

1. Logistic Regression: Training stability

(a)在数据A上训练logistic regression model很快就收敛了，在数据B上训练logistic regression model无法收敛。

(b)观察后可以发现 θ 的模长越来越大，回顾logistic regression model

$$h_{\theta}(x) = g(\theta^T x), g(z) = 1/(1 + e^{-z}), g(z)' = g(z)(1 - g(z))$$

当 θ 的模长很大时， $\theta^T x$ 的模长很大， $g(\theta^T x) \rightarrow 0$ ， $g(z)' = g(z)(1 - g(z)) \rightarrow 0$ ，从而梯度会越来越小，训练会很慢。

之所以数据B发生这个现象而A没有发生这个现象，是因为数据A线性不可分，数据B线性可分。

由数据B线性可分的

$$y_i \theta^T x_i \geq 0$$

我们的目标函数为

$$J(\theta) = -\frac{1}{m} \sum_{i=1}^m \log(h_{\theta}(y^{(i)} x^{(i)}))$$

要使得使得目标函数越小，只要 $h_{\theta}(y^{(i)} x^{(i)})$ 越大即可，由于 $y_i \theta^T x_i \geq 0$ ，所以 θ 的模长越大， $y_i \theta^T x_i$ 就会越大，由梯度下降的性质可知，迭代之后会令 θ 的模长越来越大，就会发生上述现象。

而数据A不是线性可分的，所以存在 j ，使得

$$y_j \theta^T x_j < 0$$

所以算法不会让 θ 的模长不停地增加。

(c)要解决上述问题，关键是不能让 θ 的模长不停地增长，所以(iii),(v)是最好的方法。

(d)SVM不会发生这个问题，因为SVM是最大间隔分类器，即使可分，最大距离分类器也是唯一的，不会无限迭代下去。

而logistic回归实际上是在让函数间隔变大，所以会出现无法收敛的情形。

2. Model Calibration

(a)只要考虑两个分子即可，logistic回归的输出范围为 $(0, 1)$ ，题目中的 $(a, b) = (0, 1)$ ，所以

$$\begin{aligned} \sum_{i \in I_{a,b}} P(y^{(i)} = 1 | x^{(i)}; \theta) &= \sum_{i=1}^m P(y^{(i)} = 1 | x^{(i)}; \theta) \\ \sum_{i \in I_{a,b}} 1\{y^{(i)} = 1\} &= \sum_{i=1}^m 1\{y^{(i)} = 1\} \end{aligned}$$

接下来证明这两项相等。

回顾损失函数

$$J(\theta) = -\frac{1}{m} \sum_{i=1}^m \left(y^{(i)} \log(h_{\theta}(x^{(i)})) + (1 - y^{(i)}) \log(1 - h_{\theta}(x^{(i)})) \right)$$
$$y^{(i)} \in \{0, 1\}, h_{\theta}(x) = g(\theta^T x), g(z) = 1/(1 + e^{-z})$$

回顾课本的讲义可得

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

那么

$$\nabla_{\theta} J(\theta) = -\frac{1}{m} X^T S$$

其中

$$x_k = [1, x_1^{(k)}, \dots, x_n^{(k)}]^T \in \mathbb{R}^{n+1}$$
$$X = \begin{bmatrix} x_1^T \\ \dots \\ x_m^T \end{bmatrix} = \begin{bmatrix} 1 & x_1^{(1)} & \dots & x_n^{(1)} \\ \dots & \dots & \dots & \dots \\ 1 & x_1^{(m)} & \dots & x_n^{(m)} \end{bmatrix} \in \mathbb{R}^{m \times (n+1)}$$
$$S = \begin{bmatrix} y^{(1)} - h_{\theta}(x^{(1)}) \\ \dots \\ y^{(m)} - h_{\theta}(x^{(m)}) \end{bmatrix} \in \mathbb{R}^m$$

由 θ 的选择规则可知

$$X^T S = 0$$

这里有 $n + 1$ 个等式, 注意 X^T 的第一行全为1, 所以我们考虑第一个等式

$$[1, \dots, 1] S = 0$$
$$\sum_{i=1}^m y^{(i)} - h_{\theta}(x^{(i)}) = 0$$
$$\sum_{i=1}^m y^{(i)} = \sum_{i=1}^m h_{\theta}(x^{(i)})$$

