

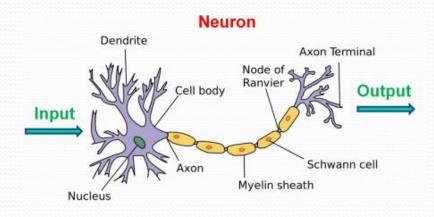
ESS2222

Lecture 3 – Bias-Variance Trade-off

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Bias-Variance Trade-off
 Overfitting & Regularization
 Ridge & Lasso Regression
 Nonlinear Transformation
 Cross-Validation
 Support Vector Machine



Outline

Review of Lecture 2

What does ν say about μ ?

For a sample of size N, ν is probably close to μ (within ϵ). P $[|\nu - \mu| > \epsilon] \le 2e^{-2\epsilon^2 N}$ Hoeffding's inequality P $[|E_{in}(g) - E_{out}(g)| > \epsilon] \le 2Me^{-2\epsilon^2 N}$ (Generalized)



Stochastic gradient decent (SGD) method, $w = w + \Delta w$ $\Delta w = -\eta \Delta J(w)$ $\Delta w j = -\eta \frac{\partial J}{\partial w_j} = \eta (y^i - \phi(z^i)) x^i_j$ Based on random sample.

a) Error is noisier , b) Convergence faster, c) Local minima can be escaped faster

Scaling is important for optimal performance,

Trade-off Between Approximation and Generalization:

A common issue in machine learning is overfitting, which occurs when the model is complex or the model not only captures the signal but also the noise in a dataset. In order to generalize the model to out-of-sample we have to avoid overfitting.

This is the difference between approximation (fitting) and learning (generalization).

Regularization:

Regularization is a powerful method for reducing overfitting. A good bias-variance trade-off can be obtained by tuning the complexity of the model via regularization. This can be achieved by introducing a penalty (bias) term for model complexity.

Trade-off Between Approximation and Generalization:

Our goal: Small $\mathbf{E}_{out} \rightarrow$ good approx. of *f* out of sample

However:

More complex $H \rightarrow$ better chance of approximating f in sample Less complex $H \rightarrow$ better chance of generalizing f out of sample

The best hypothesis is within the hypothesis set H. But the only way to navigate through this set to find the good candidate (g) is via the samples (to find the performance of one hypothesis versus another).

$$E_{out}(g^D) = E_X [(g^D(x) - f(x))^2]$$
$$E_D [E_{out}(g^D)] = ?$$

It can be shown that:

$$E_D[E_{out}(g^D)] = E_X\left[E_D\left[\left[(g^D(x) - \overline{g}(x))^2\right]\right]\right] + E_X\left[\left[(\overline{g}(x) - f(x))^2\right]\right]$$
var bias

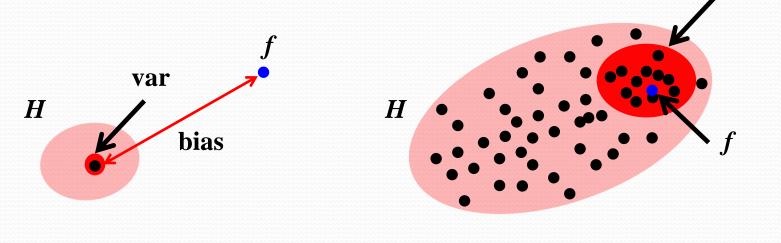
Error

where
$$\overline{g}(\mathbf{x}) = E_D[g^D(\mathbf{x})] \approx \frac{1}{k} \sum_k g^{D_k}(\mathbf{x})$$
 D: data

: Expected value over D (different data)

The Trade-off

 $bias = E_X \left[\left[(\overline{g}(x) - f(x))^2 \right] \right]$ $variance = E_X \left[E_D \left[\left[(g^D(x) - \overline{g}(x))^2 \right] \right] \right]$



small H

large H (and complex)

var

So as we go from small H to large H:



