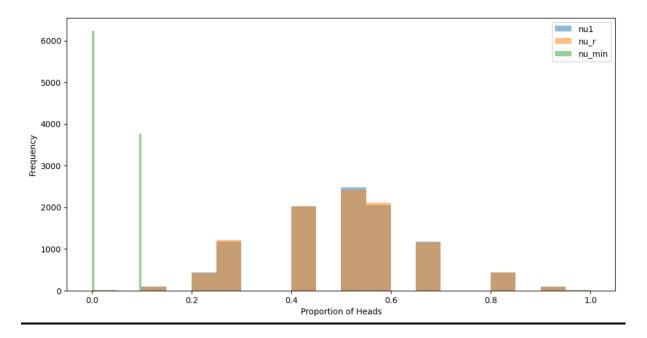
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Problem 1:

Plotting the histogram of the distribution nu1, nu_r, nu_min over the 100000 samples.

In this experiment of flipping 1000 different fair coins 10 times each and calculating the proportions of heads for the first coin, a randomly chosen coin, and the coin that came up heads the least among all of them.

The coin_experiment() code uses the numpy In [7]: ▶ import numpy as np library to generate random choices for each def coin experiment(): coin flip, and then calculates the proportion of num_flips = 10 heads for each coin. It returns the proportions # Initialize arrays to store the results results = np.zeros((num_coins, num_flips))
proportions = np.zeros((num_coins)) for the first coin (nu1), a randomly chosen for i in range(num_oins):
 # Flip the coin num flips times
flips = np.random.choice([0, 1], size=num_flips)
 results[i] = flips coin (nu r), and the coin that came up heads the least (nu_min). # Calculate the proportion of heads
proportion_heads = np.mean(flips)
proportions[i] = proportion_heads # Calculate the proportion of heads for the first coin nu1 = proportions[0] # Calculate the proportion of heads for a randomly chosen coin nu r = np.random.choice(proportions) # Calculate the proportion of heads for the coin that came up heads the least nu min = np.min(proportions) return nu1, nu_r, nu_min Next, we repeated the entire experiment In [9]: | import matplotlib.pyplot as plt 100,000 times and plotted the histograms of def repeat_experiment(num_repeats):
 nul_values = np.zeros((num_repeats))
 nu_r_values = np.zeros((num_repeats))
 nu_min_values = np.zeros((num_repeats)) the distribution of nu1, nu r, and nu min. Below is the output. for i in range(num_repeats): nu1, nu_r, nu_min = coin_experiment()
nu1_values[i] = nu1
nu_r_values[i] = nu_r nu_min_values[i] = nu_min # Plot the histograms # Plot the histograms
plt.figure(figsize=(12, 6))
plt.hist(nu1_values, bins=20, alpha=0.5, label='nu1')
plt.hist(nu_r_values, bins=20, alpha=0.5, label='nu_r')
plt.hist(nu_min_values, bins=20, alpha=0.5, label='nu_min')
plt.xlabel('Proportion of Heads')
plt.ylabel('Frequency')
plt.legend()
nlt.show() plt.show() repeat experiment(10000)



Observation from the histogram1:

The range of possible proportions of heads is divided into intervals, called bins. As we are flipping a coin 10 times, the proportion of heads ranges from 0 to 1 in increments of 0.1. Each interval represents a possible outcome of the experiment.

For each interval, we counted how many times the proportion of heads occurred in our experiment. This count represents the frequency of occurrence for that proportion.

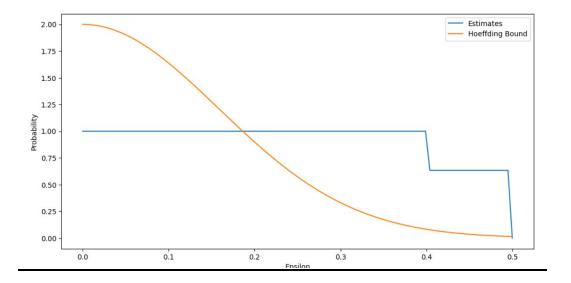
The histogram plot consists of bars, where the height of each bar represents the frequency of occurrence of the corresponding proportion of heads. The x-axis represents the range of possible proportions, while the y-axis represents the frequency of occurrence.