由于 $y^{(i)} \in \{0, 1\}, h_{\theta}(x^{(i)}) = P(y^{(i)} = 1 | x^{(i)}; \theta)$, 所以上式即为

$$\sum_{i=1}^m P(y^{(i)} = 1 | x^{(i)}; \theta) = \sum_{i=1}^m 1\{y^{(i)} = 1\}$$

从而

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

命题得证。

(b)考虑两个数据的数据集 $x^{(1)}, x^{(2)}$, 不妨设 $y^{(1)} = 1, y^{(2)} = 0$, 如果

$$P(y^{(1)} = 1 | x^{(1)}; \theta) = 0.4, P(y^{(2)} = 1 | x^{(2)}; \theta) = 0.6$$

那么我们预测 $y^{(1)} = 0, y^{(2)} = 1$, 准确率为0, 但是

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

所以perfectly calibrated无法推出perfect accuracy。

反之, 如果

$$P(y^{(1)} = 1 | x^{(1)}; \theta) = 0.6, P(y^{(2)} = 1 | x^{(2)}; \theta) = 0.3$$

那么我们预测 $y^{(1)} = 1, y^{(2)} = 0$, 此时准确率为1, 但是

$$\sum_{i \in I_{a,b}} P(y^{(i)} = 1 | x^{(i)}; \theta) = 0.9 \neq \sum_{i \in I_{a,b}} 1\{y^{(i)} = 1\} = 1$$

所以perfect accuracy无法推出perfectly calibrated。

(c)设损失函数为

$$J(\theta) = -\frac{1}{m} \sum_{i=1}^m \left(y^{(i)} \log(h_{\theta}(x^{(i)})) + (1 - y^{(i)}) \log(1 - h_{\theta}(x^{(i)})) \right) + C \sum_{i=1}^{n+1} \theta_i^2$$

记

$$\theta = \begin{bmatrix} \theta_1 \\ \dots \\ \theta_{n+1} \end{bmatrix}$$

继续使用(a)的记号, 那么

$$\begin{aligned} \nabla J(\theta) &= -\frac{1}{m} X^T S + 2C\theta = 0 \\ X^T S &= 2mC\theta \end{aligned}$$

依旧考虑第一个等式

$$\begin{aligned}
[1, \dots, 1]S &= 2mC\theta_1 \\
\sum_{i=1}^m y^{(i)} - h_{\theta}(x^{(i)}) &= 2mC\theta_1 \\
\sum_{i=1}^m y^{(i)} &= \sum_{i=1}^m h_{\theta}(x^{(i)}) + 2mC\theta_1 \\
\sum_{i=1}^m 1\{y^{(i)} = 1\} &= \sum_{i=1}^m P(y^{(i)} = 1|x^{(i)}; \theta) + 2mC\theta_1
\end{aligned}$$

从而

$$\sum_{i=1}^m 1\{y^{(i)} = 1\} = \sum_{i=1}^m P(y^{(i)} = 1|x^{(i)}; \theta) + 2mC\theta_1$$

3. Bayesian Logistic Regression and weight decay

回顾定义

$$\begin{aligned}
\theta_{\text{MAP}} &= \arg \max_{\theta} p(\theta) \prod_{i=1}^m p(y^{(i)} | x^{(i)}, \theta) \\
&= \arg \max_{\theta} \exp\left(-\frac{||\theta||^2}{2\tau^2}\right) \prod_{i=1}^m p(y^{(i)} | x^{(i)}, \theta)
\end{aligned}$$

由定义可知

$$\exp\left(-\frac{||\theta_{\text{MAP}}||^2}{2\tau^2}\right) \prod_{i=1}^m p(y^{(i)} | x^{(i)}, \theta_{\text{MAP}}) \geq \exp\left(-\frac{||\theta_{\text{ML}}||^2}{2\tau^2}\right) \prod_{i=1}^m p(y^{(i)} | x^{(i)}, \theta_{\text{ML}})$$

因为

$$\prod_{i=1}^m p(y^{(i)} | x^{(i)}, \theta_{\text{MAP}}) \leq \prod_{i=1}^m p(y^{(i)} | x^{(i)}, \theta_{\text{ML}})$$

所以

$$\exp\left(-\frac{||\theta_{\text{MAP}}||^2}{2\tau^2}\right) \geq \exp\left(-\frac{||\theta_{\text{ML}}||^2}{2\tau^2}\right)$$