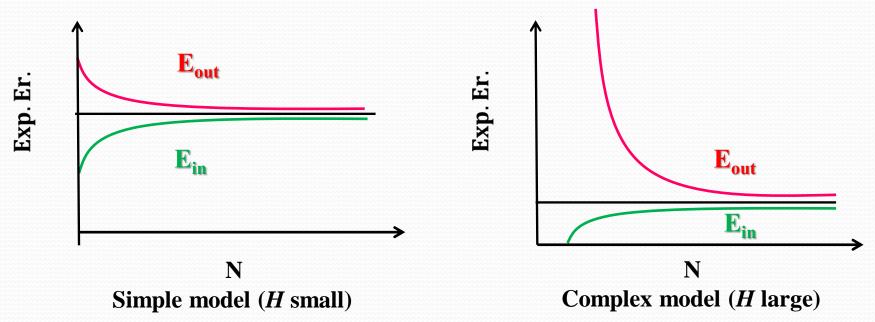
Expected E_{out} and E_{in}

Consider data set D of size N:

 $E_{D}[E_{out}(gD)]$ Expected out-of-sample error

 $\mathbf{E}_{\mathbf{D}}[\mathbf{E}_{in}(gD)]$ Expected in-sample error

How do they vary with N?



Bias-Variance Trade-off Example

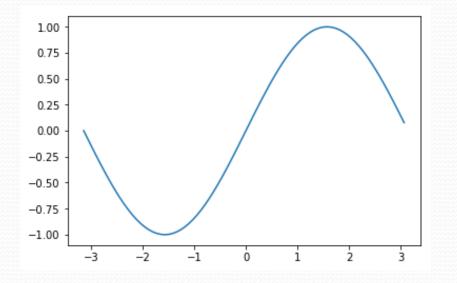
 $\mathbf{F}(\mathbf{x}) = \mathbf{Sin}(2\pi\mathbf{x})$

Suppose we have two samples! N = 2We try two models (hypotheses):

H₀: h(x) = b simple **H**₁: h(x) = ax + b complex

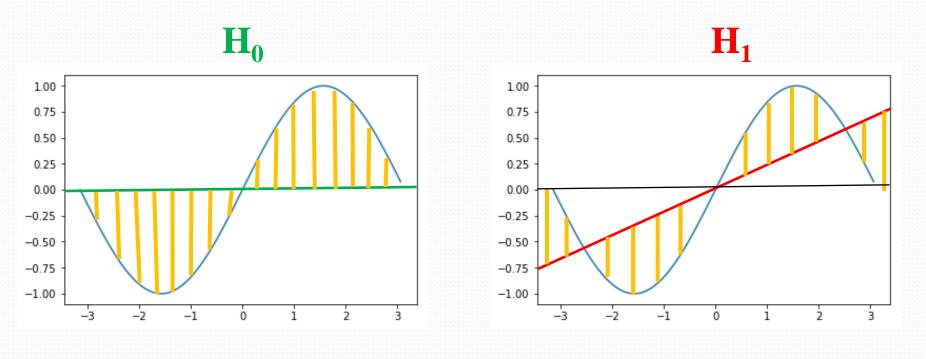
Which one is better, H_0 or H_1 ?

Better in what?

