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- This is solved by taking the cluster which has its mean closest to the test point  $x_n$

$$\arg \min_k ||x_n - \mu_k||^2$$

- If we assume the test point to be closest to a cluster mean, denoted by  $\mu_k$  then we can get the update equation for  $\mu_k$  by taking derivative of  $\mathcal{L}$  w.r.t.  $\mu_k$

$$\frac{\partial \mathcal{L}}{\partial \mu_k} = -2||x_n - \mu_k||$$

Which can be put into the update equation as

$$\mu_k = \mu_k + \eta ||x_n - \mu_k|| \tag{1}$$

- In 1 we have taken all constants to be part of the step-size  $\eta$ . A good choice of  $\eta$  would be a small value that decreases monotonically as the steps progress. By taking a small step size the cluster means will slowly progress towards the expected means and remain unaffected by noisy input datapoints.

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As  $z_n \in \mathbb{R}$  is a linear transformation of  $\mathbf{x}_n$  and  $\mathbf{w}$  we can write:

$$z_n = \mathbf{w}^T \mathbf{x}_n$$

We need to minimise distance between each projected point  $z_n$  and its cluster mean, say  $\{\mu_-, \mu_+\}$  and maximize distance between the projected means.

$$\max |\mu_- - \mu_+| \tag{2}$$

$$\min \sum_{z_n: y_n = +1} |z_n - \mu_+| + \sum_{z_n: y_n = -1} |z_n - \mu_-| \tag{3}$$

Thus the objective function is:

$$J = \max \left[ |\mu_- - \mu_+| - \sum_{z_n: y_n = +1} |z_n - \mu_+| - \sum_{z_n: y_n = -1} |z_n - \mu_-| \right] \tag{4}$$

as all values are in  $\mathbb{R}$  taking only simple absolute distance suffices.

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Let us take  $\mathbf{S}' = \frac{1}{N}\mathbf{X}\mathbf{X}^T$ . Equation 5 represents the equation to calculate eigenvalue  $\lambda'$  and eigenvector  $\mathbf{v}$  of  $\mathbf{S}'$

$$\mathbf{S}'\mathbf{v} = \lambda'\mathbf{v} \quad (5)$$

Now if we take the value of  $\mathbf{S}'$  and pre-multiply with  $\mathbf{X}^T$  and readjust the values we get:

$$\frac{1}{N}\mathbf{X}\mathbf{X}^T\mathbf{v} = \lambda'\mathbf{v} \quad (6)$$

$$\frac{1}{N}\mathbf{X}^T\mathbf{X}\mathbf{X}^T\mathbf{v} = \lambda'\mathbf{X}^T\mathbf{v} \quad (7)$$

$$\mathbf{S}\mathbf{u} = \lambda'\mathbf{u} \quad (8)$$

Thus the eigenvalue remains the same in both forms, only the eigenvectors change, which can be seen in blue from Equation 7 and 8. By computing eigenvectors this way we can reduce the complexity of calculating eigenvalues for a  $D \times D$  matrix to  $N \times N$  matrix, which is feasible in this case as  $D < N$ .

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## Part 1

In practical scenarios the input data need not be a part of a single distribution. By modelling with latent input variables  $\mathbf{z}_n$  we can leverage the fact that the input is coming from different distributions and so we can apply a combination of  $K$  different linear models to predict  $y_n$  from  $\mathbf{x}_n$ . This can also be useful in outlier detection as the outlier input point will be a part of its own cluster.

## Part 2

The latent variable model is defined in Equation 9

$$p(\mathbf{z}_n|\mathbf{x}_n, \Theta) = \frac{p(\mathbf{z}_n|\Theta)p(\mathbf{x}_n|\mathbf{z}_n, \Theta)}{p(\mathbf{x}_n|\Theta)} \quad (9)$$

Upon applying the variables and distributions from the question we get Equation 10. Now that we only have to take MLE solution we can reduce the same to 11

$$p(z_n = k|y_n, \Theta) = \frac{p(z_n = k|\Theta)p(y_n|z_n = k, \Theta)}{p(y_n|\Theta)} \quad (10)$$

$$= p(z_n = k|\Theta)p(y_n|z_n = k, \Theta) \quad (11)$$

where:

$$p(z_n = k|\Theta) = \pi_k \quad (12)$$

$$p(y_n|z_n = k, \Theta) = \mathcal{N}(w_{z_n}^T x_n, \beta^{-1}) \quad (13)$$

Taking the log of equation 11 with the provided distributions we get:

$$\log p(z_n = k|y_n, \Theta) = \log \pi_k + \log \mathcal{N}(w_{z_n}^T x_n, \beta^{-1}) \quad (14)$$

