ECE 532 Homework 8: The Hinge Loss & the Support Vector Machines

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QUESTION1

Express the standard least squares solution using the dual form using regularization parameter $\lambda=10^{-5}$. Verify that it generates the same results as the primal solution.

This is the standard least square optimization with the ridge regularizer:

$$\min_{w \in \mathbb{R}^n} = \|y - Xw\|_2^2 + \lambda \|w\|_2^2$$

by setting $w = X^T \alpha$, for some α , we obtained the dual optimization :

$$\min_{\alpha \in \mathbb{D}^m} = \|y - XX^T \alpha\|_2^2 + \lambda \alpha^T XX^T \alpha$$

Let $\hat{\alpha}$ denote the solution of this dual optimization, then

$$\hat{\alpha} = (XX^T + \lambda I)^{-1} v$$

The following code can be used to test that they provides the same weights.

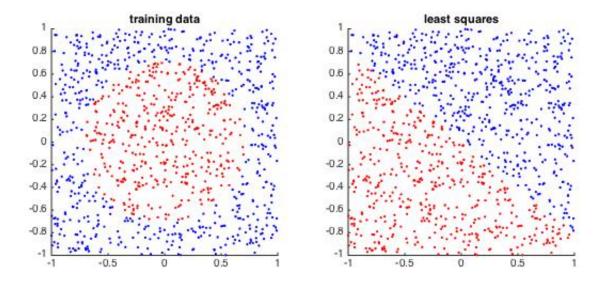
```
clear; close all;clc
training_data
% set lambda
lambda = 1e-5;
% fit ridge regression vs. the dual optimization
x.ridge = inv(A' * A + lambda * eye(size(A' * A))) * A' * b;
alpha = inv(A * A' + lambda * eye(size(A * A'))) * b;
% recover the ridge weights from the dual
x.ridgeDual = A' * alpha;
[x.ridge x.ridgeDual]
```

Design a classifier using the Gaussian kernel

$$k(a_i, a_j) = exp(-\frac{1}{2}||a_i - a_j||_2^2)$$

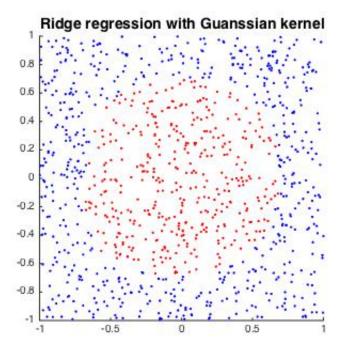
and regularization parameter $\lambda=10^{-5}$. Compare its classification performance of the training data to the least square method.

Here's a visualization of the data



On the left, the plot is showing the truth. The positive and negative cases were plotted using different colors. From this plot, we know that we need a non-linear decision boundary since there is no straight line solution is going to separate the two classes.

Here's the plot showing the predicted classification for ridgre regression with Gaussian kernel:



The accuracy of ridgre regression with Gaussian kernel is 0.9820, which is much better than the least square solution.

Here's the MATLAB code for Gaussian kernelized ridge

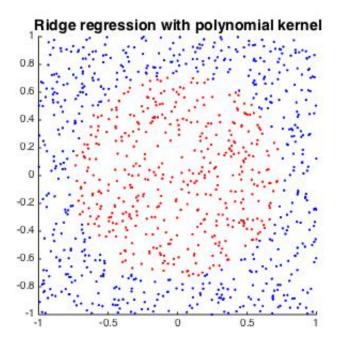
```
clear; close all;clc
training_data
% set lambda
lambda = 1e-5;
%% compute the polynomial kernal
m = size(A, 1);
gauKer = nan(m,m);
for i = 1 : m
    for j = 1 : m
        gauKer(i,j) = exp(-.5 * (norm(A(i,:) - A(j,:),2))^2);
end
%% fit the kernelized least square
alpha = inv(gauKer + lambda * eye(size(gauKer))) * b;
%% compute classification accuracy
prediction = nan(m, 1);
for i = 1 : m
    temp = 0;
    xnew = A(i,:);
    for j = 1 : m
        temp = temp + alpha(j) * exp(-.5 * (norm(xnew - A(j,:),2))^2);
    prediction(i) = sign(temp);
end
sum(prediction)
accuracy = sum(prediction == b) / length(b)
%% plot the prediction
figure;
hold on;
for i=1:m
    a = A(i,:);
    if prediction(i) == 1
        plot(a(1),a(2),'b.');
        plot(a(1),a(2),'r.');
    end
end
axis('square')
title('Ridge regression with Guanssian kernel', 'fontsize',14)
```

Design a classifier using the polynomial kernel

$$k(a_i, a_j) = (a_i^T a_j + 1)^2$$

and regularization parameter $\lambda=10^{-5}$. Compare its classification performance of the training data to the least square and the Gaussian kernel classifier.

