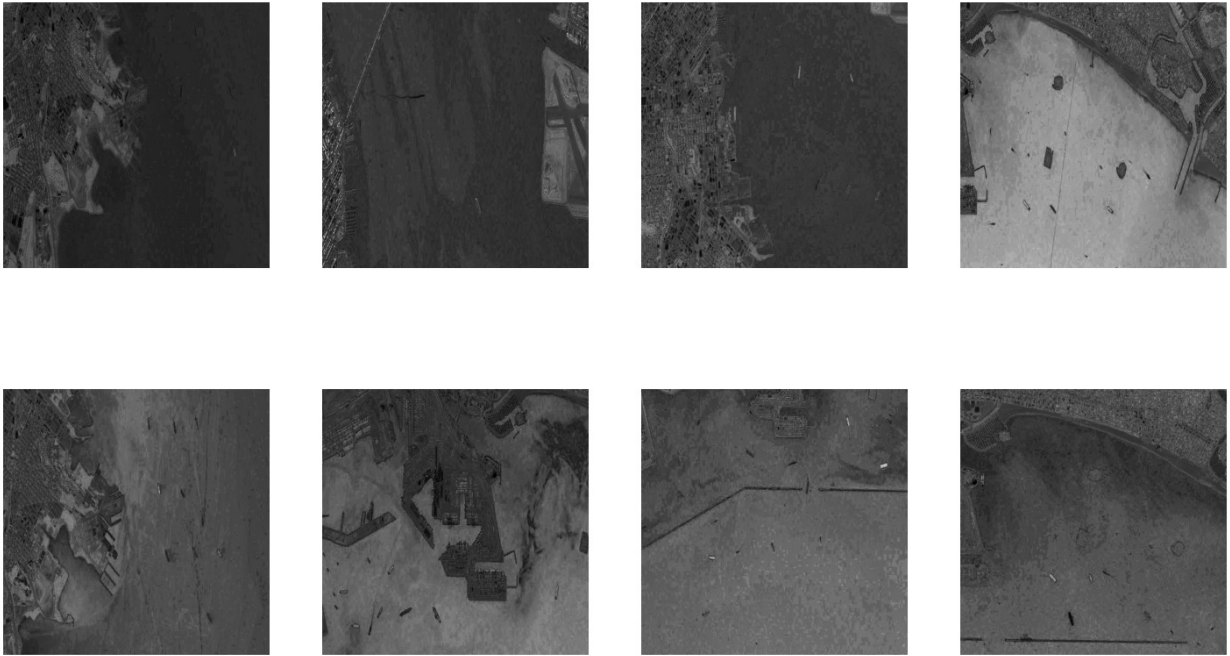
H in HSV Spectrum



Plotting the S cahnnel in HSV image to if we can gain any information.

```
plot_images([
    image[:, :, 1]
    for image in hsv_images
], title="S in HSV Spectrum", cmap='gray')
```

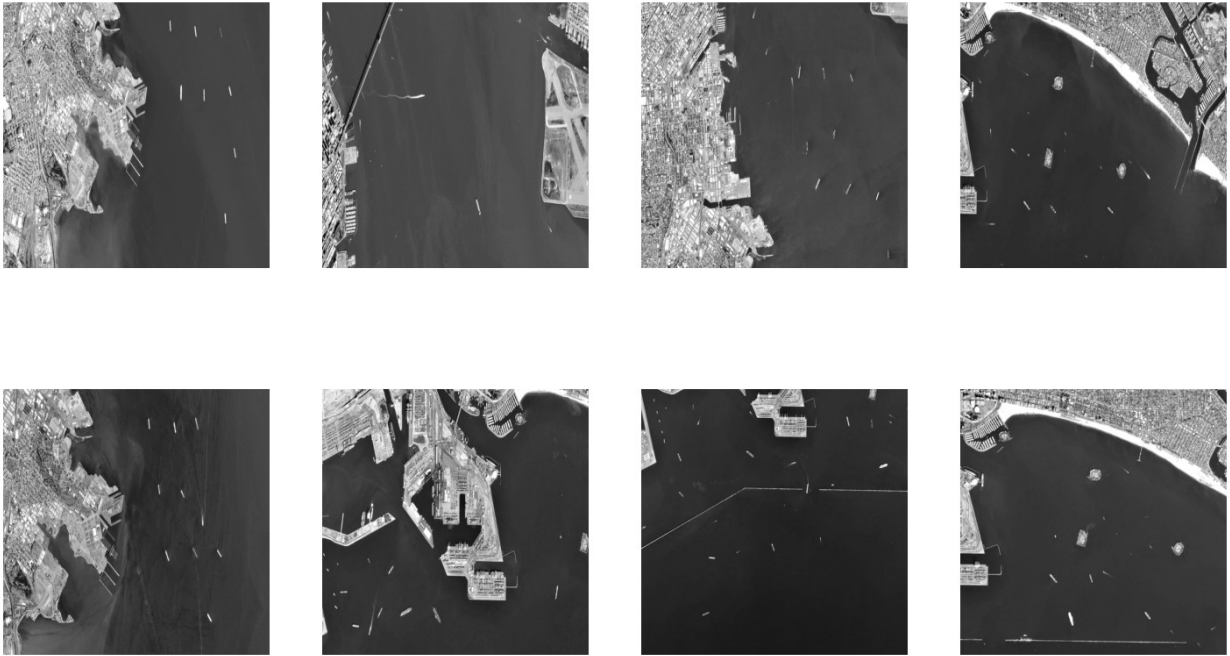
S in HSV Spectrum



Plotting the V cahnnel in HSV image to if we can gain any information.

```
plot_images([
    image[:, :, 2]
    for image in hsv_images
], title="V in HSV Spectrum", cmap='gray')
```

