

KMeans Kernel Classifier

Course: Math Behind ML

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Abstract—The least squares SVM is a kernel method for non-linear regression and classification tasks. Here we combine KMeans clustering with the least squares SVM. First KMeans clustering is used to extract a set of representative vectors for each class, and then LS-SVM uses these representative vectors as a training dataset for the classification task

- It is faster than LS-SVM.
- It is more robust.
- It is very easy to implement.

I. INTRODUCTION

The kernel methods transform a given non-linear problem into a linear one by using a similarity kernel function $\Omega(x, x')$. It is a similarity function defined over pairs of input data points (x, x') . This way the input data is mapped into a higher dimensional feature space $\phi(x)$, where the inner product $\langle \cdot, \cdot \rangle$ can be calculated using Mercer's condition:

$$\Omega(x, x') = \langle x, x' \rangle \quad (1)$$

Consider $\chi = \{x_n | n = 1, \dots, N\}$ as training dataset.

Representer theorem: Any non-linear function $f : \chi \rightarrow \mathbb{R}$ can be expressed as linear combination of kernel products on training dataset which was mentioned above earlier.

$$f(x) = \sum_{n=1}^N a_n \Omega(x, x_n) \quad (2)$$

Time complexity of LS-SVM is $O(N^3)$ where N is size of the training dataset which is too high and makes it unsuitable for large dataset. So for this reason we use KMeans clustering to extract a set of representative vectors for each class, and then LS-SVM uses these representative vectors as a training dataset for the classification task. This way we can reduce the time complexity of LS-SVM to $O(K^3)$ where K is the number of clusters. These representative vectors are called as **centroids**. These are then used by LS-SVM to classify the test data. This KMeans-LS-SVM method has some advantages:

II. KERNEL LS-SVM CLASSIFIER

We already know that in binary classification, kernel SVM method constructs an hyperplane with the maximal margin between the two classes in feature space $\phi(x)$. This can be represented as convex quadratic programming problem involving inequality constraints.

The kernel LS-SVM simplifies the optimization problem by considering equality constraints only, such that solution is obtained by solving a system of linear equations. Now this problem is similar to ridge regression problem which is formulated as follows:

$$\min_{w, b} \frac{1}{2} w^T w + \frac{\gamma}{2} \sum_{n=1}^N (\hat{y}_n - w^T \phi(x_n) - b)^2 \quad (3)$$

Assume that K classes are encoded using standard basis in \mathbb{R}^K , i.e, let $x_i \in C_k$, then output y_i is a vector with 1 in the k^{th} position and 0 elsewhere:

$$y_{ij} = \begin{cases} 1 & \text{if } x_i \in C_j \\ 0 & \text{otherwise} \end{cases} \quad (4)$$

Consider input data $\{(x_i, y_i) | x_i \in \mathbb{R}^M, y_i \in \mathbb{R}^K, i = 1, \dots, N\}$ and the feature mapping function $\phi(x)$. The kernel LS-SVM is formulated as follows:

$$\min_{w, b} S(w, b, \epsilon) = \frac{1}{2} \sum_{j=1}^K w_j^T w_j + \frac{\gamma}{2} \sum_{i=1}^N \sum_{j=1}^K (\epsilon_{ij})^2 \quad (5)$$

subject to

$$\langle \phi(x), \omega_j \rangle + b_j = y_{ij} - \epsilon_{ij}, i = 1, \dots, N; j = 1, \dots, K \quad (6)$$

$$w_j^T \phi(x_i) + b_j = y_{ij} - \epsilon_{ij}, i = 1, \dots, N; j = 1, \dots, K \quad (7)$$

where $\epsilon_{ij} \geq 0$ are approximation errors, b_j is bias coefficient, $w^{(j)}$ is the vector of weights corresponding to the j^{th} class. The objective function S is a sum of least squares errors and the regularization term. This regularization parameter γ corresponds to a multi-dimensional version of the ridge regression problem.