### ALT-OPT ALGORITHM

Step 1: Find the best  $z_n$

$$\hat{z}_n = \arg \max_{k=1,2,\dots,K} \log \pi_k + \log \mathcal{N}(w_{z_n}^T x_n, \beta^{-1}) \quad (15)$$

$$= \arg \max_{k=1,2,\dots,K} \log \pi_k + \frac{\beta}{2} (y_n - w_{z_n}^T x_n)^2 \quad (16)$$

Step 2: re-estimate the parameters

$$\pi_k = \frac{N_k}{N} \quad (17)$$

$$w_k = (X_k^T X_k)^{-1} X_k^T y_k \quad (18)$$

Where  $N_k$  is the number of points in cluster  $k$

## 1 Kernel Ridge Regression

### 1.1 Regularized Kernel Ridge Regression

I calculated the output for the test-point by using Equation 19.

$$\mathbf{Y}^* = \mathbf{K}^*(\mathbf{K} - \lambda \mathbf{I}_N)^{-1} \mathbf{Y} \quad (19)$$

where  $\mathbf{K}^*$  and  $\mathbf{K}$  are the kernel matrices:  $\mathbf{K}^* = k(x^*, x_n)$ ,  $\mathbf{K} = k(x_m, x_n)$  and  $k(x_m, x_n) = \exp(-\gamma \|x_n - x_m\|^2)$  is the rbf kernel function. The RMSE value is calculated using:

$$\sqrt{\frac{1}{N} \|y_{true} - y_{pred}\|^2}$$

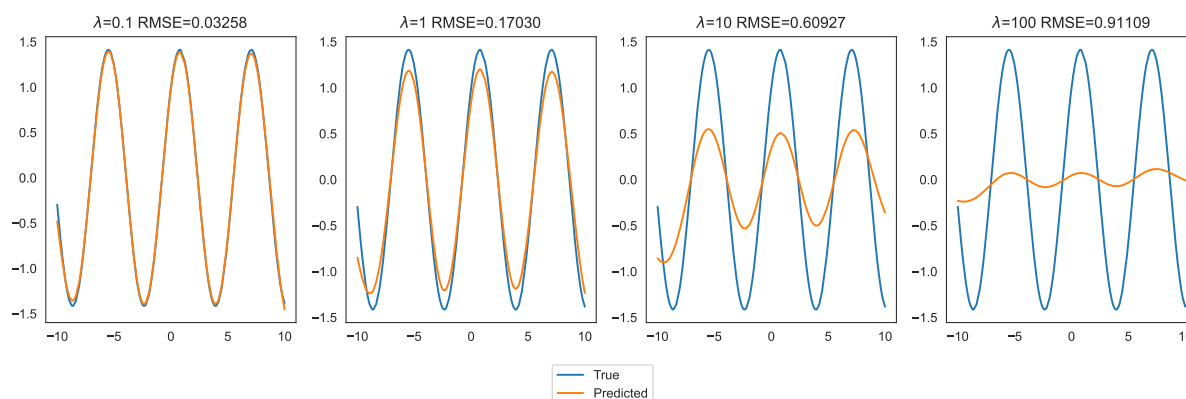


Figure 1: Kernel ridge regression with different regularizations

From Figure 1 we can see that as we increase the regularization hyperparameter  $\lambda$  from 0.1 to 100 we see that the predictions are not able to accurately predict the noiseless y-values. Adding regularization makes the model resistant to noise and improves test-accuracy, but in this case the y-values in test data are noiseless so it is better to overfit the model to reduce RMSE value.

### 1.2 Landmark Kernel Ridge Regression

Here I randomly chose 2, 5, 20, 50 and 100 landmark points from train data and ran the same kernel ridge regression code on the test data. From Figure 2 we can see that as the number of Landmark points increases the regression model starts to predict the model in a better way, the RMSE reduces.