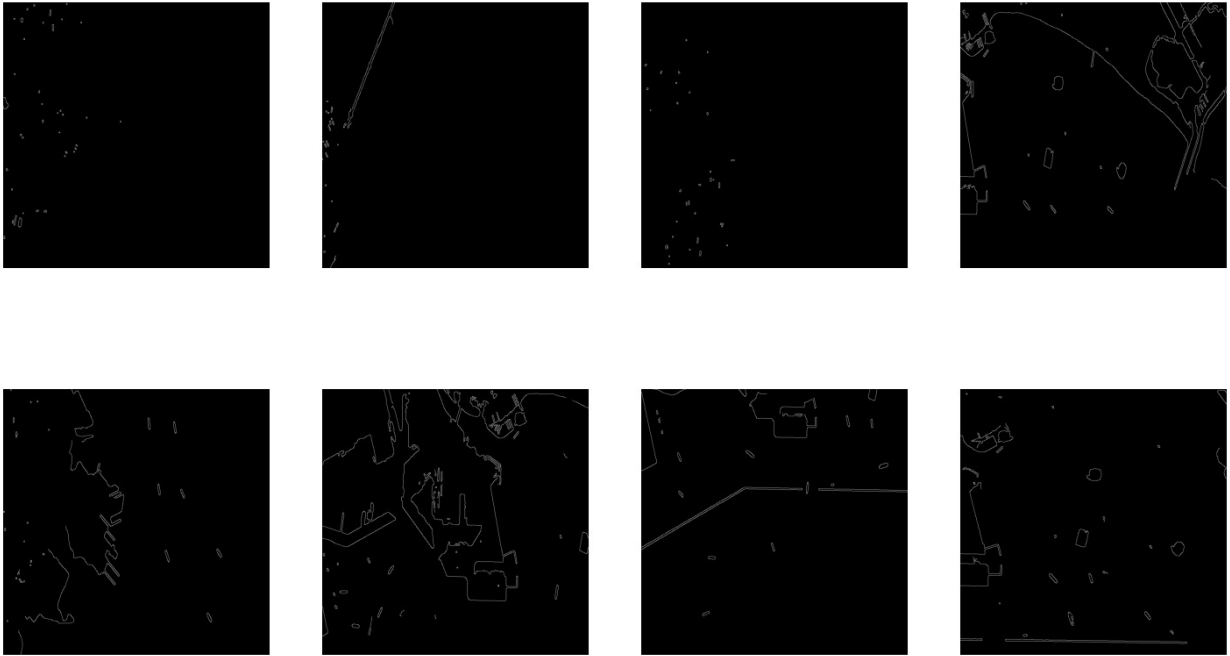

V in HSV Spectrum



Applying Gaussian Blur and Canny Edge Detection on the H channel of the images

```
H_blur = [cv.GaussianBlur(image[:, :, 0], (9, 9), 0) for image in  
hsv_images]  
canny_H = [cv.Canny(image, 10, 100) for image in H_blur]  
plot_images(canny_H, title="H in HSV Canny", cmap="gray")
```

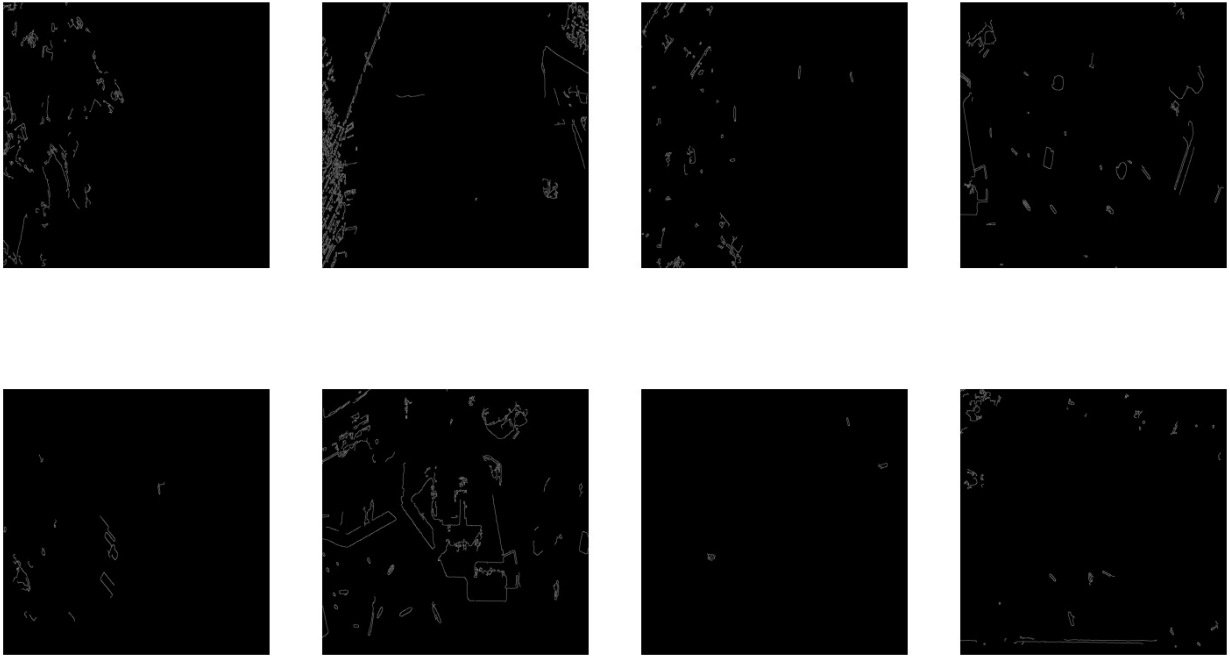
H in HSV Canny



Applying Gaussian Blur and Canny Edge Detection on the S channel of the images

```
S_blur = [cv.GaussianBlur(image[:, :, 1], (9, 9), 0) for image in  
hsv_images]  
canny_S = [cv.Canny(image, 10, 100) for image in S_blur]  
plot_images(canny_S, title="S in HSV Canny", cmap="gray")
```