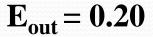


Approximation - H₀ vs H₁

$F(x) = Sin(2\pi x)$ Approximate F(x) with H_0 and H_1



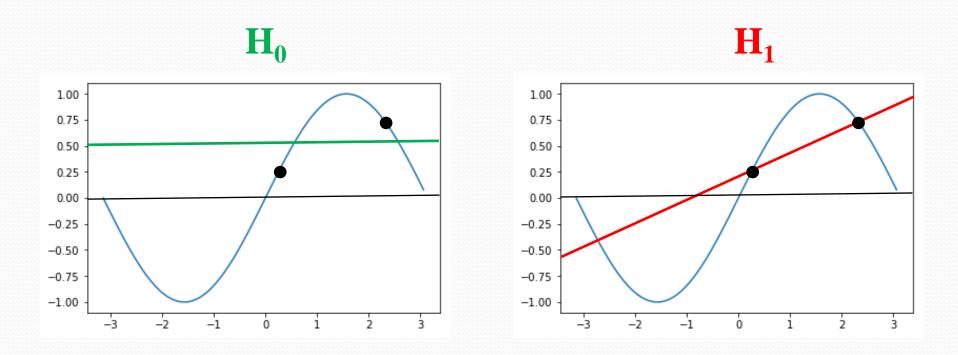
 $E_{out} = 0.5$



Error

Learning - H₀ vs H₁

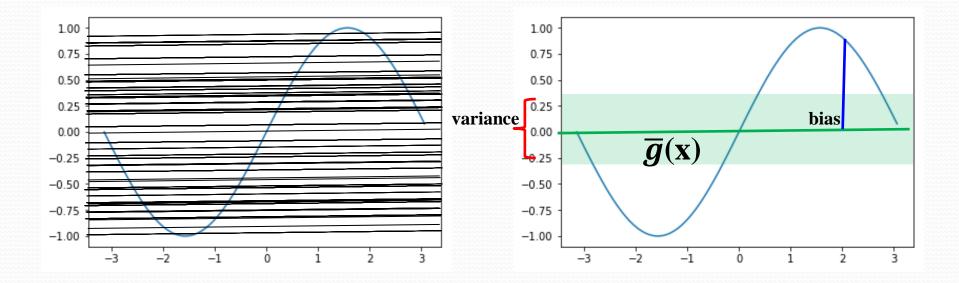
 $\mathbf{F}(\mathbf{x}) = \mathbf{Sin}(2\pi\mathbf{x})$



Final hypothesis using two samples

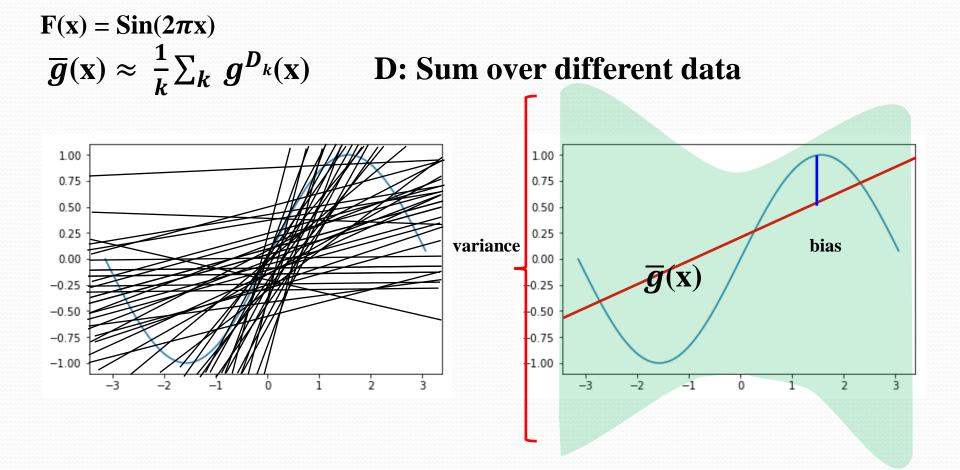
Bias and Variance - H₀

 $F(\mathbf{x}) = \operatorname{Sin}(2\pi \mathbf{x})$ $\overline{g}(\mathbf{x}) \approx \frac{1}{k} \sum_{k} g^{D_{k}}(\mathbf{x})$ D: Sum over different data



Note that $\overline{g}(x)$ is not the output of learning. The output of learning is one of these lines based on given samples.

Bias and Variance - H₁



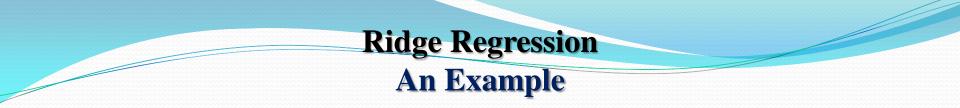
 $\mathbf{F}(\mathbf{x}) = \mathbf{Sin}(2\pi\mathbf{x})$ H_1 H₀ 1.00 1.00 0.75 0.75 0.50 0.50 0.25 0.25 $\overline{g}(\mathbf{x})$ $\overline{g}(\mathbf{x})$ 0.00 0.00 -0.25 -0.25 -0.50 -0.50 -0.75 -0.75 -1.00-1.00-3 -3 -2 -2 -1 -1 ò i ż ż. ò ż 1

