# Problem Set #2: Supervised Learning II

## Problem 1 Logistic Regression: Training stability

- (a) The most notable difference in training the logistic regression model on datasets A and B is that the algorithm does not converge on dataset B.
- (b) To investigate why the training procedure behaves unexpectedly on dataset B, but not on A, we print the value of  $\theta$  after every 10000 iterations. We notice that for data set B, although the normalized  $\frac{\theta}{\|\theta\|}$  almost stop changing after several tens of thousands of iterations, each component of the unnormalized  $\theta$  keeps increasing. We also notice that dataset A is not linearly separable while dataset B is linearly separable.

From the code, we notice that the algorithm calculates the gradient of loss function as

$$\nabla_{\theta} J(\theta) = -\frac{1}{m} \sum_{i=1}^{m} \frac{y^{(i)} x^{(i)}}{1 + \exp(y^{(i)} \theta^{T} x^{(i)})}.$$

From this, we know that the algorithm uses gradient descent to minimize the loss function

$$J(\theta) = -\frac{1}{m} \sum_{i=1}^{m} \log \frac{1}{1 + \exp(-y^{(i)}\theta^{T}x^{(i)})}.$$

Hence, for a dataset that is linearly separable, that is,  $y^{(i)}\theta^Tx^{(i)} > 0$  for all i, a  $\theta$  with larger norm always leads to a smaller loss, preventing the algorithm from converging. However, on a dataset that is not linearly separable, there exists i such that  $y^{(i)}\theta^Tx^{(i)} < 0$ . By plotting  $f(z) = \log(1 + e^{-z})$  in Figure 1, we notice that negative margin dominates when scaling  $\theta$  to a larger norm. Hence, we cannot always increase  $\theta$  to a larger norm while minimizing  $J(\theta)$ .

- (c) Consider the following modifications
  - i. Using a different constant learning rate will not make the algorithm converge on dataset B, since scaling  $\theta$  to larger norm still always decreases the loss.
  - ii. Decreasing the learning rate over time will make the algorithm converge for dataset B, since in this way the change of  $\theta$  converge to 0.

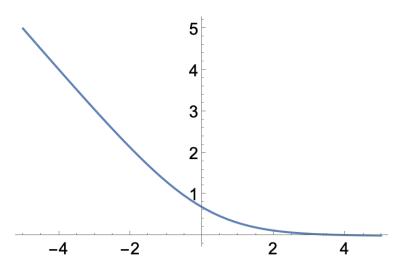


Figure 1: Plot of  $f(z) = \log(1 + e^{-z})$  for  $-5 \le z \le 5$ .

- iii. Linear scaling the input features does not help, since it does not change the dataset's linear separability.
- iv. Adding a regularization term  $\|\theta\|_2^2$  helps, since now scaling  $\theta$  to larger norm penalize the algorithm.
- v. Adding zero-mean Gaussian noise to the training data or labels helps as long as it makes the dataset not linearly separable.
- (d) Support vector machines, which uses hinge loss, are not vulnerable to datasets like B. In SVM, geometric margin is considered, instead of functional margin considered here. In other words,  $\theta$  is normalized, so for linearly separable datasets like B, the algorithm will still converge.

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#### Problem 2 Model Calibration

Try to understand the output  $h_{\theta}(x)$  of the hypothesis function of a logistic regression model, in particular why we might treat the output as a probability.

When probabilities outputted by a model match empirical observation, the model is well-calibrated. For example, if a set of examples  $x^{(i)}$  for which  $h_{\theta}(x^{(i)}) \approx 0.7$ , around 70% of those examples should have positive labels. In a well-calibrated model, this property holds true at every probability value.

Suppose training set  $\{x^{(i)}, y^{(i)}\}_{i=1}^m$  with  $x^{(i)} \in \mathbb{R}^{n+1}$  and  $y^{(i)} \in \{0, 1\}$ . Assume we have an intercept term  $x^{(i)}_0 = 1$  for all i. Let  $\theta$  be the maximum likelihood parameters learned after training logistic regression model. In order for model to be well-calibrated, given any range of probabilities (a, b) such that  $0 \le a < b \le 1$ , and trianing examples  $x^{(i)}$  where the model outputs  $h_{\theta}(x^{(i)})$  fall in the range (a, b), the fraction of positives in that set of examples should be equal to the average of the model outputs for those examples. That is,

$$\frac{\sum_{i \in I_{a,b}} P(y^{(i)} = 1 \mid x^{(i)}; \theta)}{|\{i \in I_{a,b}\}|} = \frac{\sum_{i \in I_{a,b}} \mathbb{I}\{y^{(i)} = 1\}}{|\{i \in I_{a,b}\}|},$$

where 
$$P(y^{(i)} = 1 \mid x; \theta) = h_{\theta}(x) = 1/(1 + \exp(-\theta^T x)), I_{a,b} = \{i : h_{\theta}(x^{(i)}) \in (a,b)\}.$$