从而

$$||\theta_{\text{MAP}}||_2 \leq ||\theta_{\text{ML}}||_2$$

4. Constructing kernels

假设 K_i 对应的矩阵为 M_i , K 对应矩阵为 M , 由核函数的定义可知 M_i 为半正定阵。

(a) $K(x, z) = K_1(x, z) + K_2(x, z)$ 是核, 因为

$$\begin{aligned}x^T M x &= x^T (M_1 + M_2) x \\&= x^T M_1 x + x^T M_2 x \\&\geq 0\end{aligned}$$

(b) $K(x, z) = K_1(x, z) - K_2(x, z)$ 不是核。取 $K_2(x, z) = 2K_1(x, z)$

$$x^T M x = x^T (M_1 - M_2) x = -x^T M_1 x \leq 0$$

(c) $K(x, z) = aK_1(x, z), a > 0$ 是核

$$\begin{aligned}x^T M x &= x^T (aM_1) x \\&= ax^T M_1 x \\&\geq 0\end{aligned}$$

(d) $K(x, z) = -aK_1(x, z), a > 0$ 不是核

$$\begin{aligned}x^T M x &= x^T (-aM_1) x \\&= -ax^T M_1 x \\&\leq 0\end{aligned}$$

(e) $K(x, z) = K_1(x, z)K_2(x, z)$ 是核

因为 K_1, K_2 为核, 所以设 $K_1(x, z) = \Phi_1(x)\Phi_1^T(z), K_2 = \Phi_2(x)\Phi_2^T(z)$ 。

记 $\Phi(x)$ 是 $\Phi_1(x)\Phi_2^T(x)$ 每一行拼接而成的向量, 设 $\Phi_1(x), \Phi_2(x) \in \mathbb{R}^n$, 给出以下记号

$$\begin{aligned}\Phi^i(x) &= \Phi_1^i(x)\Phi_2^T(x) \in \mathbb{R}^{1 \times n} \\ \Phi_1^i(x) &\text{为 } \Phi_1(x) \text{ 的第 } i \text{ 个分量}\end{aligned}$$

那么

$$\Phi(x) = [\Phi^1(x) \quad \Phi^2(x) \quad \dots \quad \Phi^n(x)] \in \mathbb{R}^{1 \times n^2}$$

接着计算 $\Phi(x)\Phi^T(x')$, 注意 $\Phi^i(x)$ 为行向量

$$\begin{aligned}
(\Phi(x)\Phi^T(x')) &= \sum_{i=1}^n (\Phi^i(x))\Phi^i(x')^T \\
&= \sum_{i=1}^n (\Phi_1^i(x)\Phi_2(x)^T)(\Phi_1^i(x')\Phi_2(x')^T)^T \\
&= \sum_{i=1}^n \Phi_1^i(x)\Phi_1^i(x')\Phi_2(x)^T\Phi_2(x') \\
&= \sum_{i=1}^n \Phi_1^i(x)\Phi_1^i(x')K_2(x, x') \\
&= K_2(x, x') \sum_{i=1}^n \Phi_1^i(x)\Phi_1^i(x') \\
&= K_2(x, x')K_1(x, x')
\end{aligned}$$

所以 $\Phi(x)$ 对应的核为 $K_1(x, x')K_2(x, x')$, 从而 $K(x, z) = K_1(x, z)K_2(x, z)$ 是核。

(f) $K(x, z) = f(x)f(z)$ 是核, 因为符合定义。

(g) $K(x, z) = K_3(\phi(x), \phi(z))$ 是核, 因为

$$y^T M y = y^T M_3 y \geq 0$$

(h)由(e)可知, 如果 K_1 是核, 那么 $K_1^i (i \geq 1, i \in N)$ 也是核, 又由(a)(c)可得核函数的正系数的线性组合为核, 所以 $K(x, z) = p(K_1(x, z))$ 也是核。

5. Kernelizing the Perceptron

设这里的数据为 $x^{(1)}, \dots, x^{(m)}$

(a)根据更新公式

$$\theta^{(i+1)} := \theta^{(i)} + \alpha 1\{g(\theta^{(i)T} \phi(x^{(i+1)}))y^{(i+1)} < 0\}y^{(i+1)} \phi(x^{(i+1)})$$