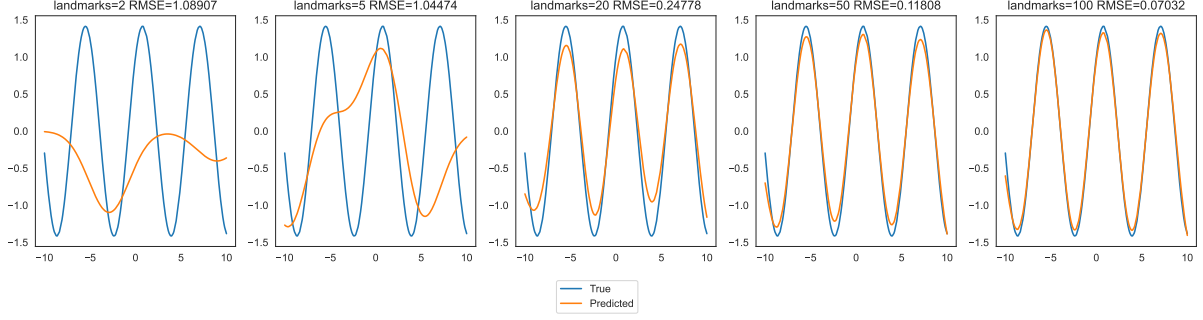


Figure 2: Landmark kernel ridge regression with different number of landmark points

## 2 K-Means Clustering

In Figure 3 we see that the data is circular and there are 2 intuitive clusters that are possible. So I calculated the radius of each point by applying 20 and 21 and then projecting all the points on 1-dimension to get Figure 4 and applying k-means on the projected data to obtain Figure 5

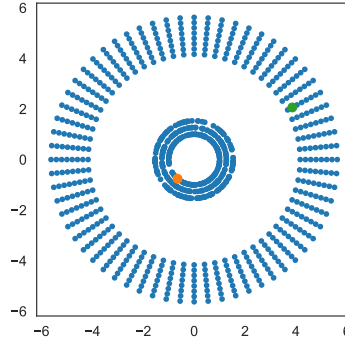


Figure 3: Scatter plot of data with initial centers highlighted

$$\theta = \tan^{-1} \left( \frac{y}{x} \right) \quad (20)$$

$$r = \frac{x}{\cos(\theta)} \quad (21)$$

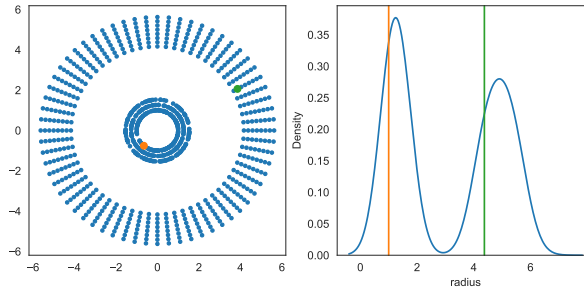


Figure 4: Projecting data points into 1-dimension by taking radius of points from origin. Also highlighting the initial centers

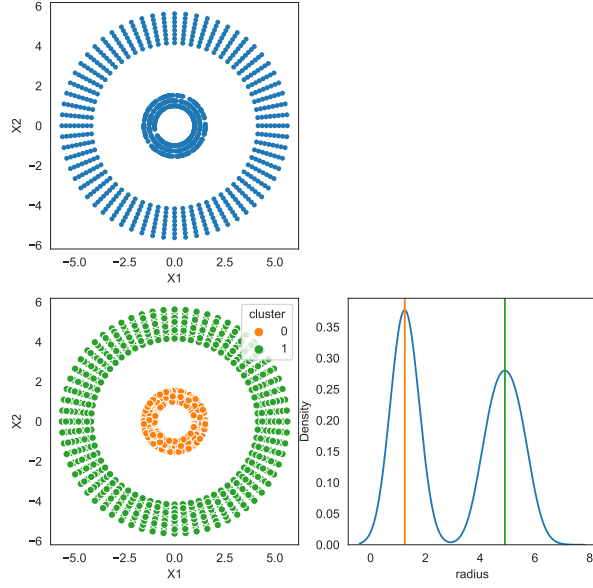


Figure 5: Clustering the data points based on radius and showing the position of the cluster centers on the 1-dimensional plot.

Next I chose 1 landmark point randomly and added another feature based on RBF kernel function. Then I ran a k-means prediction algorithm which led to the good and bad clustering. Figure 6 shows all the clustered plots.