(a) For the described logistic regression model over the range (a, b) = (0, 1), we want to show the above equality holds. Recall the gradient of log-likelihood

$$\frac{\partial \ell}{\partial \theta_j} = \sum_{i=1}^m (y^{(i)} - h_{\theta}(x^{(i)})) x^{(i)}_j.$$

For a maximum likelihood estimation,  $\frac{\partial \ell}{\partial \theta} = 0$ . Hence  $\frac{\partial \ell}{\partial \theta_0} = 0$ . Since  $x^{(i)}_{0} = 1$ , we have

$$\sum_{i=1}^{m} y^{(i)} - h_{\theta}(x^{(i)}) = 0.$$

The desired equality follows immediately since  $i \in I_{0,1}$  for all i.

(b) A perfectly calibrated model — that is, the equality holds for any  $(a, b) \subset [0, 1]$  — does not imply that the model achieves perfect accuracy. Consider  $(a, b) = (\frac{1}{2}, 1)$ , the above equality implies

$$\frac{\sum_{i \in I_{a,b}} P(y^{(i)} = 1 \mid x^{(i)}; \theta)}{|\{i \in I_{a,b}\}|} = \frac{\sum_{i \in I_{a,b}} \mathbb{I}\{y^{(i)} = 1\}}{|\{i \in I_{a,b}\}|} < 1.$$

This shows that the model does not have perfect accuracy.

For the converse direction, a perfect accuracy does not imply perfectly calibrated. Consider again  $(a, b) = (\frac{1}{2}, 1)$ , then we have

$$\frac{\sum_{i \in I_{a,b}} \mathbb{I}\{y^{(i)} = 1\}}{|\{i \in I_{a,b}\}|} = 1 > \frac{\sum_{i \in I_{a,b}} P(y^{(i)} = 1 \mid x^{(i)}; \theta)}{|\{i \in I_{a,b}\}|}.$$

(c) Discuss what effect of  $L_2$  regularization in the logistic regression objective has on model calibration. For  $L_2$  regularization in logistic regression, the gradient becomes

$$\frac{\partial \ell}{\partial \theta_j} = \sum_{i=1}^m (y^{(i)} - h_{\theta}(x^{(i)})) x^{(i)}{}_j - 2C\theta_j = 0.$$

Hence, the equality does not hold unless  $\theta_0 = 0$ .

The interval (0,1) is the only range for which logistic regression is guaranteed to be calibrated. When GLM assumptions hold, all ranges  $(a,b) \subset [0,1]$  are well calibrated. In addition, when test set has same distribution and when model has not overfit or underfit, logistic regression are well-calibrated on test data as well. Thus logistic regression is popular when we are interested in level of uncertainty in the model output.

## Problem 3 Bayesian Interpretation of Regularization

In Bayesian statistics, almost every quantity is a random variable. Joint distribution of all the random variables are called model (e.g.  $p(x, y, \theta)$ ). Every unknown quantity can be estimated by conditioning the model on all observed quantities. Such conditional distribution  $p(\theta \mid x, y)$  is called *posterior distribution*. A consequence of this approach is that we are required to endow a *prior distribution*  $p(\theta)$ .

In purest Bayesian interpretation, we are required to keep the entire posterior distribution over the parameters all the way until prediction to come up with the *posterior predictive distribution*, and the final prediction will be the EV of the posterior predictive distribution. However, this is computationally very expensive.

The compromise is to estimate a point value of the parameters instead of the full distribution, which is the mode of the posterior distribution. Estimating the mode of posterior distribution is also called *maximum a posteriori estimation* (MAP). That is,

$$\theta_{\text{MAP}} = \operatorname*{argmax}_{\theta} p(\theta \mid x, y).$$

Compare this to the maximum likelihood estimation (MLE):

$$\theta_{\text{MLE}} = \operatorname*{argmax}_{\theta} p(y \mid x, \theta).$$

In this problem, explore connections between MAP estimation and common regularization techniques that are applied with MLE estimation. In particular, we will show how choice of prior distribution over  $\theta$  is equivalent to different kinds of regularization.