H₀ vs H₁

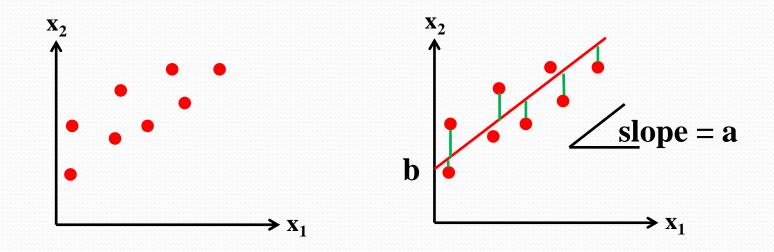
bias = 0.5 var = 0.25

bias = 0.21 var = 1.7

3



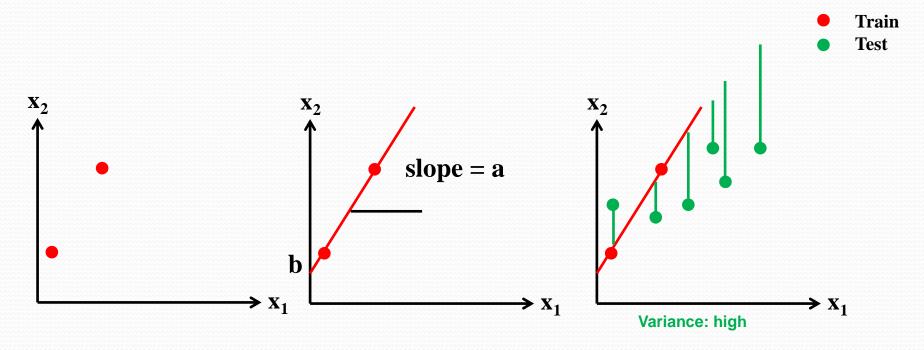
Suppose we want to train a model using a number of the training samples.



 $\mathbf{x}_2 = \mathbf{a} \ \mathbf{x}_1 + \mathbf{b} \quad (\mathbf{a} \equiv \mathbf{w}_1, \mathbf{b} \equiv \mathbf{w}_0)$

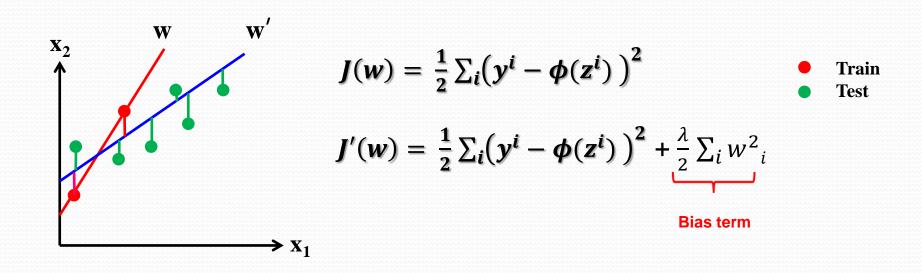
Ridge Regression An Example

Now suppose we have only two samples for training. Chose two of them as training samples and the rest as test samples.



High variance: The learned model overfits to the training data.

Overfitting & Regularization Ridge Regression



The idea behind the ridge regression is that to find a new line which doesn't fit to the training data very tightly.

Adding a small bias can highly decrease the variance.

Slightly worse fit, better generalization.

Ridge Regression An Example

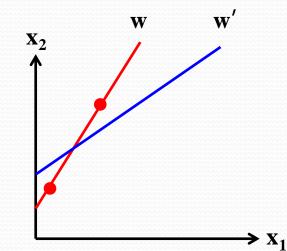
- Suppose $\Phi(z) = z$ Adeline
- $z = x.w = \sum_{i=0}^{n} w_i x_i$

then: $\Phi(z) = x \cdot w = \sum_{i=1}^{n} w_{i} x_{i}$ activation func.