By looking at the histogram we can assess the distribution of the observed proportion of heads(observed empirical risk) for random nu_r is more concentrated at 0.5 and even for the first sample it is showing at 0.5. The proportions are more common at 0.5. Hence the distribution of proportions of heads aligns with our expectations compared to the true population proportion (nu = 0.5)

Plotting the histogram for estimates of Pr(|v-u| > epsilon) as a function of epsilon.

We calculated the estimates of Pr(|v-u| > epsilon) as a function of epsilon and plotted them together with the Hoeffding bound. It uses the coin_experiment function to generate the nu_min values and checks if they exceed epsilon to update the estimates

```
def calculate hoeffding bound(epsilon, num flips):
    return 2 * np.exp(-2 * epsilon**2 * num_flips)
def plot_hoeffding_bound(num_repeats):
    epsilons = np.linspace(0, 0.5, 100)
    hoeffding_bounds = calculate_hoeffding_bound(epsilons, num_flips=10)
    nu values = np.zeros((num repeats))
    hoeffding_estimates = np.zeros_like(epsilons)
    for i in range(num repeats):
        nu1, nu_r, nu_min = coin_experiment()
        nu_values[i] = nu_min
        for j, epsilon in enumerate(epsilons):
            if abs(nu min - 0.5) > epsilon:
                hoeffding_estimates[j] += 1
    # Normalize the estimates
    hoeffding_estimates /= num_repeats
    # Plot the estimates and the Hoeffding bound
    plt.figure(figsize=(12, 6))
plt.plot(epsilons, hoeffding_estimates, label='Estimates')
    plt.plot(epsilons, hoeffding_bounds, label='Hoeffding_Bound')
plt.xlabel('Epsilon')
    plt.ylabel('Probability')
    plt.legend()
    plt.show()
plot_hoeffding_bound(10000)
```



Observations from the histogram2:

The observed proportion of heads (v) and the true proportion of heads (u = 0.5), the histogram is plotted for Pr(|v-u| > e) as a function of e.

The range of possible deviations (e) is divided into intervals that range from 0 to 0.5 into smaller intervals, each representing a different magnitude of deviation.

The observed proportion of heads for a coin (v) falling within the Hoeffding bound (as most concentration in the histogram is on 0.5), it means that deviation from the true population (u) is within the acceptable range. In other words, these coins obey the Hoeffding bound, and the observed proportions of heads are consistent with what would be expected (u = 0.5) based on random chance.

Also, for more samples, the whole epsilon term decreases therefore the probability also decreases.

Problem 3:

Frohims

(a) If t is a non-negative random variable for any
$$\alpha > 0$$
, $Pr(t > \kappa) \le F(t)$.

Read:

Pr(t > \alpha) = \int f(t) dt \quad \text{(iber f(t) is the polarity } \rightarrow eq 0)}

Now, condition the expectation on the event t > \int \text{

Eft] = \int t \text{ f(t) dt} = \int t \text{ f(t) dt} + \int t \text{ f(t) dt}.

Since t is non negative:

\[
\text{

Eft] \geq \int t \text{ f(t) dt} \geq \int \int \text{ f(t) dt} \quad \text{

From eq 0}
\]

\[
\text{

Eft] \geq \text{

Pr(t \geq \alpha) \leq \text{

From eq 0}
\]

\[

Pant (c): If u,..., un are independent and identically distributed, each with mean u and variance -, and se is the sample mean of ee, ,..., in , then for 20: Px ((u-px)2x) < =2 Apply part (b) to the random variable re with a and note that Proof: Var (u) = = to n fid means il and variance a. $P_{\lambda}((2l-\mu)^2 > x) \leq \frac{V_{\lambda}(2l)}{x} = \frac{-2^{\frac{1}{2}}}{n^{\frac{1}{2}}}$

Problem 4:

Problem 4:

Part 1

The perception prediction is given by
$$h(x) = sign(\omega \cdot x)$$
 where

 $w = (\omega_0, \omega_0, \omega_0)$ and $x = (1, x, x)$. The regions on the plane when $h(x) = +1$
 $w = (\omega_0, \omega_0, \omega_0)$ and $x = (1, x, x)$. The regions on the plane when $h(x) = +1$

and $h(x) = -1$ are separated by line.

If we set $h(x) = 0$, we can divid the equation of the decision boundary.