In the primal weight space the multi class classifier takes the form:

$$x \in C_k, \Leftrightarrow k = \arg \max_{j=1, \dots, K} g_j(x) \quad (8)$$

$$\text{where } g_j(x) = \frac{\exp(\langle \phi(x), w^{(j)} \rangle + b_j)}{\sum_{i=1}^K \exp(\langle \phi(x), w^{(i)} \rangle + b_i)} \quad (9)$$

Here g_j is the non-linear soft max function

Now applying Lagrangian to (5)

$$L(w, b, \epsilon, a) = S(w, b, \epsilon) - \sum_{i=1}^N \sum_{j=1}^K a_{ij} [\langle \phi(x), \omega_j \rangle + b_j - y_{ij} + \epsilon_{ij}]$$

where $a_{ij} \in \mathbb{R}$ is the lagrange multiplier. Now applying KKT conditions:

$$\frac{\partial L}{\partial w^{(j)}} = 0 \Rightarrow w^{(j)} = \sum_{n=1}^N a_{nj} \phi(x_n) \quad (10)$$

$$\frac{\partial L}{\partial b_{(j)}} = 0 \Rightarrow \sum_{i=1}^N a_{ij} = 0 \quad (11)$$

$$\frac{\partial L}{\partial \epsilon_{(ij)}} = 0 \Rightarrow a_{ij} = \gamma \epsilon_{ij} \quad (12)$$

$$\frac{\partial L}{\partial a_{(ij)}} = 0 \Rightarrow \langle \phi(x), \omega_j \rangle + b_j - y_{ij} + \epsilon_{ij} = 0 \quad (13)$$

Now from eq(10), eq(12) and eq(13):

$$\sum_{n=1}^N [\Omega(x_i, x_n) + \gamma^{-1} \delta_{in}] a_{nj} + b_j = y_{ij}, \quad (14)$$

Here δ_{in} is the Kronecker delta function: where $\delta_{in} = 1$ if $i = n$ and 0 otherwise

As you can see in eq(14) there are K independent system of equations with binary labels y_{ij} . Now each system can be written in the matrix form as follows:

$$\begin{bmatrix} 0 & u^T \\ u & \Omega + \gamma^{-1} I \end{bmatrix} \begin{bmatrix} b_j \\ a^{(j)} \end{bmatrix} = \begin{bmatrix} 0 \\ y_j \end{bmatrix}, j = 1, \dots, K \quad (15)$$

Here $I_{N \times N}$ is the identity matrix, $u_{N \times 1} = [1, \dots, 1]^T$ is a vector of ones, $a_{N \times 1}^{(j)} = [a_{1j}, \dots, a_{Nj}]^T$ is weights and $y_j = [y_{1j}, \dots, y_{Nj}]^T$ is the vector of binary labels for the j^{th} class.

Each system has $N + 1$ linear equations with $N + 1$ unknowns. Each system has $N + 1$ linear equations with $N + 1$ unknowns.

$$\Theta = \begin{bmatrix} 0 & u^T \\ u & \Omega + \gamma^{-1} I \end{bmatrix} \quad (16)$$

All the K systems can be written as:

$$\Theta W = Z \quad (17)$$

where

$$W_{(N+1) \times K} = \begin{bmatrix} b_1 & \dots & b_K \\ a^{(1)} & \dots & a^{(K)} \end{bmatrix}, Z_{(N+1) \times K} = \begin{bmatrix} 0 & \dots & 0 \\ y_1 & \dots & y_K \end{bmatrix}$$

Now once all the K systems are solved, we consider multi-class classifier in dual space(from eq (14)) as follows:

$$g_j(x) = \frac{\exp(\langle \phi(x), w^{(j)} \rangle + b_j)}{\sum_{i=1}^K \exp(\langle \phi(x), w^{(i)} \rangle + b_i)}$$

From eq(9) and eq(10), we get:

$$g_j(x) = \frac{\sum_{n=1}^N \exp(\Omega(x, x_n) a_{nj} + b_j)}{\sum_{i=1}^K \sum_{n=1}^N \exp(\Omega(x, x_n) a_{ni} + b_i)}$$

Now our problem becomes:

$$x \in C_k, \Leftrightarrow k = \arg \max_{j=1, \dots, K} g_j(x) \quad (18)$$

where

$$g_j(x) = \frac{\sum_{n=1}^N \exp(\Omega(x, x_n) a_{nj} + b_j)}{\sum_{i=1}^K \sum_{n=1}^N \exp(\Omega(x, x_n) a_{ni} + b_i)}$$