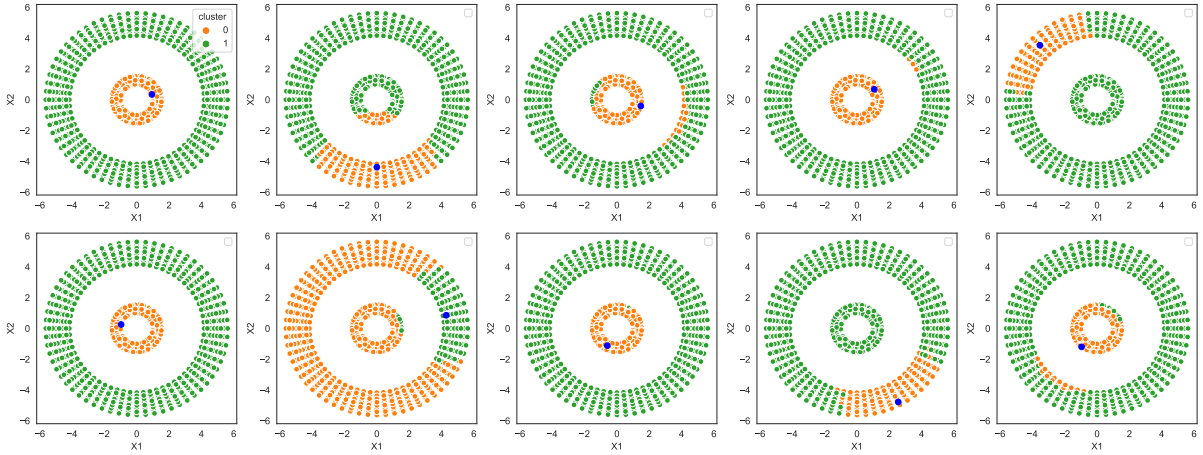


Figure 6: Applying k-means with 1 training landmark point.

### 3 PCA and TSNE

TSNE is able to make clusters which are naturally observable on 2 dimensions in a better way compared to PCA. In PCA, we can see significant overlap and mixing of points from different clusters. Thus, TSNE is a better projection technique to visualize high-dimensional data.

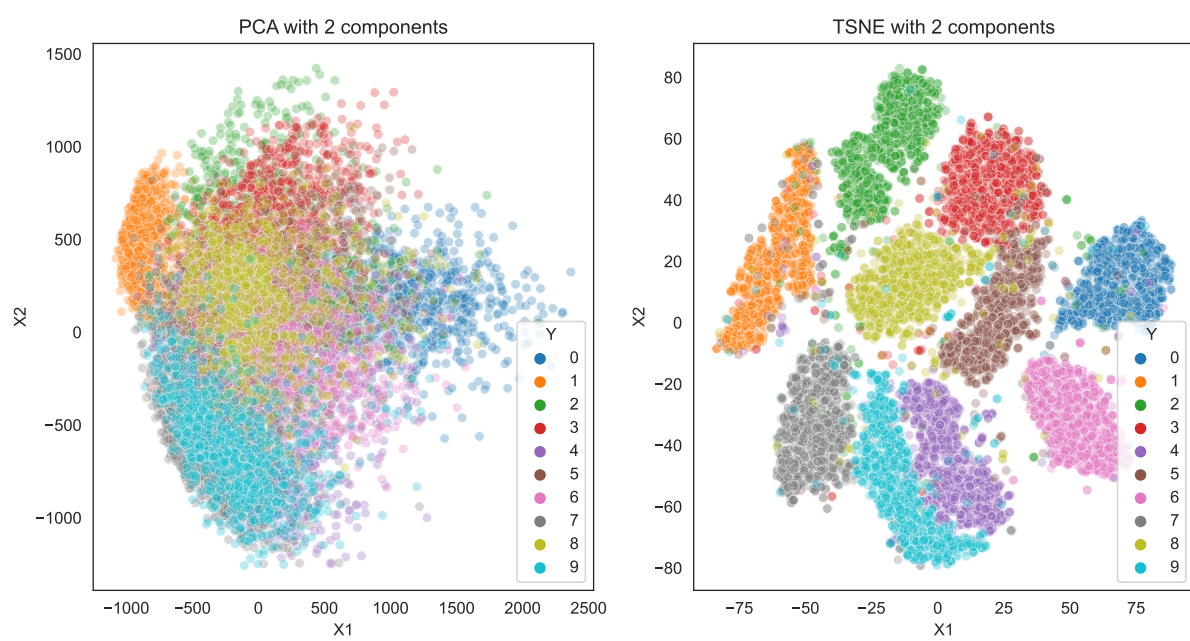


Figure 7: 2-Dimensional PCA and TSNE plots of MNIST