 $h(x) = sign(\omega \cdot x)$
 $0 = \omega \cdot x$
 $0 = \omega \cdot x$
 $0 = \omega \cdot x$
 $0 = \omega \cdot x + \omega_0 \cdot x$

Part (b)

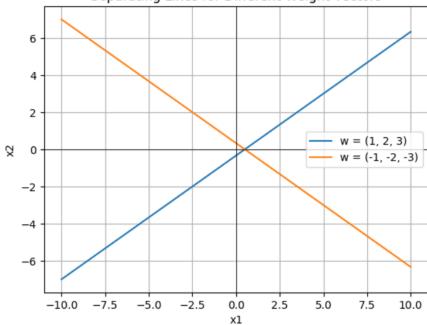
The plot will show two lines, one for each weight vector. The lines separate the regions where h(x) = +1 and h(x) = -1. The lines have opposite slopes and intercepts, illustrating the difference between the two weight vectors of their signs and magnitude.





1 # Problem 4

Separating Lines for Different Weight Vectors



Problem 5
To derive the estimaters we and = 2 that maximize the log-likelihood dunction of a given distribution, we mill find the maximum by taking partial derivatives with usped to we and = 2, setting them to zero, and solving for the parameters.

Estimating 11:

$$\ln p(x|u, \alpha^{2}) = \frac{-1}{2\sigma^{2}} \sum_{i=1}^{n} (x_{i} - u)^{2} - \frac{n \ln 2}{2} - \frac{n \ln (2\pi)}{2}$$

$$= \frac{1}{2\sigma^{2}} \left(x_{i} - u\right) - \frac{n \ln 2}{2} - \frac{n \ln (2\pi)}{2}$$

$$= \frac{1}{2\sigma^{2}} \left(x_{i} - u\right) - 0 - 0 = 0$$

$$= \frac{1}{2\sigma^{2}} \left(x_{i} - u\right) = 0$$

$$\Rightarrow \begin{cases} \lambda \\ \leq \lambda \\ \leq 1 \end{cases} (\lambda_0 - \lambda_0) = 0$$

$$\sum_{i=1}^{n} x_i - nu = 0 \Rightarrow u = \frac{1}{n} \sum_{i=1}^{n} x_i$$

so the maximum likelihood estimate for it is the sample men

Estimating of:

$$\ln p(x|u,\sigma) = \frac{1}{2\sigma^2} \int_{\sigma_0}^{\pi} (x_0 - u)^2 - \frac{n}{2} \ln \sigma^2 - \frac{n}{2} \ln(2\pi)$$
partial desirvative w.r. f σ^2

$$\frac{\partial}{\partial z^{2}} \ln p(x|\mu,\sigma^{2}) = -\frac{1}{2} (-2) \sigma^{-3} \frac{x}{2} (x_{0} - \mu)^{2} - \frac{1}{2} \frac{1}{2^{2}} (2\sigma^{2} - 0)$$

$$-\frac{1}{2}(-\frac{1}{2})^{\frac{1}{2}} = \frac{1}{2}(x_{0} - \mu)^{2} - \frac{n}{2}(\frac{1}{2})^{2} = 0$$

$$-\frac{1}{2}(-\frac{1}{2})^{\frac{1}{2}} = \frac{n}{2}(x_{0} - \mu)^{2} = \frac{n}{2}$$

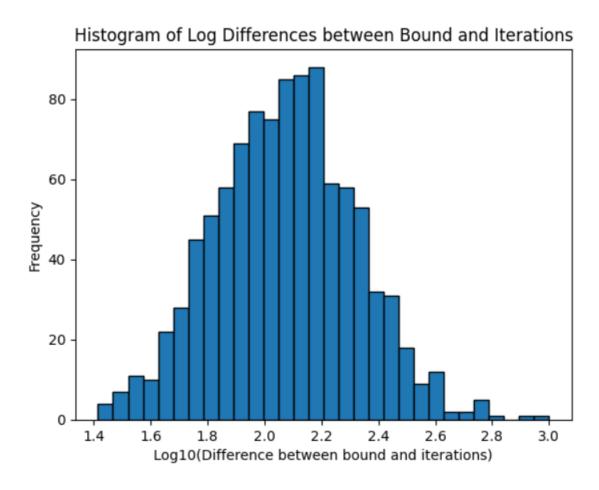
$$-\frac{1}{2}(x_{0} - \mu)^{2} = \frac{1}{2}(x_{0} - \mu)^{2}$$