如果初始化 $\theta^{(0)} = 0$, 那么 $\theta^{(i)}$ 可以表示为 $\phi(x^{(j)})$ 的线性组合, 从而

$$\theta^{(i)} = \sum_{j=1}^i \beta_j \phi(x^{(j)})$$

(b)计算 $g(\theta^{(i)T} \phi(x^{(i+1)}))$

$$g(\theta^{(i)T} \phi(x^{(i+1)})) = g\left(\sum_{j=1}^i \beta_j \phi(x^{(j)})^T \phi(x^{(i+1)})\right) = g\left(\sum_{j=1}^i \beta_j M_{j, i+1}\right)$$

(c)由上述公式可知, 第 i 次我们只要更新 β_i 即可, 更新公式如下

$$\beta_{i+1} = \alpha 1\{g(\theta^{(i)T} \phi(x^{(i+1)}))y^{(i+1)} < 0\}y^{(i+1)}$$

6.Spam classification

(a)(b)(c)

代码的注释比较详细，这里只说明一点，在朴素贝叶斯中我们需要计算：

$$\prod_{i=1}^m \left(\prod_{j=1}^{n_i} p(x_j^{(i)} | y^{(i)}; \phi_{k|y=0}, \phi_{k|y=1}) p(y^{(i)}; \phi_y) \right)$$

题目中计算的对数概率，这是为了防止数值过小变为0

$$\sum_{i=1}^n \left(\sum_{j=1}^{n_i} \log p(x_j^{(i)} | y^{(i)}; \phi_{k|y=0}, \phi_{k|y=1}) + \log p(y^{(i)}; \phi_y) \right)$$

```
# -*- coding: utf-8 -*-
"""
Created on Thu Mar 14 15:58:26 2019

@author: qinzhen
"""

import numpy as np
import matplotlib.pyplot as plt

def readMatrix(file):
    fd = open(file, 'r')
    #读取第一行
    hdr = fd.readline()
    #读取行列
    rows, cols = [int(s) for s in fd.readline().strip().split()]
    #读取单词
    tokens = fd.readline().strip().split()
    #构造空矩阵
    matrix = np.zeros((rows, cols))
    Y = []
    #line为每行的元素
    for i, line in enumerate(fd):
        nums = [int(x) for x in line.strip().split()]
        #第一个元素表示是否为垃圾邮件
        Y.append(nums[0])
        #将后续数据读入
        kv = np.array(nums[1:])
        #从第一个开始每两个累加
        k = np.cumsum(kv[:-1:2])
        #从第二个开始每隔一个取出
        v = kv[1::2]
        #这里应该是一种特殊的存储格式，我们直接使用即可
        matrix[i, k] = v
    return matrix, tokens, np.array(Y)

def nb_train(matrix, category):
    #matrix(i, j)表示第i个邮件中第j个元素出现了几次
```

```

state = {}
#token的数量
N = matrix.shape[1]
#邮件数量
M = matrix.shape[0]
#####

#垃圾邮件的数量
y1 = matrix[category==1]
n1 = np.sum(y1)
#非垃圾邮件的数量
y0 = matrix[category==0]
n0 = np.sum(y0)

#P(y=1)
p1 = y1.shape[0] / M
#P(y=0)
p0 = y0.shape[0] / M
state[-1] = [p0, p1]

for i in range(N):
    #找到第i个token
    #垃圾邮件中第i个token出现的数量
    s1 = matrix[category==1][:, i]
    #拉普拉斯平滑
    u1 = (s1.sum() + 1) / (n1 + N)
    #非垃圾邮件中第i个token出现的数量
    s0 = matrix[category==0][:, i]
    #拉普拉斯平滑
    u0 = (s0.sum() + 1) / (n0 + N)
    #存入字典
    state[i] = [u0, u1]

#####
return state

def nb_test(matrix, state):
    output = np.zeros(matrix.shape[0])
    #####
    #邮件数量
    M = matrix.shape[0]
    #token的数量
    N = matrix.shape[1]
    for i in range(M):
        #第i个邮件
        vector = matrix[i]
        s1 = np.log(state[-1][1])
        s0 = np.log(state[-1][0])

        for j in range(N):
            #对第j个token的对数概率做累加
            s1 += vector[j] * np.log(state[j][1])
            s0 += vector[j] * np.log(state[j][0])