 $\Phi(\mathbf{z}) = w_1 x_1 + w_0$

$$J(w) = \frac{1}{2} \sum_{i} (y^{i} - \phi(z^{i}))^{2}$$

$$J'(w) = \frac{1}{2} \sum_{i} (y^{i} - \phi(z^{i}))^{2} + \frac{\lambda}{2} \sum_{i} w^{2}_{i}$$



$$J(w) = \frac{1}{2} \sum_{i} (y^{i} - \phi(z^{i}))^{2} = J(w) = \frac{1}{2} [l_{x1} - w_{1} x_{1} - w_{0}]^{2} + [l_{x2} - w_{1} x_{2} - w_{0}]^{2}$$

$$J'(w) = \frac{1}{2} \sum_{i} (y^{i} - \phi(z^{i}))^{2} = J(w) = \frac{1}{2} [l_{x1} - w_{1} x_{1} - w_{0}]^{2} + [l_{x2} - w_{1} x_{2} - w_{0}]^{2}$$

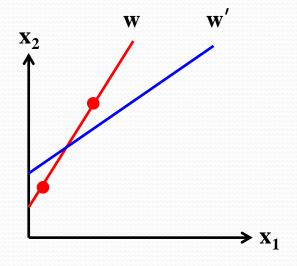
$$+ \frac{\lambda}{2} [w^{2}_{0} + w^{2}_{1}]$$

l_{xi} : label

 $\frac{\partial J'}{\partial w_0}, \frac{\partial J'}{\partial w_1}$

Ridge Regression An Example

- Note that our goal is minimizing the cost function J
- With ridge regression penalty:
- $J(w) = \frac{1}{2} \sum_{i} (y^{i} \phi(z^{i}))^{2}$ $J'(w) = \frac{1}{2} \sum_{i} (y^{i} - \phi(z^{i}))^{2} + \frac{\lambda}{2}$ $J'(w)_{\min} < J'(w)_{\min} \text{ for blue line}$
- So we chose ridge regression line over least squares line.

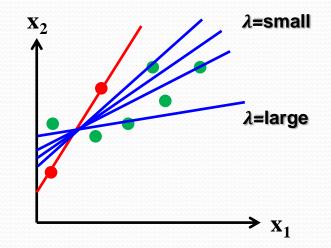


Ridge Regression λ -parameter

 $0 \le \lambda < \infty$

How to chose lambda?

We try different values for λ and use cross-Validation, typically 10-fold cross-validation to determine which one results in the lowest variance.



Cross-Validation

16%	16%	16%	16%	16%	20%
Training data	Training data	Training data	Training data	Validation	Testing data
Training data	Training data	Training data	Validation	Training data	Testing data
Training data	Training data	Validation	Training data	Training data	Testing data
Training data	Validation	Training data	Training data	Training data	Testing data
Validation	Training data	Training data	Training data	Training data	Testing data

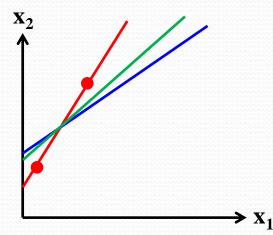
5-fold cross-validation

Overfitting & Regularization Lasso Regression

Lasso regression is similar to the ridge regression a except for an important difference!

- L2-rgression (Ridge) $J'_{r}(w) = \frac{1}{2} \sum_{i} (y^{i} - \phi(z^{i}))^{2} + \frac{\lambda}{2} \sum_{i} w^{2}_{i}$
- L1-rgression (Lasso) $J'_{l}(w) = \frac{1}{2} \sum_{i} (y^{i} - \phi(z^{i}))^{2} + \frac{\lambda}{2} \sum_{i} |w_{i}|$

In lasso regression the weight factors of less important (relevant) features shrink faster which is good.