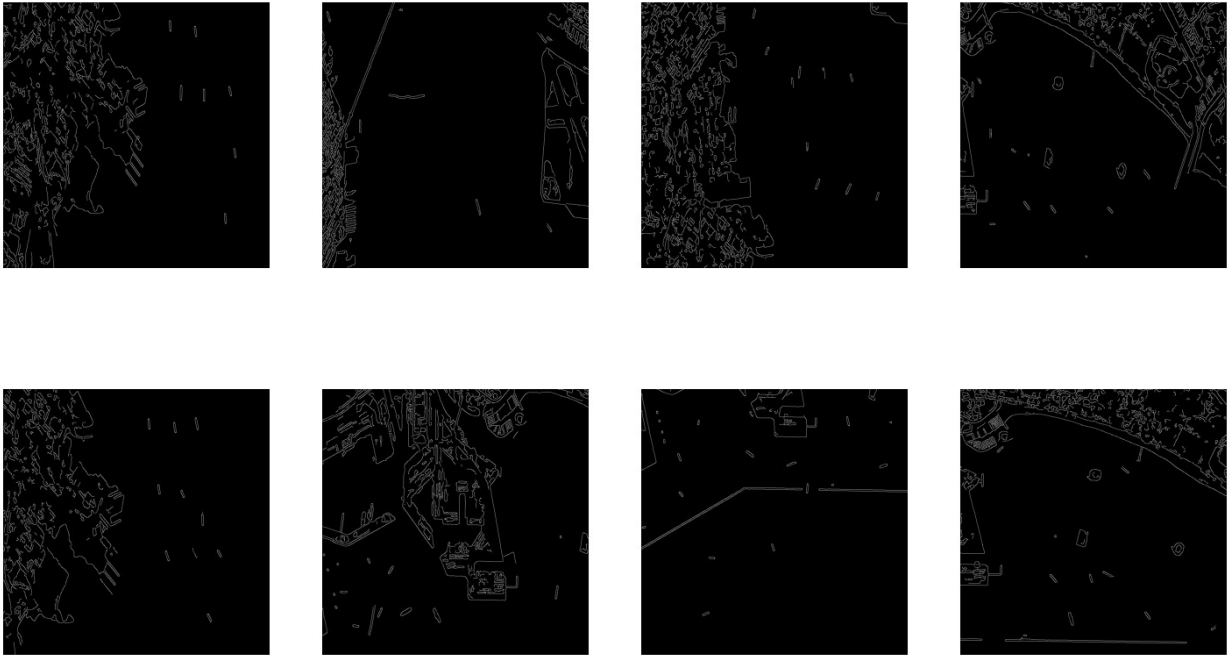

S in HSV Canny



Applying Gaussian Blur and Canny Edge Detection on the V channel of the images

```
V_blur = [cv.GaussianBlur(image[:, :, 2], (15, 15), 0) for image in  
hsv_images]  
canny_V = [cv.Canny(image, 10, 100) for image in V_blur]  
plot_images(canny_V, title="V in HSV Canny", cmap="gray")
```

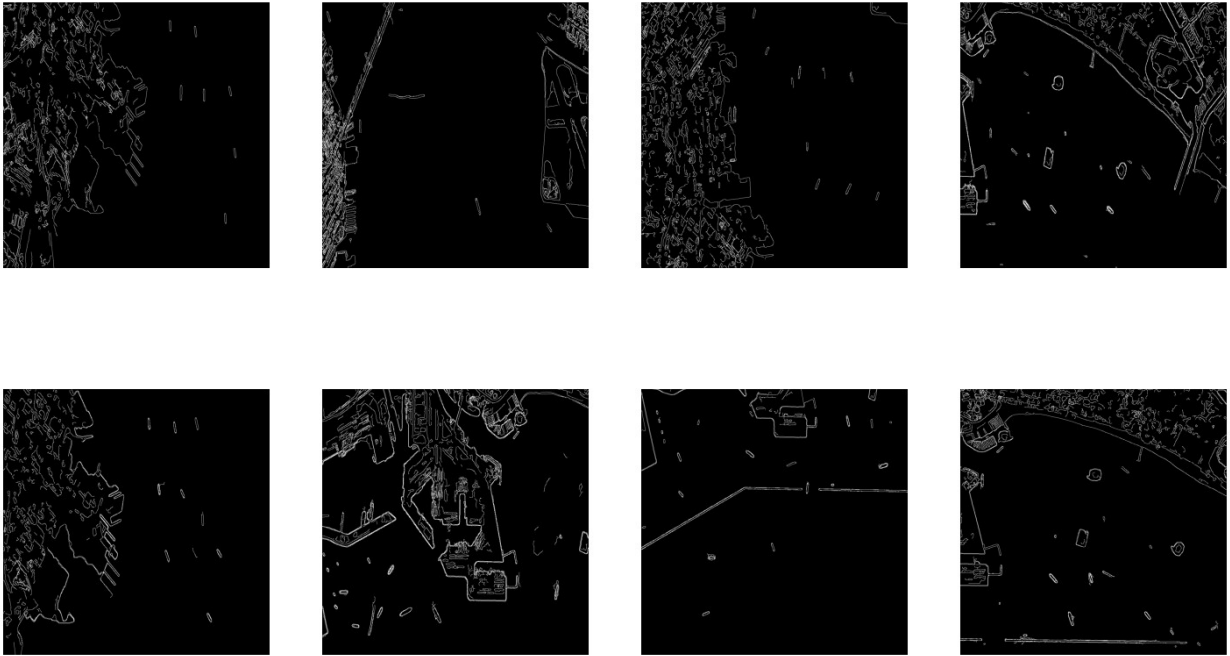
V in HSV Canny



```
## Combining the feature maps from the Canny Edge Detection of H,S and V  
## channels by super imposing them on each other. This gave us the final  
## Edge Detection map shown below.
```

```
HSV_canny = zip(canny_H, canny_S, canny_V)  
HSV_canny = [  
    np.clip(sum(list(HSV)), 0, 1)  
    for HSV in HSV_canny  
]  
  
plot_images(HSV_canny, title="Canny HSV", cmap='gray')
```