```



```

        if s1 > s0:
            output[i] = 1

#####
return output

def evaluate(output, label):
    error = (output != label).sum() * 1. / len(output)
    print('Error: %1.4f' % error)
    return error

def nb(file):
    trainMatrix, tokenlist, trainCategory = readMatrix(file)
    testMatrix, tokenlist, testCategory = readMatrix('MATRIX.TEST')

    state = nb_train(trainMatrix, trainCategory)
    output = nb_test(testMatrix, state)

    return evaluate(output, testCategory)

trainMatrix, tokenlist, trainCategory = readMatrix('MATRIX.TRAIN')
testMatrix, tokenlist, testCategory = readMatrix('MATRIX.TEST')

state = nb_train(trainMatrix, trainCategory)
output = nb_test(testMatrix, state)

evaluate(output, testCategory)

#problem b
b=[]
for i in range(1448):
    b.append((i,np.log(state[i][1])-np.log(state[i][0])))

b.sort(key=lambda i:i[-1],reverse=True)
key = b[:5]

word = []
for i in key:
    word.append(tokenlist[i[0]])

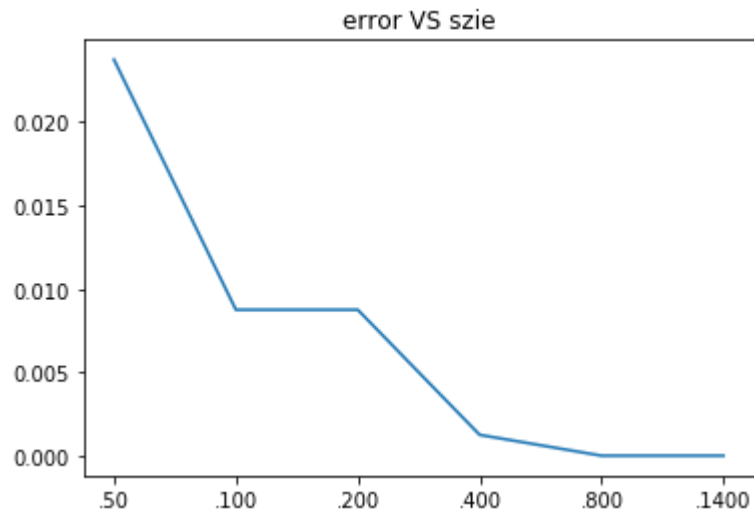
print(word)

#problem c
size = ['.50', '.100', '.200', '.400', '.800', '.1400']
size1 = [50, 100, 200, 400, 800, 1400]
train = "MATRIX.TRAIN"
error = []
for i in size:
    file = train+i
    error.append(nb(file))

plt.plot(size, error)
plt.title("error VS szie")

```

```
Error: 0.0025
Error: 0.0238
Error: 0.0088
Error: 0.0088
Error: 0.0013
Error: 0.0000
Error: 0.0000
```



(d)这部分老师已经提供，但这里还是详细解释下。

首先这题需要计算高斯核矩阵，所以我们需要计算 $[\|x^{(i)} - x^{(j)}\|^2]_{i,j}$ ，下面介绍向量化计算的方法：
假设

$$X = \begin{bmatrix} -(x^{(1)})^T \\ -(x^{(2)})^T \\ \vdots \\ -(x^{(m)})^T \end{bmatrix} \in \mathbb{R}^{m \times d}, Y = \begin{bmatrix} -(y^{(1)})^T \\ -(y^{(2)})^T \\ \vdots \\ -(y^{(n)})^T \end{bmatrix} \in \mathbb{R}^{n \times d}$$

其中 $x^{(i)}, y^{(i)} \in \mathbb{R}^d$ ，现在的问题是如何高效计算矩阵 $D \in \mathbb{R}^{m \times n}$ ，其中

$$D_{i,j} = \|x^{(i)} - y^{(j)}\|^2$$

首先对 $D_{i,j}$ 进行处理

$$\begin{aligned} D_{i,j} &= \|x^{(i)} - y^{(j)}\|^2 \\ &= (x^{(i)} - y^{(j)})^T (x^{(i)} - y^{(j)}) \\ &= (x^{(i)})^T x^{(i)} - 2(x^{(i)})^T y^{(j)} + (y^{(j)})^T y^{(j)} \end{aligned}$$