(a) Assume that  $p(\theta) = p(\theta \mid x)$ , we have

$$p(\theta \mid x, y) = \frac{p(y \mid \theta, x)p(\theta \mid x)}{p(y \mid x)} = \frac{p(y \mid \theta, x)p(\theta)}{p(y \mid x)}.$$

Since  $p(y \mid x)$  does not depend on  $\theta$ ,

$$\theta_{\text{MAP}} = \operatorname*{argmax}_{\theta} p(y \mid \theta, x) p(\theta).$$

Note that the assumption  $p(\theta) = p(\theta \mid x)$  will be valid for models such as linear regression where the input x are not explicitly modeled by  $\theta$ . Note also that this means x and  $\theta$  are marginally independent, but not conditionally independent when y is given.

(b) Now we show that MAP estimation with a zero-mean Gaussian priori over  $\theta$ , specifically  $\theta \sim \mathcal{N}(0, \eta^2 I)$ , is equivalent to applying  $L_2$  regularization with MLE estimation. Specifically, we need to show

$$\theta_{\text{MAP}} = \underset{\theta}{\operatorname{argmin}} - \log p(y \mid x, \theta) + \lambda \|\theta\|_{2}^{2}.$$

Recall the definition of multivariate normal, we have

$$\begin{split} \theta_{\text{MAP}} &= \operatorname*{argmax}_{\theta} p(\theta \mid x, y) \\ &= \operatorname*{argmax}_{\theta} p(y \mid \theta, x) p(\theta) \\ &= \operatorname*{argmax}_{\theta} p(y \mid \theta, x) \exp \left( -\frac{1}{2\eta^2} \|\theta\|_2^2 \right), \end{split}$$

where we have ignored some of the constants. Taking the negative log on both sides, it follows that

$$\theta_{\text{MAP}} = \underset{\theta}{\operatorname{argmin}} - \log p(y \mid x, \theta) + \frac{1}{2\eta^2} \|\theta\|_2^2,$$

as desired, where  $\lambda = \frac{1}{2\eta^2}$ .

(c) Now consider a specific instance, a linear regression model given by  $y = \theta^T x + \varepsilon$ , where  $\varepsilon \sim \mathcal{N}(0, \sigma^2)$ . Like before, assume a Gaussian prior on this model such that  $\theta \sim \mathcal{N}(0, \eta^2 I)$ . Let X be the design matrix of all training examples where each row is one example input, and y be the column vector of all the example outputs. We want to derive a closed form expression for  $\theta_{\text{MAP}}$ .

For this model, the likelihood of an example  $(x^{(i)}, y^{(i)})$  is

$$p(y^{(i)} \mid x^{(i)}, \theta) = \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(y^{(i)} - \theta^T x^{(i)})^2}{2\sigma^2}\right).$$

Hence,

$$\log p(y^{(i)} \mid x^{(i)}, \theta) = -\frac{(y^{(i)} - \theta^T x^{(i)})^2}{2\sigma^2} + C,$$

where C is some constant, and  $\theta_{\text{MAP}}$  is given by

$$\theta_{\text{MAP}} = \underset{\theta}{\operatorname{argmin}} \sum_{i=1}^{m} \frac{(y^{(i)} - \theta^{T} x^{(i)})^{2}}{2\sigma^{2}} + \frac{1}{2\eta^{2}} \|\theta\|_{2}^{2}$$
$$= \underset{\theta}{\operatorname{argmin}} \frac{1}{2\sigma^{2}} (y - X\theta)^{T} (y - X\theta) + \frac{1}{2\eta^{2}} \theta^{T} \theta.$$

Set the gradient of the function to 0, we have

$$0 = -\frac{1}{\sigma^2} X^T (y - X\theta_{\text{MAP}}) + \frac{1}{\eta^2} \theta_{\text{MAP}}$$