Here g_j is the non-linear soft max function

III. KMEANS CLUSTERING

First we use KMeans clustering algorithm to extract a set of representative vectors for each class. Now this representative vectors will be passed into LS-SVM kernel model as training dataset. KMeans clustering algorithm is as follows:

- 1) Take $\{x_i^k | x_i^k \in \mathbb{R}^M, i = 1, \dots, N_k\}$ as training samples for class C_k where N_k is the number of training samples for the class C_k and $N = \sum_{k=1}^K N_k$ is the total number of training samples.
- 2) Take $\{\mu_q^k | \mu_q^k \in \mathbb{R}^M, q = 1, \dots, Q\}$ as initial centroids for class C_k where $Q < N_K$ is the number of centroids for class C_k .
- 3) Build a matrix $X_k = [x_{im}^k]_{N_k \times M}$ where each row is a training sample for class C_k .
- 4) Build a matrix $\Xi_k = [\xi_{qm}^k]_{Q \times M}$ where each row is a randomly initialized centroid ' for class C_k .
- 5) Let $R_k = X_k \Xi_k^T = [r_{iq}^k]_{N_k \times Q}$
- 6) Let $\hat{R}_k = [\hat{r}_{iq}^k]_{N_k \times Q}$ be transformed sparse matrix of R_k where:
$$\hat{r}_{iq}^k = \begin{cases} 1 & \text{if } q = \arg \max_q r_{iq}^k \\ 0 & \text{otherwise} \end{cases} \quad i = 1, \dots, N_k$$
Each sample is assigned to the nearest centroid.
- 7) $\hat{\Xi}_k = \hat{R}_k^T X_k = [\hat{\xi}_{qm}^k]_{Q \times M}$.
This is the new set of centroids.

8) Normalizing new set of centroids:

$$\hat{\xi}_q^k = \frac{\xi_q^k}{\|\xi_q^k\|},$$

$$q = 1, \dots, Q$$

9) Computing alignment deviation between new set and old set of centroids:

$$\delta = 1 - \frac{\sum_{q=1}^Q \langle \hat{\xi}_q^k, \xi_q^k \rangle}{Q}$$

10) $\Xi_k = \hat{\Xi}_k$

11) Repeat steps 5 to 10 until $\delta < \beta$ where β is the tolerance.

12) Return Ξ_k

IV. KMEANS KERNEL LS-SVM CLASSIFIER

After extracting a set of representative vectors for each class $C_k, k = 1, \dots, K$ using KMeans clustering, we pass these KQ centroids into LS-SVM kernel model as training dataset.

Training dataset for LS-SVM before KMeans clustering:

$$\{(x_i^k, y_i^k) | x_i^k \in \mathbb{R}^M, y_i^k \in \mathbb{R}^K, i = 1, \dots, N\}$$

Training dataset for LS-SVM after KMeans clustering:

$$\{(\xi_q^k, y_q^k) | \xi_q^k \in \mathbb{R}^M, y_q^k \in \mathbb{R}^K, q = 1, \dots, KQ\}$$

As you can see the training dataset size is reduced from N to KQ where $KQ < N$.

Previously there were $N + 1$ linear equations with $N + 1$ unknowns and $O(N^3)$ time complexity.

Now there are $KQ + 1$ linear equations with $KQ + 1$ unknowns and $O((KQ)^3)$ time complexity.

As we discussed earlier our problem previously was:

$$x \in C_k, \Leftrightarrow k = \arg \max_{j=1, \dots, K} g_j(x)$$

$$\text{where } g_j(x) = \frac{\sum_{n=1}^N \exp(\Omega(x, x_n) a_{nj} + b_j)}{\sum_{i=1}^K \sum_{n=1}^N \exp(\Omega(x, x_n) a_{ni} + b_i)}$$

Now our problem becomes:

$$x \in C_k, \Leftrightarrow k = \arg \max_{j=1, \dots, K} g_j(x)$$

$$\text{where } g_j(x) = \frac{\sum_{n=1}^{KQ} \exp(\Omega(x, \xi_n^k) a_{nj} + b_j)}{\sum_{i=1}^K \sum_{n=1}^{KQ} \exp(\Omega(x, \xi_n^k) a_{ni} + b_i)}$$

Here g_j is the non-linear soft max function

V. APPLICATION

VI. CONCLUSION