Canny HSV



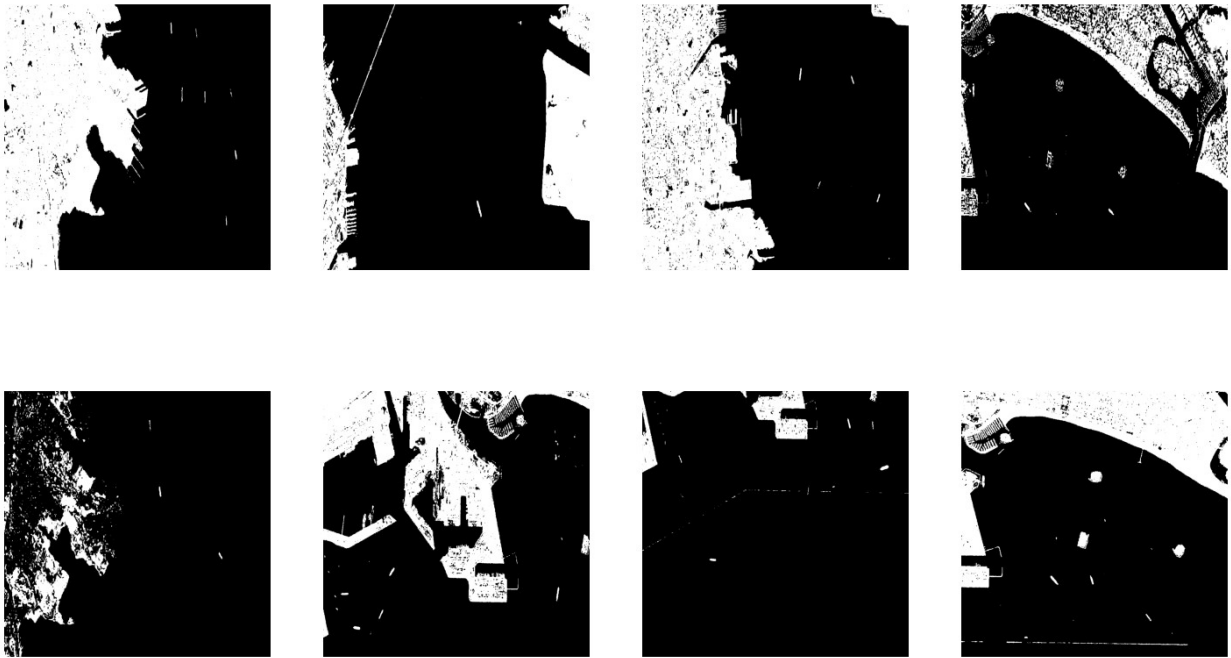
Applying Thresholding

```
## The edge detection on H,S and V channels gave better results, so  
## we are moving forward in that direction. Now we have applied  
thresholding  
## on separate channels of HSV and clipped them in that range.
```

```
## This part of the code clips H channel in the range of [0 - 20]
```

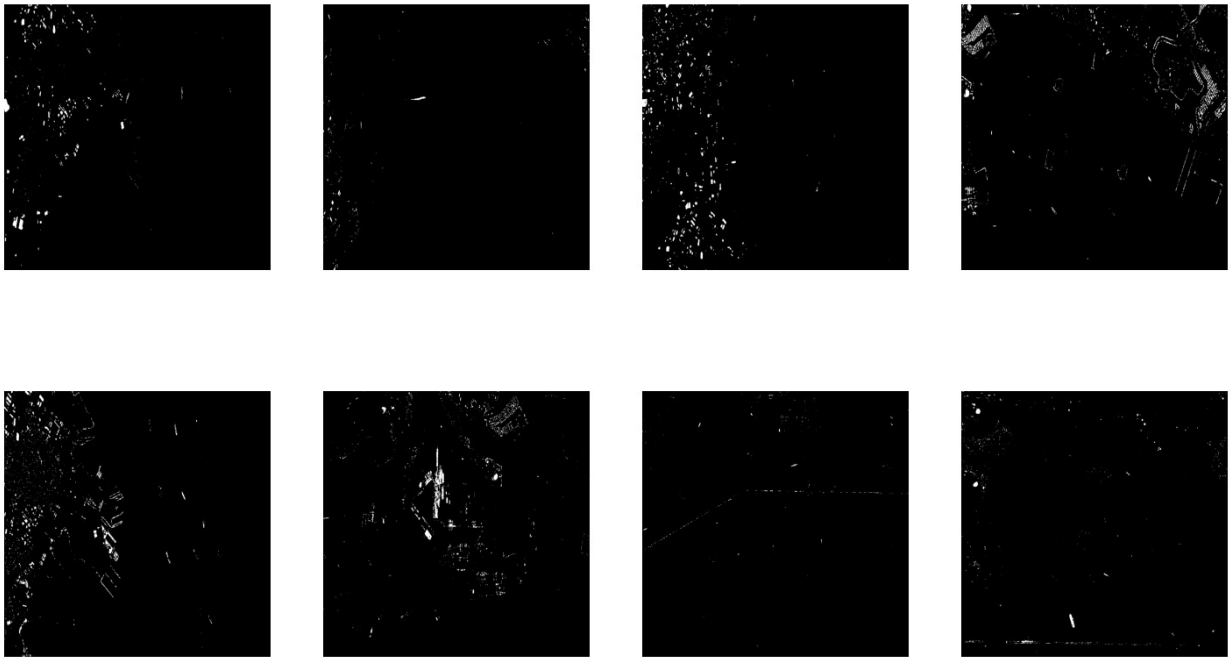
```
plot_images([  
    image[:, :, 0] < 20  
    for image in hsv_images  
], title="Thresholding H in HSV Spectrum", cmap='gray')
```