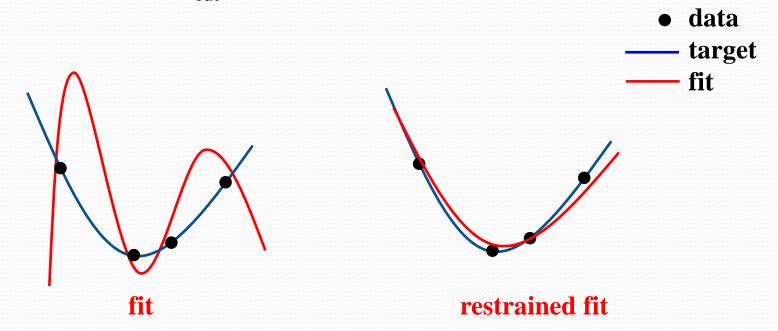


When we train a model, we are matching the model complexity to the data resources, not the target complexity (which should be avoided).

Lesson

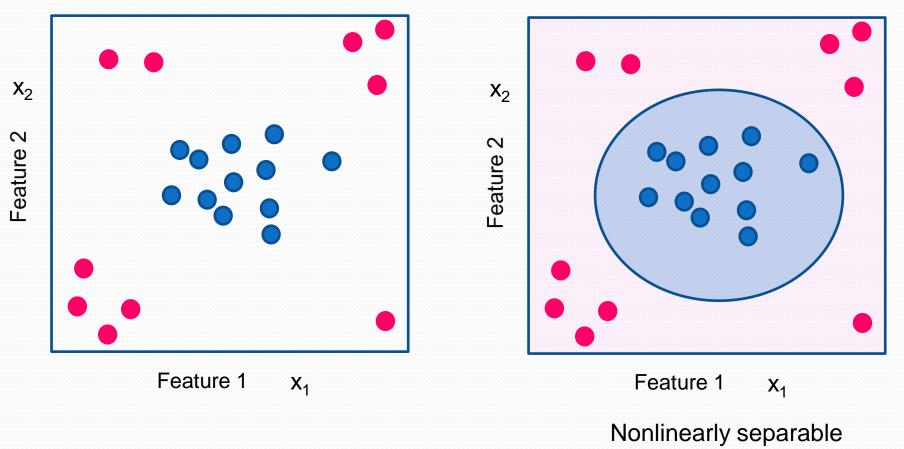
When the number of sample data is not large, we have to avoid training complex models.

Allowing $E_{in} \rightarrow E_{in} + \in$ (e.g., $\in \sim 1\% E_{in}$) can highly improve the out-of-sample performance (E_{out} and variance).



Can we apply linear leaning algorithms for nonlinear problems? Note that feature is a higher level representation of raw input.

Nonlinearity





Can we apply linear leaning algorithms to nonlinear problems?

 $h(x) = \sum_{i=0}^{m} w_i x_i$ is linear in both w and x.

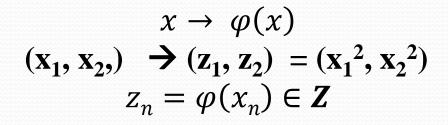
Being linear in w is important because the algorithm works because of linearity in the weights.

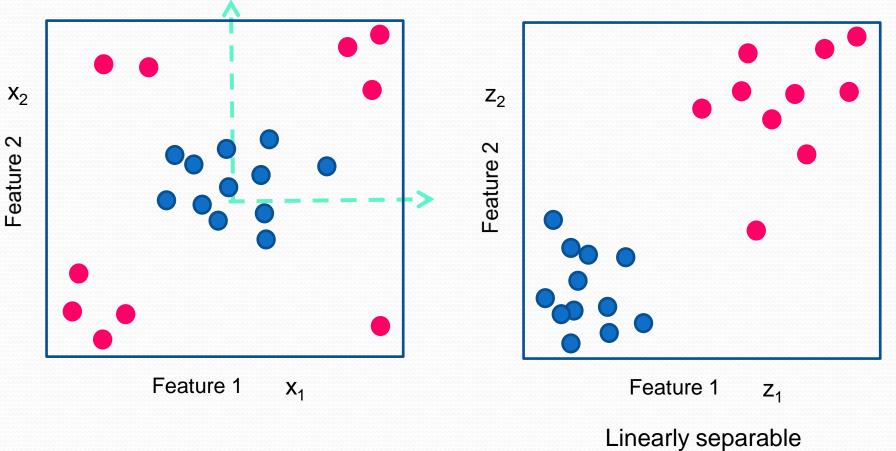
We still can use linear algorithm for nonlinear problems.



