

# CS 524: Introduction to Optimization

## Lecture 31 : SVM & cross-validation

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## Choosing a value of the tuning parameter

Each regularization method has an associated **tuning parameter**: e.g., this was  $\lambda$  in the smoothing spline problem, and  $\lambda$  for lasso and ridge regression in the penalized forms (or  $t$  in the constrained forms)

The tuning parameter controls the amount of regularization, so choosing a good value of the tuning parameter is crucial. Because each tuning parameter value corresponds to a fitted model, we also refer to this task as **model selection**

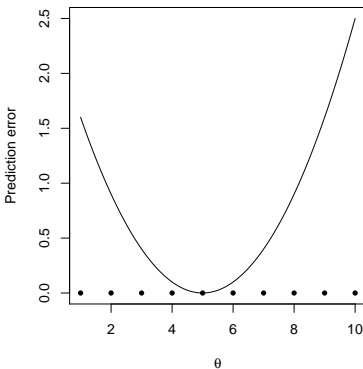
What we might consider a good choice of tuning parameter, however, depends on whether our goal is **prediction accuracy** or **recovering the right model** for interpretation purposes. We'll cover choosing the tuning parameter for the purposes of prediction; choosing the tuning parameter for the latter purpose is a harder problem

# Cross-validation

**Cross-validation** is a simple, intuitive way to estimate prediction error

Given training data  $(a_i, y_i)$ ,  $i = 1, \dots, m$  and an estimator  $\phi_\theta$  depending on a tuning parameter  $\theta$

Even if  $\theta$  is a continuous parameter, it's usually not practically feasible to consider all possible values of  $\theta$ , so we discretize the range and consider choosing  $\theta$  over some discrete set  $\{\theta_1, \dots, \theta_p\}$



For a number  $K$ , we split the training pairs into  $K$  parts or “folds” (commonly  $K = 5$  or  $K = 10$ )

1	2	3	4	5
Train	Train	Validation	Train	Train

**$K$ -fold cross validation** considers training on all but the  $k$ th part, and then validating on the  $k$ th part, iterating over  $k = 1, \dots, K$

(When  $K = m$ , we call this **leave-one-out cross-validation**, because we leave out one data point at a time)

## $K$ -fold cross validation procedure:

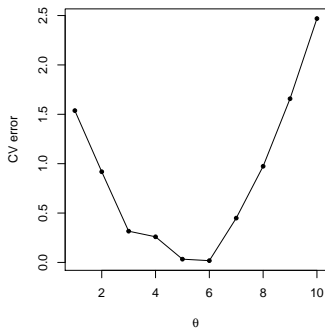
- Divide the set  $\{1, \dots, m\}$  into  $K$  subsets (i.e., folds) of roughly equal size,  $F_1, \dots, F_K$
- For  $k = 1, \dots, K$ :
  - ▶ Consider training on  $(a_i, y_i)$ ,  $i \notin F_k$ , and validating on  $(a_i, y_i)$ ,  $i \in F_k$
  - ▶ For each value of the tuning parameter  $\theta \in \{\theta_1, \dots, \theta_p\}$ , compute the estimate  $\phi_\theta^{-k}$  on the training set, and record the total error on the validation set:

$$e_k(\theta) = \sum_{i \in F_k} (y_i - \phi_\theta^{-k}(a_i))^2$$

- For each tuning parameter value  $\theta$ , compute the average error over all folds,

$$CV(\theta) = \frac{1}{m} \sum_{k=1}^K e_k(\theta) = \frac{1}{m} \sum_{k=1}^K \sum_{i \in F_k} (y_i - \phi_\theta^{-k}(a_i))^2$$

Having done this, we get a **cross-validation error curve**  $CV(\theta)$  (this curve is a function of  $\theta$ ), e.g.,



and we choose the value of tuning parameter that minimizes this curve,

$$\hat{\theta} = \arg \min_{\theta \in \{\theta_1, \dots, \theta_p\}} CV(\theta)$$

## Recap: cross validation

**Training error**, the error of an estimator as measured by the data used to fit it, is not a good surrogate for prediction error. It just keeps decreasing with increasing model complexity

**Cross-validation**, on the other hand, much more accurately reflects prediction error. If we want to choose a value for the tuning parameter of a generic estimator (and minimizing prediction error is our goal), then cross-validation is the standard tool

We usually pick the tuning parameter  $\theta$  that minimizes the cross-validation error curve. Sometimes called “Hyper-parameter estimation”

# Cross-validation is a general tool

So far we've looked at cross-validation for estimation under squared error loss, but it applies much **more broadly** than this.

For an arbitrary loss  $\ell(y_i - \phi(a_i))$ , the cross-validation estimate of prediction error under  $\ell$  is

$$\frac{1}{n} \sum_{k=1}^K \sum_{i \in F_k} \ell(y_i - \phi^{-k}(a_i))$$

E.g., for classification, each  $y_i \in \{0, 1\}$ , and we might want to use the 0-1 loss

$$\ell(y_i - \phi(a_i)) = \begin{cases} 0 & \text{if } y_i = \phi(a_i) \\ 1 & \text{if } y_i \neq \phi(a_i) \end{cases}$$

Cross-validation now gives us an estimate of misclassification error for a new observation. Usually in cross-validation for classification we try to **balance the folds**.