Thresholding H in HSV Spectrum



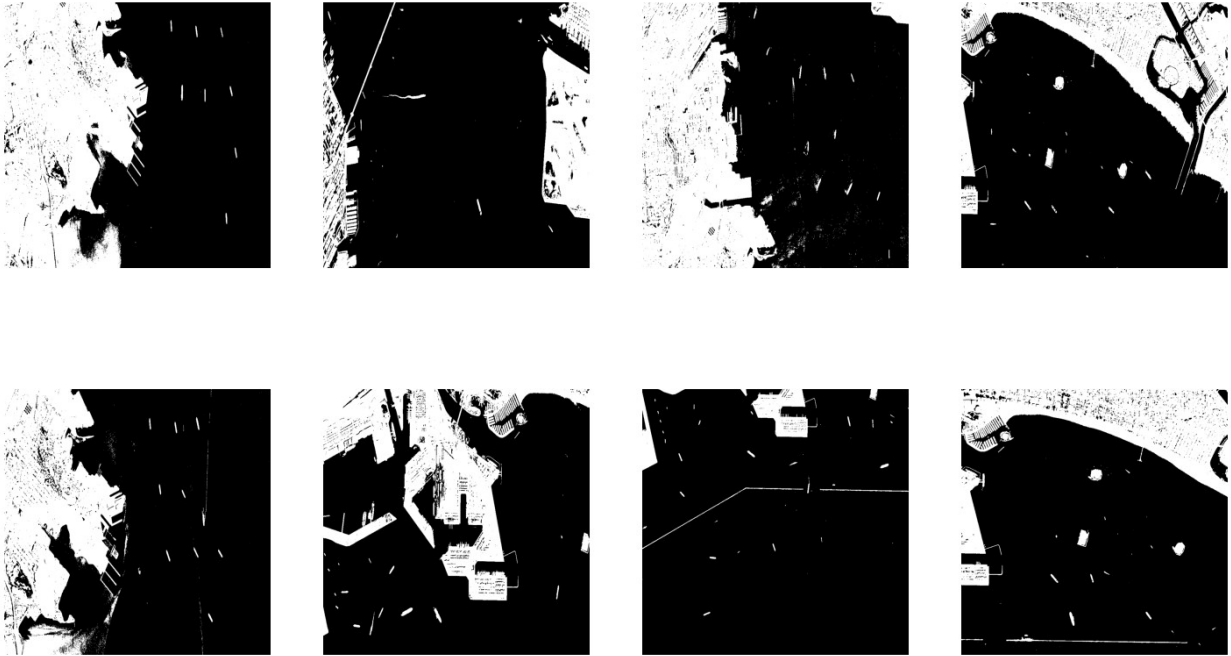
```
## This part of the code clips S channel in the range of [0 - 25]
plot_images([
    image[:, :, 1] < 25
    for image in hsv_images
], title="Thresholding S in HSV Spectrum", cmap='gray')
```

Thresholding S in HSV Spectrum



```
## This part of the code clips H channel in the range of [100 - 255]  
plot_images([  
    image[:, :, 2] > 100  
    for image in hsv_images  
], title="Thresholding V in HSV Spectrum", cmap='gray')
```