那么

$$\begin{aligned}
D &= \begin{bmatrix} D_{1,1} & \dots & D_{1,n} \\ \dots & \dots & \dots \\ D_{m,1} & \dots & D_{m,n} \end{bmatrix} \\
&= \begin{bmatrix} (x^{(1)})^T x^{(1)} - 2(x^{(1)})^T y^{(1)} + (y^{(1)})^T y^{(1)} & \dots & (x^{(1)})^T x^{(n)} - 2(x^{(1)})^T y^{(n)} + (y^{(n)})^T y^{(n)} \\ \dots & \dots & \dots \\ (x^{(m)})^T x^{(m)} - 2(x^{(m)})^T y^{(1)} + (y^{(1)})^T y^{(1)} & \dots & (x^{(m)})^T x^{(n)} - 2(x^{(m)})^T y^{(n)} + (y^{(n)})^T y^{(n)} \end{bmatrix} \\
&= \begin{bmatrix} (x^{(1)})^T x^{(1)} & \dots & (x^{(1)})^T x^{(n)} \\ \dots & \dots & \dots \\ (x^{(m)})^T x^{(m)} & \dots & (x^{(m)})^T x^{(n)} \end{bmatrix} + \begin{bmatrix} (y^{(1)})^T y^{(1)} & \dots & (y^{(n)})^T y^{(n)} \\ \dots & \dots & \dots \\ (y^{(1)})^T y^{(1)} & \dots & (y^{(n)})^T y^{(n)} \end{bmatrix} - 2 \begin{bmatrix} (x^{(1)})^T y^{(1)} & \dots & (x^{(1)})^T y^{(n)} \\ \dots & \dots & \dots \\ (x^{(m)})^T y^{(1)} & \dots & (x^{(m)})^T y^{(n)} \end{bmatrix} \\
&= \begin{bmatrix} (x^{(1)})^T x^{(1)} \\ \dots \\ (x^{(m)})^T x^{(m)} \end{bmatrix} \underbrace{\begin{bmatrix} 1 & \dots & 1 \end{bmatrix}}_{1 \times n \text{ 矩阵}} + \underbrace{\begin{bmatrix} 1 \\ \vdots \\ 1 \end{bmatrix}}_{m \times 1 \text{ 矩阵}} \begin{bmatrix} (y^{(1)})^T y^{(1)} & \dots & (y^{(n)})^T y^{(n)} \end{bmatrix} - 2XY^T
\end{aligned}$$

特别的，这里 $X = Y$ ，所以上述代码如下：

```
d1 = np.sum(matrix ** 2, axis=1).reshape(-1, 1)
d2 = d1.T
dist = d1 + d2 - 2 * matrix.dot(matrix.T)
```

带入高斯核的计算公式可得：

```
k = np.exp(- dist / (2 * (tau ** 2)))
```

老师给出的代码为：

```
gram = matrix.dot(matrix.T)
squared = np.sum(matrix*matrix, axis=1)
k = np.exp(-(squared.reshape((-1,1)) + squared.reshape((1,-1)) - 2 * gram) / (2 * (tau ** 2)))
```

为了解释剩余代码，首先介绍SVM对应的合页损失函数（参考统计学习方法113页）：

$$\sum_{i=1}^N [1 - y^{(i)}(w^T x^{(i)} + b)]_+ + \lambda ||w||^2$$

其中

$$[x]_+ = \begin{cases} 1 & x > 0 \\ 0 & x \leq 0 \end{cases}$$

假设SVM对应的特征转换为 $\phi(x)$ ，那么由课上的讨论可知， w 的形式如下：

$$w = \sum_{j=1}^N \alpha_j \phi(x^{(j)}) \quad (1)$$

与之对应的损失函数为

$$\begin{aligned} \sum_{i=1}^N [1 - y^{(i)}(w^T \phi(x^{(i)}) + b)]_+ + \lambda \|w\|^2 &= \sum_{i=1}^N \left[1 - y^{(i)} \left(\left(\sum_{j=1}^N \alpha_j \phi(x^{(j)}) \right)^T \phi(x^{(i)}) + b \right) \right]_+ \\ &\quad + \lambda \left(\sum_{j=1}^N \alpha_j \phi(x^{(j)}) \right)^T \left(\sum_{j=1}^N \alpha_j \phi(x^{(j)}) \right) \\ &= \sum_{i=1}^N \left[1 - y^{(i)} \left(\sum_{j=1}^N \alpha_j \phi^T(x^{(j)}) \phi(x^{(i)}) + b \right) \right]_+ \\ &\quad + \lambda \sum_{i=1}^N \sum_{j=1}^N \alpha_i \alpha_j \phi^T(x^{(i)}) \phi(x^{(j)}) \end{aligned}$$

