COMPSCI 371D Homework 2

```
In [1]: %matplotlib inline
    import numpy as np
    from matplotlib import pyplot as plt
    from mpl_toolkits.mplot3d import Axes3D # Do not remove this import
    from math import floor, ceil
    import warnings
```

Part 1: Functions in Machine Learning

Problem 1.1 (Exam-Style)

```
Domain: X = [x_0, x_2, \dots, x_{24}] where each x_i is limited to 0 < x_i < 2^{16} - 1
```

Range: Y = [Covered in Vegetation, Not Covered in Vegetation]

Problem 1.2 (Exam-Style)

h is a Classifier.

Problem 1.3 (Exam-Style)

We can use the zero-one loss function

$$\ell(y,\hat{y}) = \left\{egin{array}{ll} 0 & ext{for } y = \hat{y} \ 1 & ext{otherwise} \end{array}
ight.$$

because our range Y is binary so we can only care about if the classifier is correct or not.

Problem 1.4 (Exam-Style)

The predictor will most likely work well since the hardest part of getting a training set is labeling the training set. Since the set samples provided are already annotated, and there are a lot of them, the classifier will have a lot of samples to test with, which will allow for it to better deal with new data that it has not come across.

Part 2: The L_1 Loss

Problem 2.1

```
In [2]: def plot_function(function, ax, variable_range=(-1, 1), samples=101):
    t = np.linspace(variable_range[0], variable_range[1], num=samples)
    x, y = np.meshgrid(t, t)
    ax.plot_surface(x, y, function(x, y), cmap=plt.get_cmap('viridis'))
    ax.set_xticks([])
    ax.set_yticks([])
    ax.set_zticks([])
```

```
In [3]: fig = plt.figure(figsize=(12, 5))
    subplot_1 = fig.add_subplot(1, 2, 1, projection='3d')
    variable_range=(-1, 1)
    samples=101
    def l(x, y):
        return abs(0 - x) + abs(0-y)
    plot_function(1, subplot_1, variable_range=variable_range, samples=samples)
```



Problem 2.2 (Exam-Style)

The roots of the first derivative of $L_1(m)$ cannot be bound using standard calculas techniques.

Problem 2.3 (Exam-Style)

 $L_1(m)$ can be minimized one component at a time because you are essentially finding the gradient of $L_1(m)$. By minimizing each point $L_1(m)$ with respect to each dimention, you can find the overall minimum point, which will minimize the function the most.

Problem 2.4 (Exam-Style)

Yes

Problem 2.5 (Exam-Style)

Because there are more points on the right of m than on the left of m, as m moves from x_2 towards x_3 , the new distances of the points on the right of m will affect $L_1(m)$ more than the new distances from the points left of m. When m moves a Δx distance to the right, for the 5 points on the right of m, their average distance will change by $-\Delta x$, while the two points to the left of m will have their lengths changed by $+\Delta x$, so the overall change will be $-3\Delta x$, which is a negative linear amount.

Problem 2.6

Givien a set of size N, the $\mathop{argmin}_{m} L_1(m)$ can be found when $m=(x_1+x_2+x_3+\ldots+x_n)/N$

Part 3: Intuition in Many Dimensions

```
In [4]: from math import gamma, sqrt

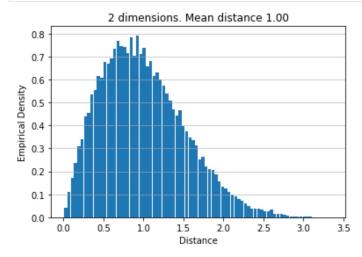
def gaussian_sample(n, d):
    mean = np.zeros(d)
    mean_distance = 2 * gamma((d + 1) / 2) / gamma(d / 2) if d < 200 else sqrt(2 * d)
    covariance = np.eye(d) / pow(mean_distance, 2)
    return np.random.multivariate_normal(mean, covariance, size=n)

In [5]: from matplotlib import pyplot as plt

def show_histogram(x, d):
    density, _, _ = plt.hist(x=x, bins='auto', density=True, rwidth=0.9)
    mean = np.mean(x)
    plt.grid(axis='y', alpha=0.75)
    plt.xlabel('Distance')
    plt.ylabel('Empirical Density')
    format_string = '{} dimension{}. Mean distance {:.2f}'
    plt.title(format_string.format(d, 's' if d > 1 else '', mean))
```

Problem 3.1

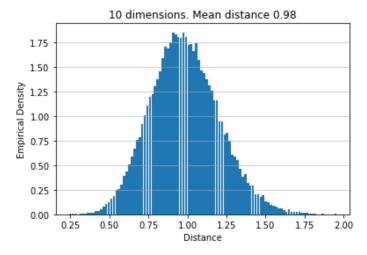
```
In [7]: points = gaussian_sample(300,2)
    dis = distances(points)
    show_histogram(dis, 2)
```



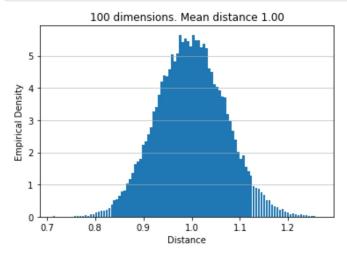
Problem 3.2

```
In [8]: def show_histogram(x, d):
    density, _, _ = plt.hist(x=x, bins='auto', density=True, rwidth=0.9)
    mean = np.mean(x)
    plt.grid(axis='y', alpha=0.75)
    plt.xlabel('Distance')
    plt.ylabel('Empirical Density')
    format_string = '{} dimension{}. Mean distance {:.2f}'
    plt.title(format_string.format(d, 's' if d > 1 else '', mean))
```