$$-\frac{1}{2}(x_{0} - \mu)^{2} = \frac{1}{2}(x_{0} - \mu)^{2}$$

6. Problem 6 The likelihood function L measurer the likelihood of observing the given data { (x;, y;)}; given the parameters of the model. we model the probability as P(y=+1/x) using a function h(x)The lihelihood function $\phi(x)$ a single data point (x_i^0, y_i^0) is given by: $L(\phi) = \begin{cases} h(x_i^0) & \text{if } y_i^0 = -1 \\ 1 - h(x_i^0) & \text{if } y_i^0 = -1 \end{cases}$ $(0) = \underbrace{2}_{i=1} \left(| (y_i = + \overline{y_i}) + | (x_i) \right) + | (y_i = -\overline{y_i}) | (y_i = + \overline{y_i})$ $h(z) = \sigma(w \cdot z)$, where $\sigma(z)$ is the sigmond function. where I[A] is the indicated genetism. -(t) = 1 1+e-t $l(0) = \sum_{i=1}^{n} \left[l(y_i = +1) \log(-(w \cdot x_i)) + l(y_i = -1) \log(1 - -(w \cdot x_i)) \right]$ $= \sum_{i=1}^{n} \left[1 \left[y_{i}^{n} = + i \right] \log \left(\frac{1}{1 + e^{\omega_{i} \cdot x_{i}^{n}}} \right) + 1 \left[y_{i}^{n} = -1 \right] \log \left(1 - \frac{1}{1 + e^{\omega_{i} \cdot x_{i}^{n}}} \right) \right]$ $= \sum_{i=1}^{\infty} \left[i \left[y_i = +1 \right] \log \left(\frac{1}{1 + e^{-\omega \cdot x_i}} \right) + i \left[y_i = -1 \right] \log \left(\frac{e^{-\omega \cdot x_i}}{1 + e^{-\omega \cdot x_i}} \right) \right]$ = \(\frac{1}{2} \left[\left[\frac{1}{2} \cdot \frac{1}{1 + e^{-\omega_1 \cdot \chi_1}} \right] + 1 \left[\frac{1}{2} \cdot \cdot \left[\frac{1}{1 + e^{-\omega_1 \chi_2}} \right] \right] $= \sum_{i=1}^{n} \left(\left[y_i = + i \right] \log \left(\frac{1}{1 + e^{\omega_i \cdot x_i}} \right) - \left[\left[y_i = - i \right] \log \left(1 + e^{\omega_i \cdot x_i} \right) \right]$ This expression represents the minimized when h(x) = -(w.x).

Problem 2:

- 1. Generated an 11-dimensional weight vector w*, where the first dimension is 0 and the other 10 dimensions are sampled independently at random from the uniform (0, 1) distribution.
- 2. Repeated the steps 1000 times:
 - a. Generated a random training set with 100 examples, where each dimension of each training example is sampled independently at random from the uniform (-1, 1) distribution, and the examples are all classified by w*.
 - b. By running the perceptron learning algorithm on the training set, starting with the zero-weight vector, and keeping track of the number of iterations it takes to learn a hypothesis that correctly separates the training data.
- 3. Plotted a histogram of the number of iterations the algorithm takes to learn a linear separator.
- 4. Calculate the difference between the number of iterations and the bound on the number of errors derived in class for each experiment.
- 5. Plot a histogram of the logarithm of this difference.



The code generated a new weight vector and training set for each experiment. By running the perceptron learning algorithm on the training set and keeping track of the number of iterations required to learn a hypothesis that correctly separates the training data. It then plots a histogram of the number of iterations and calculates the difference between the number of iterations and the bound on the number of errors derived in class. Finally, plotted a histogram of the logarithm of this difference.

By analyzing the histograms, we examined the distribution of the number of iterations and the difference between the iterations and the bound. This analysis shows how the perceptron learning algorithm performs in different scenarios where the algorithm takes more iterations to correctly learn a hypothesis that separates the training data. The algorithm took more iterations between 2.0 and 2.2 to correctly separate the training data.

Note that the results may vary between different runs of the experiment due to the randomness involved in generating the weight vector and training set. Therefore, to run the experiment multiple times and analyzed the distribution of the results to draw meaningful conclusions.