由(1)可知，我们只要求出 $\alpha = (\alpha_1, \dots, \alpha_N)^T$ 即可，所以可以关于 α 做梯度随机梯度下降法，注意一个样本（不妨设为 $x^{(i)}$ ）对应的损失函数为

$$L_i \triangleq \left[1 - y^{(i)} \left(\sum_{j=1}^N \alpha_j \phi^T(x^{(j)}) \phi(x^{(i)}) + b \right) \right]_+ + \frac{\lambda}{N} \sum_{s=1}^N \sum_{t=1}^N \alpha_s \alpha_t \phi^T(x^{(s)}) \phi(x^{(t)})$$

记核矩阵为 K ，对上式关于 α_k 求偏导可得

$$\begin{aligned} \frac{\partial L}{\partial \alpha_k} &= 1\{[1 - y^{(i)}(\sum_{j=1}^N \alpha_j \phi^T(x^{(j)}) \phi(x^{(i)}) + b)] > 0\} \times \left(-y^{(i)} \phi^T(x^{(k)}) \phi(x^{(i)}) \right) + 2 \frac{\lambda}{N} \left(\sum_{j=1}^N \alpha_j \phi^T(x^{(j)}) \phi(x^{(k)}) \right) \\ &= 1\{[1 - y^{(i)}(w^T \phi(x^{(i)}) + b)] > 0\} \times \left(-y^{(i)} \phi^T(x^{(k)}) \phi(x^{(i)}) \right) + 2 \frac{\lambda}{N} \left(\phi(x^{(i)}) w^T \right) \\ &= 1\{[1 - y^{(i)}(w^T \phi(x^{(i)}) + b)] > 0\} \times \left(-y^{(i)} K_{k,i} \right) + 2 \frac{\lambda}{N} \left(\phi(x^{(i)}) w^T \right) \end{aligned}$$

因此梯度为

$$\nabla_{\alpha} L = 1\{[1 - y^{(i)}(w^T \phi(x^{(i)}) + b)] > 0\} (-y^{(i)}) \begin{bmatrix} K_{1,i} \\ \vdots \\ K_{N,i} \end{bmatrix} + \frac{2\lambda}{N} K \alpha$$

所以计算梯度的过程如下，首先随机选个更新的样本（M为样本的数量）：

```
i = int(np.random.rand() * M)
```

接着计算函数间隔（k为之前计算的核矩阵）：

```
margin = category[i] * (k[i, :].dot(alpha))
```

然后计算正则项的梯度，注意这里我和老师给的计算式不大一样，老师给的式子如下：

```
grad = M * L * k[:, i] * alpha[i]
```

我的计算式如下：

```
grad = L * k.dot(alpha)
```

这里系数没有那么重要，实验结果表明两种方法的效果都尚可。

最后根据函数间隔是否大于1决定是否更新前一项的梯度：

```
if(margin < 1):  
    grad -= category[i] * k[:, i]
```

代码中的梯度除以了迭代次数j的一个函数，这是为了让梯度逐渐变小：

```
alpha -= grad / ((np.sqrt(j+1)))
```

完整代码如下：

```
# -*- coding: utf-8 -*-  
"""  
Created on Thu Mar 14 16:04:08 2019  
  
@author: qinzhen  
"""  
  
import numpy as np  
import matplotlib.pyplot as plt  
  
tau = 8.  
  
def readMatrix(file):  
    fd = open(file, 'r')  
    hdr = fd.readline()  
    rows, cols = [int(s) for s in fd.readline().strip().split()]  
    tokens = fd.readline().strip().split()  
    matrix = np.zeros((rows, cols))  
    Y = []  
    for i, line in enumerate(fd):  
        nums = [int(x) for x in line.strip().split()]  
        Y.append(nums[0])  
        kv = np.array(nums[1:])  
        k = np.cumsum(kv[:-1:2])  
        v = kv[1::2]  
        matrix[i, k] = v  
    #化为正负1  
    category = (np.array(Y) * 2) - 1  
    return matrix, tokens, category  
  
def svm_train(matrix, category):  
    state = {}  
    M, N = matrix.shape  
    #####
```

```

#大于0的化为1
matrix = 1.0 * (matrix > 0)
'''

#构造kernel矩阵
d1 = np.sum(matrix ** 2, axis=1).reshape(-1, 1)
d2 = d1.T
squared = matrix.dot(matrix.T)
dist = d1 + d2 - 2 * squared
k = np.exp(- dist / (2 * (tau ** 2)))
'''

gram = matrix.dot(matrix.T)
squared = np.sum(matrix*matrix, axis=1)
k = np.exp(-(squared.reshape((-1,1)) + squared.reshape((1,-1)) - 2 * gram) / (2 * (tau
** 2)))

#初始化向量
alpha = np.zeros(M)
#循环次数
n = 40
#系数
L = 1. / (64 * M)
#平均值
alpha_avg = np.zeros(M)

for j in range(n * M):
    #随机取一个样本
    i = int(np.random.rand() * M)
    #计算函数间隔
    margin = category[i] * (k[i, :].dot(alpha))
    #grad = M * L * k[:, i] * alpha[i]
    grad = L * k.dot(alpha)
    if(margin < 1):
        grad -= category[i] * k[:, i]
    alpha -= grad / ((np.sqrt(j+1)))
    alpha_avg += alpha

alpha_avg /= (n * M)

state['alpha'] = alpha
state['alpha_avg'] = alpha_avg
state['Xtrain'] = matrix
state['Sqtrain'] = squared

#####
return state

def svm_test(matrix, state):
    M, N = matrix.shape
    output = np.zeros(M)
    #####
    Xtrain = state['Xtrain']
    Sqtrain = state['Sqtrain']