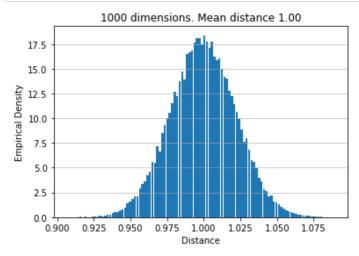
```
In [9]: d1 = distances(gaussian_sample(300,10))
    show_histogram(d1,10)
```



In [10]: d2 = distances(gaussian_sample(300,100))
show_histogram(d2,100)



In [11]: d3 = distances(gaussian_sample(300,1000))
show_histogram(d3,1000)



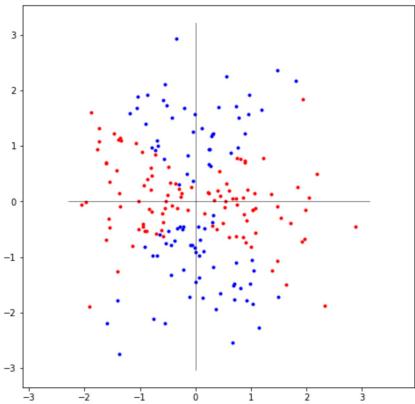
Problem 3.3

While the four plots may look the same, the amount of data behind each plot is very different, with the d=1000 data set having exponentially more data than the d=2 data set.

Problem 3.4

Despite each plot having the same number of points, when comparing the distances between individual points, since the higher dimension points have more data per point to compare against, it greatly increases the amount of time it takes to compute the distance.

Part 4: Data Transformations



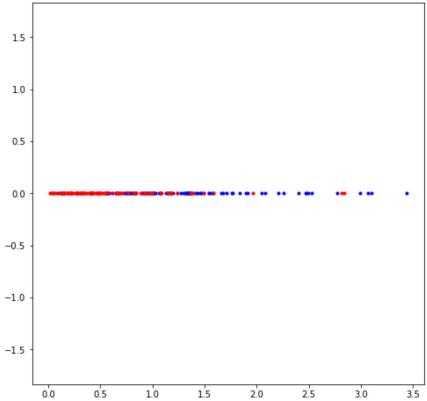
Problem 4.1

Since the negative points tend to follow the line $y=-\frac{1}{2}x$, we can use a transformation that takes the vertical distance from the line. Then be finding the distance that most negative points are away from the line, we can set that value as the threshold.

The transformation for a point $x=(x_1,y_1)$ would look like $z=\sqrt{(-rac{1}{2}x_1-y_1)^2}$

```
In [13]: def transform(x):
    z = []
    for i in x:
        n = sqrt(np.power(-.5*i[0] - i[1],2))
        z.append(n)
    return np.array(z)

z = transform(x)
plt.figure(figsize=(8, 8))
positive = z[y > 0]
    negative = z[y <= 0]
plt.plot(positive, np.zeros(len(positive)), '.b')
plt.plot(negative, np.zeros(len(negative)), '.r')
y_limits = plt.ylim(-1,1)
plt.axis('equal')
plt.show()</pre>
```



Problem 4.2

(i)

```
In [14]: def h(z, threshold):
    y_hat = []
    for i in z:
        if i > threshold:
            y_hat.append(1)
        else:
            y_hat.append(-1)
    return np.array(y_hat)
```

(ii)

```
In [15]: def risk(y, y_hat):
    risk = 0
    for i in range(len(y)):
        if y[i] != y_hat[i]:
            risk += 1
    return risk
```

(iii)

```
In [16]: def find_threshold(z, y):
             z sorted = np.sort(z)
             best_t = -10000
             min_risk = 10000
             t_and_r = [[]]
             t_his = []
             r_his = []
             for i in range(len(z)-1):
                 t = (z_sorted[i] + z_sorted[i+1])/2
                  y_hat = h(z_sorted, t)
                  new_risk = risk(y, y_hat)
                  t_and_r.append([t, new_risk])
                  t_his.append(t)
                  r his.append(new risk)
                  if new risk < min risk:</pre>
                      min_risk = new_risk
                     best_t = t
             return best t, min risk, t his, r his
```

```
In [17]: t, r, t_his, r_his = find_threshold(z, y)
```

```
In [18]: print("Best Threshold: ",round(t,4))
```

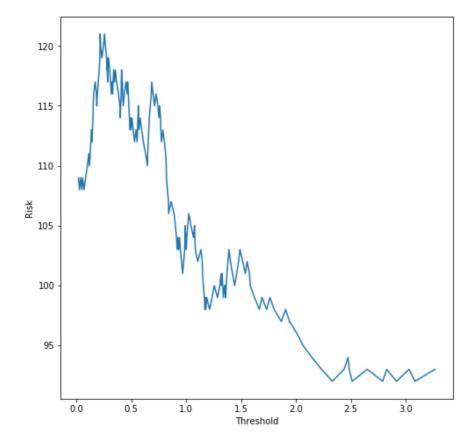
Best Threshold: 2.3319

```
In [19]: print("Minimum Risk: ", round(r,4))
```

Minimum Risk: 92

```
In [20]: plt.figure(figsize=(8, 8))
    plt.plot(t_his, r_his)
    plt.xlabel('Threshold')
    plt.ylabel('Risk')
```

Out[20]: Text(0, 0.5, 'Risk')



```
In []:
In []:
```