Here's the plot showing the predicted classification:



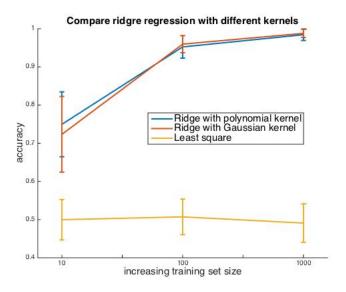
The accuracy of ridgre regression with polynomial kernel is 0.9650, which is much better than the least square solution. It is slightly worse than the Gaussian kernel but I am not sure if the statistically difference is significant or not.

Here's the MATLAB code for polynomial kernelized ridge

```
clear; close all; clc
training_data
% set lambda
lambda = 1e-5;
%% compute the polynomial kernal
m = size(A, 1);
ker = nan(m,m);
for i=1:m
    for j=1:m
       ker(i,j) = (dot(A(i,:), A(j,:)) +1)^2;
end
% %% test kernel
% kert = phi * phi';
% [ker(:,1) kert(:,1)]
%% fit the kernelized least square
alpha = inv(ker + lambda * eye(size(ker))) * b;
%% compute classification accuracy
prediction = nan(m, 1);
predict_t = nan(m,1);
% compute the kernel LS prediction 1 by 1
for i = 1 : m
   temp = 0;
   xnew = A(i,:);
    % compute the kernel LS prediction for 1 example
    for j = 1 : m
       temp = temp + alpha(j) * (dot(xnew, A(j,:)) +1 )^2;
   prediction(i) = sign(temp);
end
accuracy = sum(prediction == b) / length(b)
%% plot the prediction
figure;
hold on;
for i = 1 : m
   a = A(i, :);
   if prediction(i) == 1
       plot(a(1),a(2),'b.');
       plot(a(1),a(2),'r.');
   end
end
axis('square')
title('Ridge regression with polynomial kernel', 'fontsize', 14)
```

Now experiment with these methods in the following way. Generate different sized sets of training data: m = 10, 100, 1000. Design the three types of classifiers using these data. Test the performance of the classifiers using independent sets of data of size 100. For each training data size, repeat the training and testing 100 times and average the results. This will provide a good indication of how the different approaches perform with different amounts of training data. Summarize your results by reporting the average test error for each of the three methods and each of the three training set sizes.

Here's the plot that shows the comparison of the performance.



The Y axis shows the accuracy, and the X axis is showing increasing size of the training set, ranges from 10 to 1000. I am comparing standard least square and ridge regression with polynomial kernel or Gaussian kernel. They were visualized by different colored lines. The error bar is showing one standard deviation.

We can see that when there are more training examples, ridge regression with either Gaussian or polynomial kernel performed better. In particular, the performance of both kernelized ridge reached the ceiling in this setting when the training size is 1000. Compare to the polynomial kernel, it seems that Gassian kernel is better when there were 1000 training examples, and it is worse when there were 10 training examples but the statistical difference was insignificant.

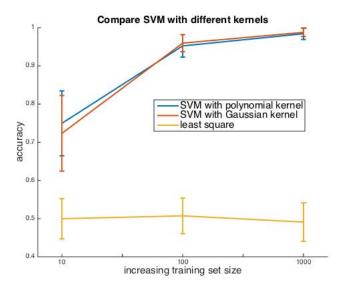
On the other hand, the performance of standard least square did not increase when there are more training examples.