It follows that

$$\theta_{\text{MAP}} = \eta^2 (\eta^2 X^T X + \sigma^2 I)^{-1} X^T y.$$

(d) Now consider the Laplace distribution, whose density

$$f(z \mid \mu, b) = \frac{1}{2b} \exp\left(-\frac{|z - \mu|}{b}\right).$$

As before, consider a linear regression model given by  $y = \theta^T x + \varepsilon$  where  $\varepsilon \sim \mathcal{N}(0, \sigma^2)$ Auumer a Laplace prior on this model where  $\theta \sim \mathcal{L}(0, bI)$ . We want to show that  $\theta_{\text{MAP}}$ in this case is equivalent to the solution of linear regression with  $L_1$  regularization, whose loss is specified as

$$J(\theta) = \|X\theta - y\|_{2}^{2} + \gamma \|\theta\|_{1}.$$

Following the same approach,  $\theta_{\text{MAP}}$  for this model is given by

$$\theta_{\text{MAP}} = \underset{\theta}{\operatorname{argmin}} \frac{1}{2\sigma^2} ||y - X\theta||_2^2 + \frac{1}{b} ||\theta||_1.$$

Hence, this is equivalent to the solution of linear regression with  $L_1$  regularization, with  $\gamma = \frac{2\sigma^2}{b}$  in the above loss function  $J(\theta)$ .

**Note:** closed form solution for linear regression problem with  $L_1$  regularization does not exist. To optimize this, use gradient descent with a random initialization and solve it numerically.

• Linear regression with  $L_2$  regularization is also called *Ridge regression*, and  $L_1$  regularization is called *Lasso regression*. These regularizations can be applied to any generalized linear models just as above. Regularization techniques are also called *weight decay* and *shrinkage*. The Gaussian and Laplace priors encourages the parameter values to be closer to their mean (i.e. zero), which results in the shrinkage effect.

• Lasso regression is known to result in sparse parameters.

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## Problem 4 Constructing kernels

Choosing kernel  $K(x, z) = \phi(x)^T \phi(z)$ . Mercer's theorem tells us K is a Mercer kernel iff for any finite set  $\{x^{(i)}\}_{i=1}^m$ , the square matrix  $K \in \mathbb{R}^{m \times m}$  whose entries are  $K_{ij} = K(x^{(i)}, x^{(j)})$  is symmetric and positive semidefinite.

Let  $K_1$ ,  $K_2$  be kernels over  $\mathbb{R}^n \times \mathbb{R}^n$ , let  $a \in \mathbb{R}^+$  be a positive real number, let  $f : \mathbb{R}^n \to \mathbb{R}$  be a real-valued function, let  $\phi : \mathbb{R}^n \to \mathbb{R}^d$  be a function mapping from  $\mathbb{R}^n$  to  $\mathbb{R}^d$ , let  $K_3$  be a kernel over  $\mathbb{R}^d \times \mathbb{R}^d$ , and let p a polynomial with *positive* coefficients.

For each K below, prove or give counter examle to show whether it is necessarily a kernel.

- (a)  $K(x,z) = K_1(x,z) + K_2(x,z)$  is a kernel. For any finite set  $\{x^{(i)}\}_{i=1}^m$ , the matrix  $K_1$  and  $K_2$  are both symmetric and PSD. Hence, for  $K = K_1 + K_2$  is also symmetric and PSD.
- (b)  $K(x,z) = K_1(x,z) K_2(x,z)$  is not necessarily a kernel. Consider matrix

$$K_1 = \text{diag}(1, 1, \dots, 1), \quad K_2 = \text{diag}(2, 2, \dots, 2).$$

Then

$$K = \operatorname{diag}(-1, -1, \dots, -1)$$

is not PSD.

- (c)  $K(x,z) = aK_1(x,z)$  is a kernel.
- (d)  $K(x,z) = -aK_1(x,z)$  is not a kernel.
- (e)  $K(x,z) = K_1(x,z)K_2(x,z)$  is a kernel. For any finite set  $\{x^{(i)}\}_{i=1}^m$ , the matrix K will be an elementwise product of  $K_1$  and  $K_2$ . Hence K is symmetric. Now we prove K is PSD. We have

$$z^{T}Kz = \sum_{i} \sum_{j} z_{i}K_{ij}z_{j}$$

$$= \sum_{i} \sum_{j} z_{i}K_{1_{ij}}K_{2_{ij}}z_{j}$$

$$= \sum_{i} \sum_{j} z_{i} \sum_{k} \phi_{k}^{1}(x^{(i)})\phi_{k}^{1}(x^{(j)}) \sum_{l} \phi_{l}^{2}(x^{(i)})\phi_{l}^{2}(x^{(j)})z_{j}$$

$$= \sum_{k} \sum_{l} (z_{i}\phi_{k}^{1}(x^{(i)})\phi_{l}^{2}(x^{(i)}))^{2}$$

$$\geq 0,$$

as desired.