```

```

#大于0的化为1
matrix = 1.0 * (matrix > 0)
#做测试集的kernel
gram = matrix.dot(Xtrain.T)
squared = np.sum(matrix * matrix, axis=1)
k = np.exp(-(squared.reshape((-1, 1)) + Sqtrain.reshape((1, -1)) - 2 * gram) / (2 *
(tau**2)))
#读取alpha
alpha_avg = state['alpha_avg']
#预测
pred = k.dot(alpha_avg)
output = np.sign(pred)

#####
return output

def evaluate(output, label):
    error = (output != label).sum() * 1. / len(output)
    print('Error: %1.4f' % error)
    return error

def svm(file):
    trainMatrix, tokenlist, trainCategory = readMatrix(file)
    testMatrix, tokenlist, testCategory = readMatrix('MATRIX.TEST')

    state = svm_train(trainMatrix, trainCategory)
    output = svm_test(testMatrix, state)

    return evaluate(output, testCategory)

trainMatrix, tokenlist, trainCategory = readMatrix('MATRIX.TRAIN.400')
testMatrix, tokenlist, testCategory = readMatrix('MATRIX.TEST')

state = svm_train(trainMatrix, trainCategory)
output = svm_test(testMatrix, state)

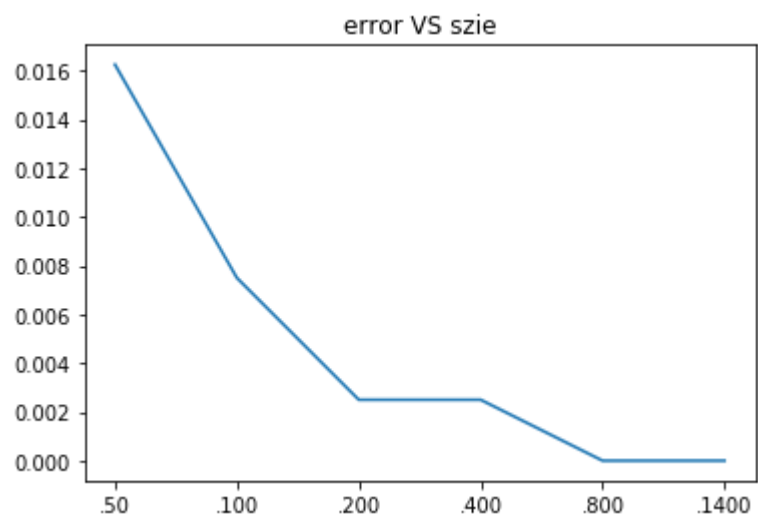
evaluate(output, testCategory)

size = ['.50', '.100', '.200', '.400', '.800', '.1400']
size1 = [50, 100, 200, 400, 800, 1400]
train = "MATRIX.TRAIN"
error = []
for i in size:
    file = train+i
    error.append(svm(file))

plt.plot(size, error)
plt.title("error VS szie")

```

Error: 0.0025
Error: 0.0163
Error: 0.0075
Error: 0.0025
Error: 0.0025
Error: 0.0000
Error: 0.0000



(e)对比两图可以发现，和NB算法相比，SVM算法的效果更好。