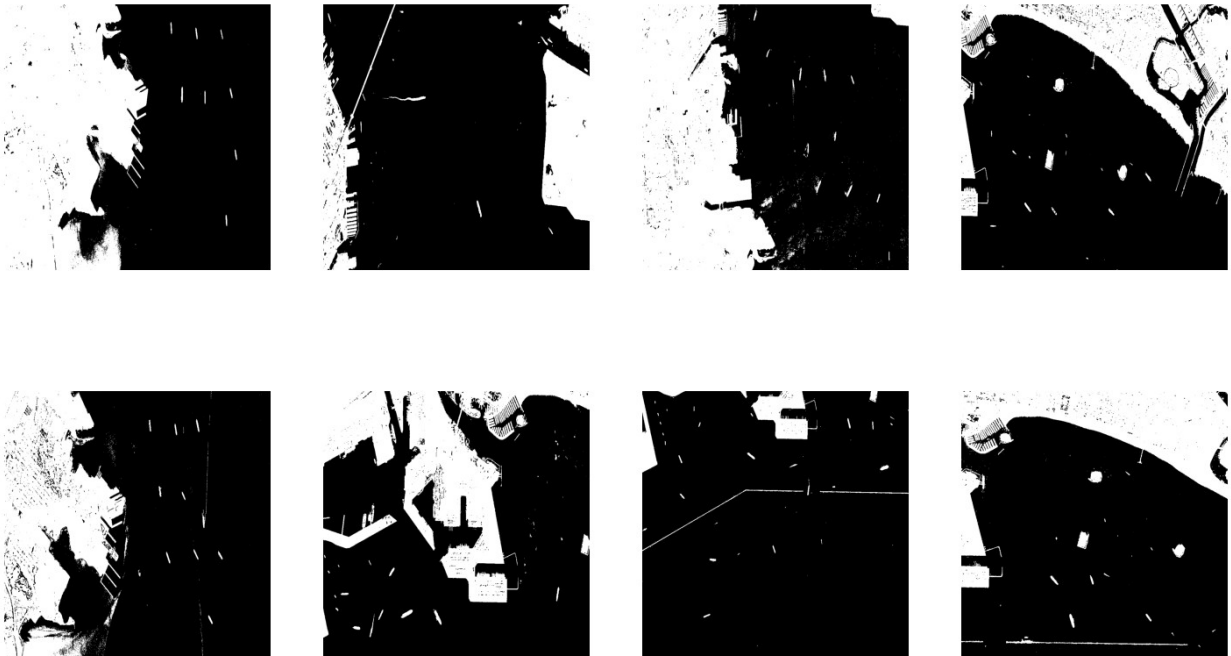
Thresholding V in HSV Spectrum



*## We are combining the thresholded feature maps of each channel
by super imposing them on each other. This gave us the final
feature maps.*

```
feature_maps = []  
for image in hsv_images:  
    map = (image[:, :, 0] < 20) + (image[:, :, 1] < 25) + (image[:, :, 2] >  
100)  
    map = np.clip(map, 0, 1)  
    feature_maps.append(map)  
  
plot_images(feature_maps, title="Feature Maps", cmap='gray')
```

Feature Maps



```
from sklearn.cluster import KMeans

kmeans = dict()
data = np.array(hsv_images)
data = data.reshape((-1, 3))
preds = dict()
for i in range(2, 6):
    kmeans[i] = KMeans(n_clusters=i)
    kmeans[i].fit(data)
    preds[i] = kmeans[i].predict(data)

/usr/local/lib/python3.10/dist-packages/sklearn/cluster/_kmeans.py:870: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/cluster/_kmeans.py:870: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/cluster/_kmeans.py:870: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning
  warnings.warn(
```



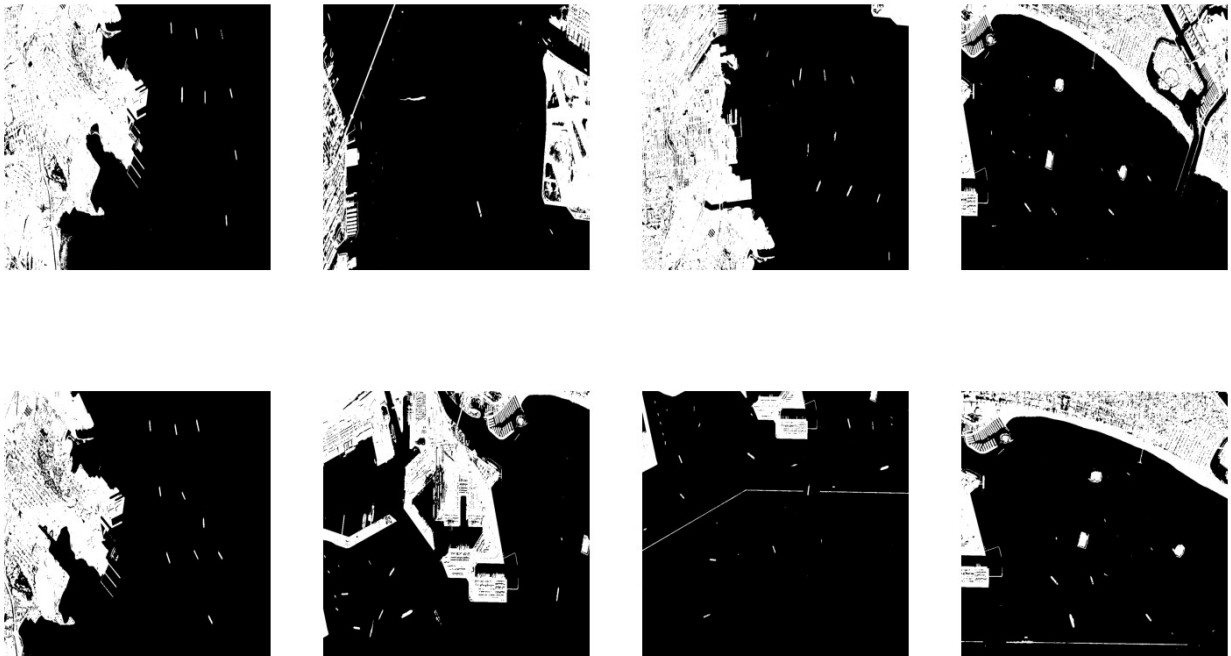
```

/usr/local/lib/python3.10/dist-packages/sklearn/cluster/_kmeans.py:870
: FutureWarning: The default value of `n_init` will change from 10 to
'auto' in 1.4. Set the value of `n_init` explicitly to suppress the
warning
    warnings.warn(

# preds[2] = preds[2] - 1
plot_images(preds[2].reshape((8,1000,1000)), title="KMeans with 2
Clusters", cmap='gray')