Here's the MATLAB code I used for comparing the 3 method on one train-test session.

```
%% compare 2 kernel methods
function [accuracy] = compare3methods(trainingSetSize)
% generate training and testing data
[A, b] = genTraining_data(trainingSetSize, 0);
[Atest, btest] = genTraining_data(100, 0);
% set lambda
lambda = 1e-5;
%% compute the polynomial kernal
m = size(A, 1);
ker.poly = nan(m,m);
ker.gaus = nan(m,m);
for i=1:m
    for j=1:m
        ker.poly(i,j) = (dot(A(i,:), A(j,:)) +1)^2;
        ker.gaus(i,j) = exp(-.5 * (norm(A(i,:) - A(j,:),2))^2);
    end
end
%% fit the kernelized least square
alpha.poly = inv(ker.poly + lambda * eye(size(ker.poly))) * b;
alpha.gaus = inv(ker.gaus + lambda * eye(size(ker.gaus))) * b;
%% compute classification accuracy
prediction.poly = nan(size(Atest, 1), 1);
prediction.gaus = nan(size(Atest, 1), 1);
% compute the kernel LS prediction 1 by 1
for i = 1 : size(Atest, 1)
   xnew = Atest(i,:);
    % compute kernel LS prediction for 1 example (polynomial)
    temp = 0;
    for j = 1 : m
        temp = temp + alpha.poly(j) * (dot(xnew, A(j,:)) +1 )^2;
    prediction.poly(i) = sign(temp);
    % compute kernel LS prediction for 1 example (gaussian)
    temp = 0;
    for j = 1 : m
        temp = temp + alpha.gaus(j) * exp(-.5 * (norm(xnew - A(j,:),2))^2);
   prediction.gaus(i) = sign(temp);
end
%% fit standard LS
wts.ls = pinv(A)*b;
prediction.ls = sign(Atest * wts.ls);
%% compute accuracy for the 2 methods
accuracy.p = sum(prediction.poly == btest) / length(btest);
accuracy.g = sum(prediction.gaus == btest) / length(btest);
accuracy.ls = sum(prediction.ls == btest) / length(btest);
end
```

Now tackle the problem using hinge loss instead of squared error loss. You do this using the Matlab function symtrain, another package, or by writing your own code (e.g., GD or SGD). Generate different sized sets of training data: m=10,100,1000. Design SVM classifiers using both Gaussian and polynomial kernels. Test the performance of the classifiers using independent sets of data of size 100. For each training data size, repeat the training and testing 100 times and average the results. Compare your results to those obtained using least squares.

Here's the plot showing the three methods I tried:



The Y axis shows the accuracy, and the X axis is showing increasing size of the training set, ranges from 10 to 1000. I am comparing the least square solution with support vector machine with polynomial kernel and Gaussian kernel. They were visualized by different colored lines. The error bar is showing one standard deviation.

The story is pretty similar with the what we saw in question 4. SVM with both kernels performed better when there are more training examples. The performance of both kernelized SVM reached the ceiling when the training size is 1000. On the other hand, the performance of standard least square did not increase when there are more training examples. Indeed, standard least square always give us chance performance.

Here's the MATLAB code for comparing the 3 methods, used in question 5. I used matlab symtrain package.

```
function accuracy = compare3kernels_svm(m)
% generate training and test data
% m = 100;
m_{test} = 100;
[A,b] = genTraining_data(m, false);
[Atest,btest] = genTraining_data(m_test,false);
%% compute svm with rbf or poly kernel
% subplot(1,2,1)
svm.rbf = svmtrain(A,b, 'ShowPlot', 0, 'kernel_function', 'rbf');
% subplot (1, 2, 2)
svm.poly = svmtrain(A,b, 'ShowPlot', 0, 'kernel_function', 'polynomial');
\%\% compute the accuracy for rbf and poly kernelized \ensuremath{\mathrm{svm}}
prediction.rbf = nan(m_test,1);
prediction.poly = nan(m_test,1);
for i = 1 : m_test
    Xnew = Atest(i,:);
    prediction.rbf(i) = svmclassify(svm.rbf, Xnew, 'ShowPlot', 0);
    prediction.poly(i) = svmclassify(svm.poly, Xnew, 'ShowPlot', 0);
% compute accuracy
accuracy.rbf = sum(prediction.rbf == btest) / length(btest);
accuracy.poly = sum(prediction.poly == btest) / length(btest);
%% fit standard least square model and compute the accuracy
wts.ls = pinv(A) * b;
prediction.ls = sign(Atest*wts.ls);
accuracy.ls = sum(prediction.ls == btest)/length(btest);
```

Consider the problem of trying to predict whether a person is a basketball player based on height. The training data consists of four people with heights of 5'10", 5'11', 6'1" and 6'10", and the latter two are basketball players. What classification rule do you obtain by minimizing hinge loss instead of squared error loss?