(f) K(x,z) = f(x)f(z) is a kernel. For any finite set  $\{x^{(i)}\}_{i=1}^m$ , the matrix K is cleraly symmetric. Now, we have

$$z^{T}Kz = \sum_{i} \sum_{j} z_{i}K_{ij}z_{j}$$

$$= \sum_{i} \sum_{j} z_{i}f(x^{(i)})f(x^{(j)})z_{j}$$

$$= \left(\sum_{i} z_{i}f(x^{(i)})\right)^{2}$$

$$\geq 0.$$

Hence, K is also PSD.

(g)  $K(x,z) = K_3(\phi(x),\phi(z))$  is a kernel. Suppose the feature map corresponds to  $K_3$  is  $\psi$ , then

$$K(x,z) = (\psi \circ \phi(x))^T (\psi \circ \phi(z)).$$

Hence, K is a kernel corresponds to feature map  $\psi \circ \phi$ .

(h)  $K(x,z) = p(K_1(x,z))$  is a kernel. Let  $p(t) = \sum_k a_k t^k$ . Then,  $K(x,z) = \sum_k a_k K_1(x,z)^k$ . By part (e),  $K_1(x,z)^k$  is a kernel for each k. Since  $a_k > 0$ ,  $K(x,z) = \sum_k a_k K_1(x,z)^k$  is a kernel.

## Problem 5 Kernelizing the Perceptron

Binary classification problem with  $y \in \{0, 1\}$ . The perceptron uses hypothese of the form  $h_{\theta}(x) = g(\theta^T x)$ , where  $g(z) = \operatorname{sgn} z = 1$  if  $z \geq 0$ , 0 otherwise. Consider stochastic gradient descent-like implementation of perceptron algorithm, where each update to  $\theta$  is made using one training example. However, unlike stochastic gradient descent, the perceptron algorithm will only make one pass through the entire training set. Update rule is

$$\theta^{(i+1)} := \theta^{(i)} + \alpha (y^{(i+1)} - h_{\theta^i}(x^{(i+1)})) x^{(i+1)},$$

where  $\theta^{(i)}$  is the value of parameters after the algorithm has seen the first *i* training examples. Prior to seeing any training examples,  $\theta^{(0)}$  is initialized to 0.

(a) Let K be a Mercer kenel corresponding to some very high-dimensional feature mapping  $\phi$ . Suppose  $\phi$  is so high-dimensional that it is infeasible to ever represent  $\phi(x)$  explicitly. Descirbe how to apply "kernel trick" to the perceptron to make it work in the high-dimensional feature space  $\phi$ , but without ever explicitly computering  $\phi(x)$ .

No need to worry about the intercept term. If you like, think of  $\phi$  as having the property that  $\phi_0(x) = 1$  so that this is taken care of.

i. To represent the high-dimensional parameter vector  $\theta$  implicitly, we will represent it as a linear combination of  $\phi(x^{(i)})$ , where  $x^{(i)}$  is the training examples. That is,

$$\theta^{(i)} = \sum_{k} \beta_k^{(i)} \phi(x^{(k)}).$$

In particular, the initial value  $\theta^{(0)}$  is represented by setting  $\beta_k = 0$  for all i.

ii. To efficiently make a prediction on a new input  $x^{(i+1)}$ , we compute

$$\begin{split} h_{\theta^{(i)}}(x^{(i+1)}) &= g\left((\theta^{(i)})^T \phi(x^{(i+1)})\right) \\ &= g\left((\sum_k \beta_k^{(i)} \phi(x^{(k)}))^T \phi(x^{(i+1)})\right) \\ &= g\left(\sum_k \beta_k^{(i)} K(x^{(k)}, x^{(i+1)})\right). \end{split}$$

iii. For a new training example  $(x^{(i+1)}, y^{(i+1)})$ , consider the original update rule

$$\theta^{(i+1)} := \theta^{(i)} + \alpha (y^{(i+1)} - h_{\theta^{(i)}}(x^{(i+1)})) x^{(i+1)}.$$

For our representation, we have

$$\sum_{k} \beta_{k}^{(i+1)} \phi(x^{(k)}) := \sum_{k} \beta_{k}^{(i)} \phi(x^{(k)}) + \alpha (y^{(i+1)} - h_{\theta^{(i)}}(x^{(i+1)})) \phi(x^{(i+1)}).$$

Hence, the modified update rule is

$$\beta_{i+1}^{(i+1)} := \beta_{i+1}^{(i)} + \alpha \left( y^{(i+1)} - g \left( \sum_{k} \beta_{k}^{(i)} K(x^{(k)}, x^{(i+1)}) \right) \right)$$

for  $\beta_{i+1}$ , and

$$\beta_k^{(i+1)} := \beta_k^{(i)}$$

for  $\beta_k$  with  $k \neq i+1$ .