```

KMeans with 2 Clusters

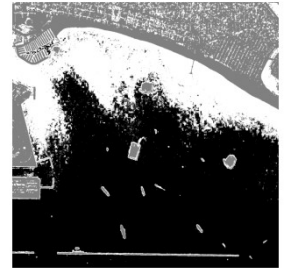
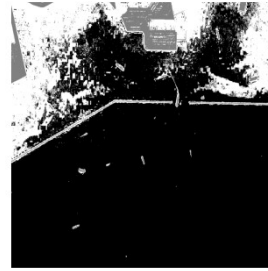


```

plot_images(preds[3].reshape((8,1000,1000)), title="KMeans with 3
Clusters", cmap='gray')

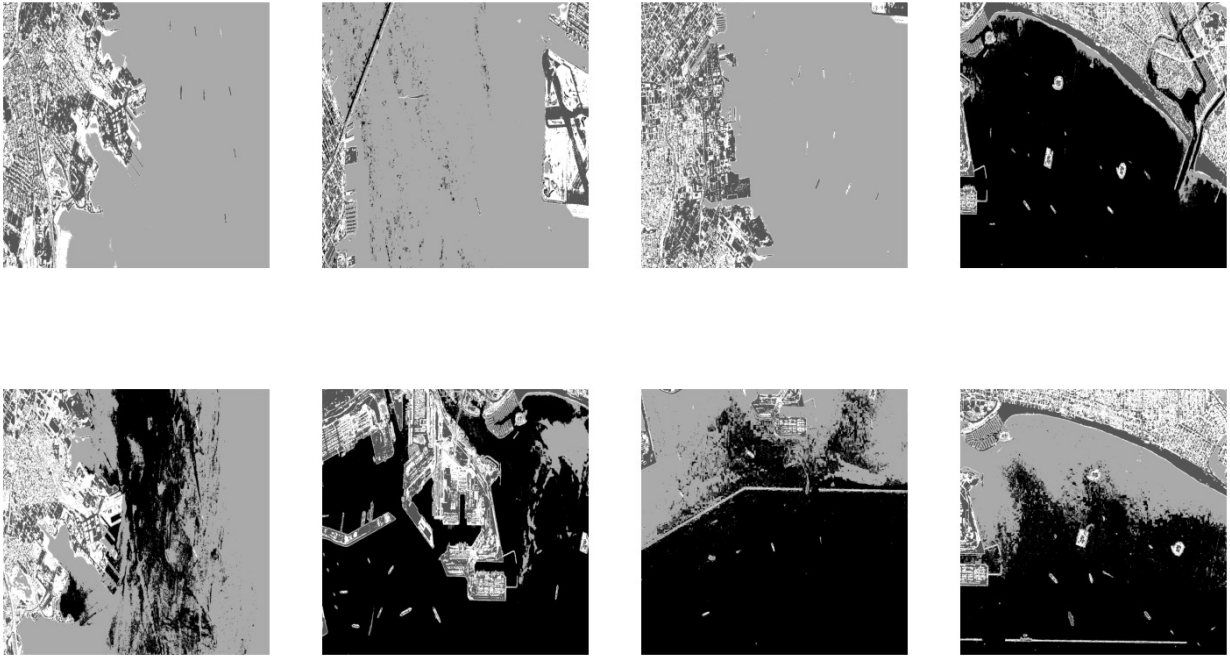
```

KMeans with 3 Clusters



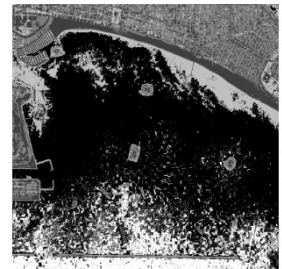
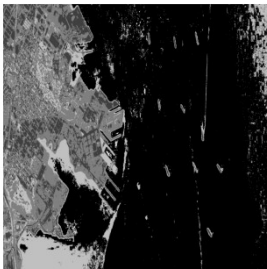
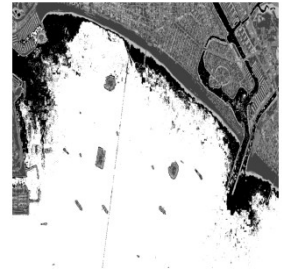
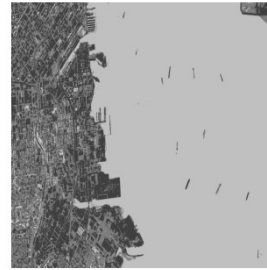
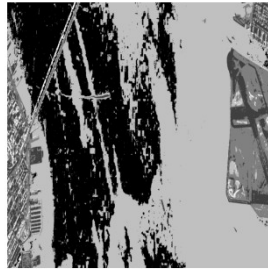
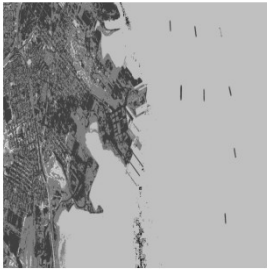
```
plot_images(preds[4].reshape((8,1000,1000)), title="KMeans with 4  
Clusters", cmap='gray')
```

KMeans with 4 Clusters



```
plot_images(preds[5].reshape((8,1000,1000)), title="KMeans with 5  
Clusters", cmap='gray')
```

