## Task 3. Relevance Vector Machine

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## 1 Introduction

SVM are a popular choice in classification problems. However they are inherently not probabilistic. A technique that uses the same intuition is the Relevance Vector Machine. It uses a full Bayesian approach with a prior that encourages sparseness.

## 2 The problem

First a dual model is used, previously we would model the data as a normal distribution. Where w is the world state, x is the data point,  $\phi$  are the paremeters of a linear function of the data and  $\sigma$  is the standard deviation. We now apply a transformation on the parameters as shown in Equation 1, so  $\phi$  is now a weighted sum over the observed data points, where  $\psi$  is a vector with the weights. If we are working with few data of high dimensionality, this sum is faster to calculate, as it has less terms, that the original linear dependency.

$$Pr(w_i|\mathbf{x}_i) = Norm_{x_i}[\phi^T \mathbf{x}_i, \sigma^2], \tag{1}$$

$$\phi = \mathbf{X}\psi,\tag{2}$$

Using the dual parameters  $\psi$ , we encourage sparsernes in the model by using the prior defined in Equation 3. Where I is the number of data points, Stud is a Student's t-distribution with  $\nu$  degrees of freedom and  $\Gamma$  is the Gamma function. Using a product of t-distributions produces the desired sparseness since the areas with higher probability density are in the origin and along the axis.

$$Pr(\psi) = \prod_{i=1}^{I} Stud_{\psi_i}[0, 1, \nu],$$
 (3)

$$= \prod_{i=1}^{I} \frac{\Gamma\left(\frac{\nu+1}{2}\right)}{\sqrt{\nu\pi}\Gamma\left(\frac{\nu}{2}\right)} \left(1 + \frac{\psi_i^2}{\nu}\right)^{-(\nu+1)/2},\tag{4}$$

A t-distribution is not conjugate to a Normal distribution, so there is no simple closed form solution for the posterior. The solution will be to approximate the t-distributions by maximizing with respect to their hidden variables h, introduced in Equation 5, where  $\mathbf{w}$  is a one dimensional array with the world state. This leads to the marginal likelihood shown in Equation 6, where  $\mathbf{I}$  is the identity matrix,  $\mathbf{X}$  is a matrix with the data points and  $\mathbf{H}$  is a matrix with all the hidden variables.

$$Pr(\psi) = \prod_{i=1}^{I} \int Norm_{\mathbf{w}}[0, 1/h_i] Gam_{h_i}[\nu/2, \nu/2] dh_i,$$
 (5)

$$Pr(\mathbf{w}|\mathbf{X}, \sigma^2) \approx \max_{\mathbf{H}} \left[ Norm_{\mathbf{w}} \left[ \mathbf{0}, \mathbf{X}^T \mathbf{X} \mathbf{H}^{-1} \mathbf{X}^T \mathbf{X} + \sigma^2 \mathbf{I} \right] \prod_{d=1}^{D} Gam_{h_d} \left[ \nu/2, \nu/2 \right] \right],$$
 (6)

The optimization is performed in three steps:

1. Optimize the marginal likelihood with respect to the hidden variables, using Equation 7.

$$h_i^{new} = \frac{1 - h_i \sum_{ii} + \nu}{\mu_i^2 + \nu},\tag{7}$$

2. Update  $\mu$  and  $\Sigma$ , using Equation 8.

$$\mu = \frac{1}{\sigma^2} \mathbf{A}^{-1} \mathbf{X}^T \mathbf{X} \mathbf{w},$$

$$\Sigma = \mathbf{A}^{-1},$$

$$\mathbf{A} = \frac{1}{\sigma^2} \mathbf{X}^T \mathbf{X} \mathbf{X}^T \mathbf{X} + \mathbf{H},$$
(8)

3. Optimize the marginal likelihood with respect to the variance parameter, using Equation 9.

$$(\sigma^2)^{new} = \frac{1}{\mathbf{I} - \sum_{i} (1 - h_i \sum_{i})} \left( \mathbf{w} - \mathbf{X}^T \mathbf{X} \boldsymbol{\mu} \right)^T \left( \mathbf{w} - \mathbf{X}^T \mathbf{X} \boldsymbol{\mu} \right), \tag{9}$$

After the training process, we only take the data points whose hidden variable  $h_i$  is smaller than a threshold. Since a larger one means a small  $\psi_i$ , and hence no significance in the weighted sum. In the implementation, we use a Gaussian kernel  $k[x_i, x_j]$ , as shown in Equation 10, instead of calculating the inner products  $\mathbf{X}^T\mathbf{X}$ , where  $\lambda$  controls the scale of the output. This is more efficient than computing the inner product.

$$k[x_i, x_j] = exp\left[-\frac{1}{2}\left(\frac{(x_i - x_j)^T(x_i - x_j)}{\lambda^2}\right)\right],\tag{10}$$

Since our world states  $\mathbf{w_i}$  are multivariate, a separate regressor is used for each dimension. However the hidden variables matrix  $\mathbf{H}$  is shared among all of them.

## 3 Results

In Figures 1 and 2 we show results for a font data set. The RVM is trained with pictures of n in different fonts and their corresponding m in the same font. For testing, n with new fonts are used.

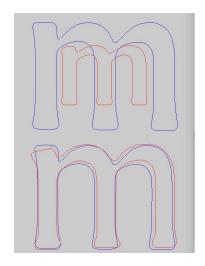


Figure 1: Blue is ground truth and red is RVM result. Top image corresponds to a kernel  $\lambda = 2$ , bottom image has  $\lambda = 7$ .

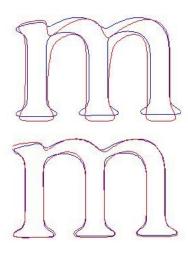


Figure 2: Blue is ground truth and red is RVM result. Top image corresponds  $\nu=0.001$ , bottom image has  $\nu=1$ 

In Figure 1 we see the effects of changing the value of  $\lambda$ . Since it controls the scale, we can see how the m varies in size. In Figure 2 variations in  $\nu$  are shown. Smaller  $\nu$  encourages more sparsity, and so less relevance vectors been used. For the top image 5 relevance vectors are used, while for the bottom one 10 vectors are used. Less vectors means a more general and sparse model used. However the error in the prediction increases as we take fewer relevance vectors.