- (b) Coding problem.
- (c) The dot kernel performs extremely poorly, since there is no feature mapping and it is equivalent to logistic regression, while the dataset is not linearly separable.

#### Problem 6 Spam classification

Use Bayes algorithm and an SVM to build a spam classifier to detect SMS spam messages.

(a) Coding problem. Implement code for processing the spam messages into numpy arrays.

(b) **Coding problem.** Implement a naive Bayes classifier for spam classification with multinomial event model and Laplace smoothing.

For multinomial event model, the model assume the probability that word k appears in some location is  $\phi_{k|y=1}$  given y=1 and  $\phi_{k|y=0}$  given y=0. That is

$$\phi_{k|y=1} = p(x_j = k \mid y = 1)$$
  $\phi_{k|y=0} = p(x_j = k \mid y = 0),$ 

for k that index into the dictionary size |V| and j that index into the length of the message. Also, the model assume the probability of a spam email  $\phi_y = p(y)$ . Hence, the log-likelihood for Naive Bayes algorithm

$$\ell(\phi_y, \phi_{k|y=1}, \phi_{k|y=0}) = \log \prod_{i=1}^m \left( \prod_{j=1}^{d_i} p(x_j^{(i)} \mid y) \right) p(y^{(i)})$$
$$= \sum_{i=1}^m \sum_{j=1}^{d_i} \left( \log p(x_j^{(i)} \mid y) \right) + \log p(y^{(i)}).$$

and the maximum likelihood estimation with Laplace smoothing

$$\phi_{y} = \frac{\sum_{i=1}^{m} \mathbb{I}\{y^{(i)} = 1\}}{m},$$

$$\phi_{k|y=0} = \frac{\sum_{i=1}^{m} \sum_{j=1}^{d_{i}} \mathbb{I}\{x_{j}^{(i)} = k \wedge y^{(i)} = 1\} + 1}{\sum_{i=1}^{m} \mathbb{I}\{y^{(i)} = 1\}d_{i} + |V|},$$

$$\phi_{k|y=0} = \frac{\sum_{i=1}^{m} \sum_{j=1}^{d_{i}} \mathbb{I}\{x_{j}^{(i)} = k \wedge y^{(i)} = 0\} + 1}{\sum_{i=1}^{m} \mathbb{I}\{y^{(i)} = 0\}d_{i} + |V|}.$$

Notice that  $\sum_{j=1}^{d_i} \mathbb{I}\{x_j^{(i)} = k \wedge y^{(i)} = 1\}$  is just the count of word k in dictionary for message  $x^{(i)}$ , so we only need to count the number of each words in each message.

To predict from the Naive Bayes model, we need to compare

$$p(y = 1 \mid x) = \frac{p(x \mid y = 1)p(y = 1)}{p(x)}, \qquad p(y = 0 \mid x) = \frac{p(x \mid y = 0)p(y = 0)}{p(x)}.$$

When calculating  $p(x \mid y) = \prod_j p(x_j \mid y)$ , underflow may happen since every term is smaller than one. We resolve this issue by using logarithm

$$\log p(x \mid y)p(y) = \log \prod_{j} p(x_j \mid y)p(y)$$
$$= \sum_{j} \log p(x_j \mid y) + \log p(y).$$

and compare

$$\log p(y = 1 \mid x) = \sum_{j} \log p(x_j \mid y = 1) + \log p(y = 1) - \log p(x),$$
$$\log p(y = 0 \mid x) = \sum_{j} \log p(x_j \mid y = 0) + \log p(y = 0) - \log p(x).$$

by taking the difference.

(c) Coding problem. Some tokens may be particularly indicative of an SMS being in a particular class. Try to get an informal sense of how indicative token k is for the SPAM class by looking at

$$\log \frac{p(x_j = k \mid y = 1)}{p(x_j = k \mid y = 0)} = \log \frac{P(\text{token } k \mid \text{email is SPAM})}{P(\text{token } k \mid \text{email is NOTSPAM})}.$$

(d) **Coding problem.** Select the appropriate kernel radius for SVM parameterized by an RBF kernel. That is,

$$K(x, z) = \exp(-\gamma ||x - z||^2).$$