KMeans with 5 Clusters



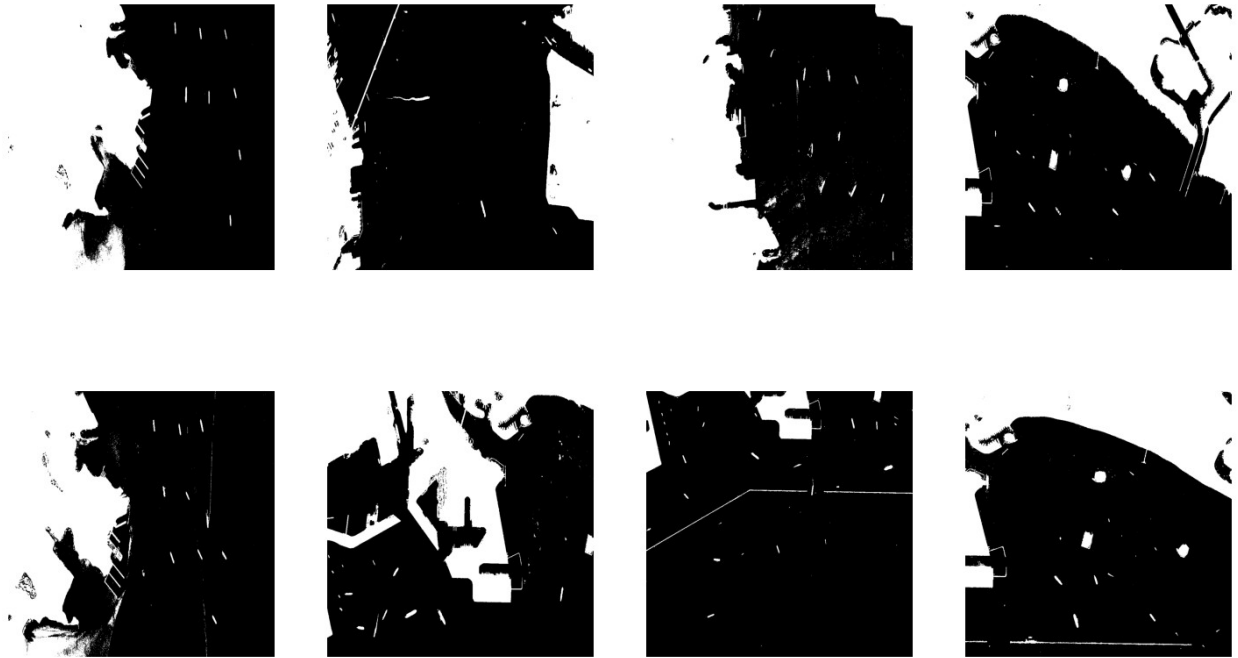
```
feature_maps = np.array(feature_maps)
cluster2 = preds[2].reshape((8,1000,1000))
blur_maps = [cv.blur(fmap, (25,25)).astype(np.int64) for fmap in
cluster2]
plot_images(blur_maps, title="Blur Maps", cmap="gray")
```

Blur Maps



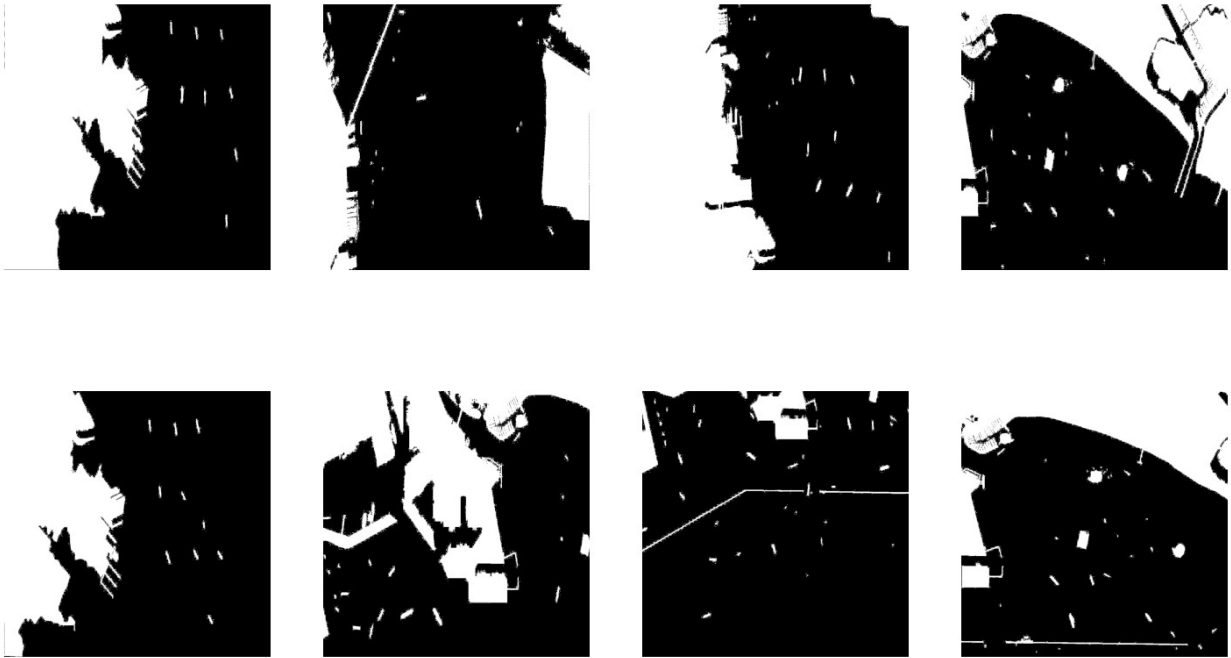
```
final_maps = np.maximum(feature_maps, blur_maps)
plot_images(final_maps, title="Final Predictions", cmap="gray")
```

Final Predictions



```
targets = load_images("/content/Masks", gray=True)
targets = [(target.astype(np.int64) > 1).astype(np.int64) for target
in targets]
plot_images(targets, title="Ground Truths", cmap='gray')
```

Ground Truths



```
from sklearn.metrics import precision_score, recall_score,
accuracy_score

for idx, (pred, truth) in enumerate(zip(final_maps, targets)):
    print(f"Image {idx+1} Scores:")
    print(f"Precision: {precision_score(truth, pred, average='micro')}\t"
    Recall: {recall_score(truth, pred, average='micro')})"
```

```
Image 1 Scores:
Precision: 0.907109507948744      Recall: 0.9826370561774739
Image 2 Scores:
Precision: 0.9903656714802226    Recall: 0.9121162876484252
Image 3 Scores:
Precision: 0.984941538363287     Recall: 0.9800828776826023
Image 4 Scores:
Precision: 0.9743328118268598    Recall: 0.9315600069372182
Image 5 Scores:
Precision: 0.9178759490063394    Recall: 0.9578681993609163
Image 6 Scores:
Precision: 0.9910150738843098    Recall: 0.9220914241345503
Image 7 Scores:
Precision: 0.9925112694488876    Recall: 0.7790629101908041
Image 8 Scores:
Precision: 0.9905091239287003    Recall: 0.956658243663726
```