First of all, let me convert feet to meters, and define the resulting matrix as my data matrix

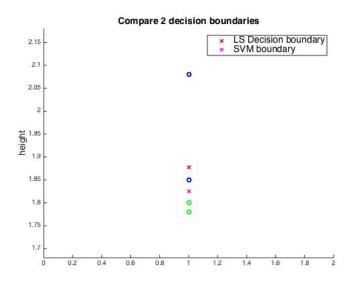
$$\begin{bmatrix} 5'10" \\ 5'11' \\ 6'1" \\ 6'10" \end{bmatrix} \rightarrow \begin{bmatrix} 1.78 \\ 1.8 \\ 1.85 \\ 2.08 \end{bmatrix} =: X$$

And define y as the following:

$$\begin{bmatrix} \text{not a basketball player} \\ \text{not a basketball player} \\ \text{basketball player} \\ \text{basketball player} \end{bmatrix} \rightarrow \begin{bmatrix} -1 \\ -1 \\ 1 \\ 1 \end{bmatrix} =: y$$

Then we can use X and y to do classification. Now I want to compare LS boundary with the SVM boundary.

In the following plot, the LS and SVM boundaries were plotted with 'x' in different colors and data were shown using circles with different colors.



Because LS has positive cost for examples that are classified too well, the very tall guy is pulling the decision boundary up, resulting a bad boundary. The SVM boundry has zero cost for correctly classified examples so the decision boundary is insensitive to the very tall guy.

Here's the matlab code that generate the simulation in question 6.

```
clear all;close all;clc;
X = [1 \ 1 \ 1 \ 1; \ 1.78 \ 1.8 \ 1.85 \ 2.08]';
y = [-1 \ -1 \ 1 \ 1]';
%% fitting LS
w.ls = inv(X' * X) * X' * y;
sign(X * w.ls)
%% fit svm
tau = 0.03;
lambda = 0;
w.svm = zeros(size(X, 2), 1);
numIters = 100000;
% implement gradient decent
for i = 1 : numIters
    change = zeros(size(X, 2), 1);
    % accumulate the gradient for all training examples
    for j = 1 : length(y)
        if 1 - y(j) * X(j,:) * w.svm > 0
            change(:) = change(:) - y(j) * X(j,:)';
    end
    % gradient decent update
    w.svm = w.svm - tau * change;
    fprintf('%d %f \n', i, norm(y - X * w.svm));
end
%% plot the decision boundary
hold on
plot(-w.ls(1)/w.ls(2), 'rx', 'linewidth', 2)
plot(-w.svm(1)/w.svm(2), 'mx', 'linewidth', 2)
title('Compare 2 decision boundaries', 'fontsize', 14)
ylabel('height', 'fontsize', 14)
legend({'LS Decision boundary', 'SVM boundary'} ,'fontsize', 14)
ylim([min(X(:,2))-.1 max(X(:,2))+.1])
% plot the points
for i = 1 : size(X, 1)
    if y(i) > 0
        plot(X(i,2), 'bo', 'linewidth', 2)
        plot(X(i,2), 'go', 'linewidth', 2)
    end
end
hold off
```

If you consider the dual solution in this case, which values of $\alpha \in \mathbb{R}^4$ are nonzero (i.e. what are the support vectors?)

The alpha associated with the 2nd and the 3rd examples will be non-zero. Namely, the example that is closest from the sym decision boundary on each side are the support vectors.

The standard hinge loss is linearly decreasing up to 1 and then exactly 0 after. Let t_0 denote the point where the hinge loss first becomes 0. Does the solution change if you shift the t_0 within the range $t_0 \in [0, 1]$?

Yes it would change.

When t_0 is 1, things larger than 1 has 0 loss, things less 0 than has a high cost, and things with in [0,1] has a small loss. Therefore, we are acutally penalizing things that are classified correctly with in [0,1]. This is really saying that we want more than merely classifying things correctly, we want to be right by at least 1, which creates the margin.

If t_0 shrinks, then the margin would shrink. Consider the extreme when t_0 is 0, then everything classified corrected would have no cost. Assume the data are linearly separable, there can exist many decision boundaries with zero loss, and the classifier no longer care about maximizing the margin.