Nonlinearity

 $x_n \in \boldsymbol{X}$





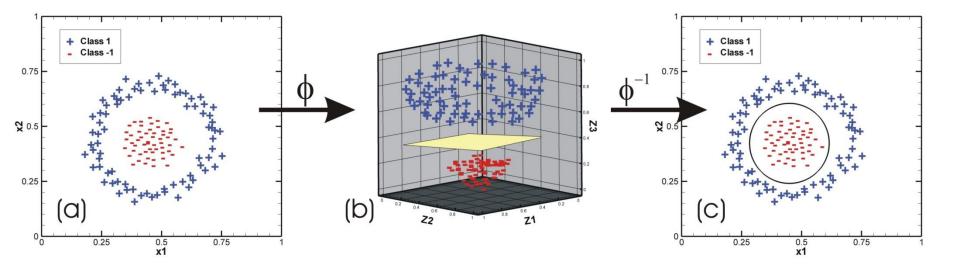
26

Separate in Z-space Classify in X-space $g(z) = sign(\widetilde{w}^T z) =$ $g(x) = \tilde{g}(\varphi(x)) = \operatorname{sign}(w^T \varphi(x)) \quad x \leftarrow \varphi^{-1}(x)$ sign($\tilde{w}^T \varphi(x)$)) **X**₂ Z_2 Feature 2 2 Feature 3 Feature 1 Feature 1 **X**₁ Z_1

Nonlinearity

Mapping onto a Higher Dimensional Feature Space

Nonlinearity



 $\varphi \colon R^2 \to R^3 \qquad x \to \varphi(x) \qquad \mathbf{z} = \mathbf{x} \cdot \mathbf{w} \to \mathbf{z} = \varphi(x) \cdot \mathbf{w}$

 $\varphi(x_1, x_2) = (z_1, z_2, z_3) = (x_1, x_2, x_1^2 + x_2^2).$

$$z = \varphi(x).w$$

$$w = \sum_{i=1}^{m} \alpha_i \varphi(x_i)$$
 Assumption

$$z = \sum_{i=1}^m \alpha_i \varphi(x_i) \cdot \varphi(x)$$

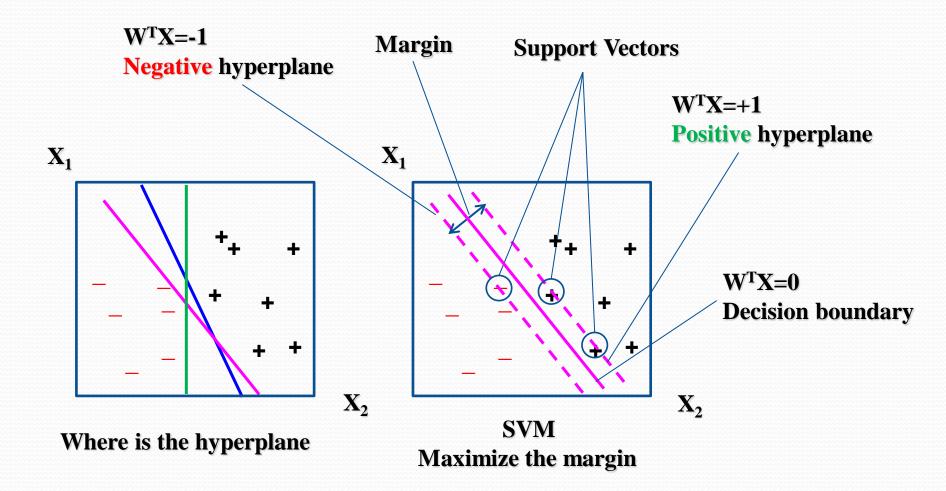
$$K(x_i, x) = \varphi(x_i) \cdot \varphi(x)$$

$$\mathbf{z} = \sum_{i=1}^{m} \alpha_i K(x_i, x)$$

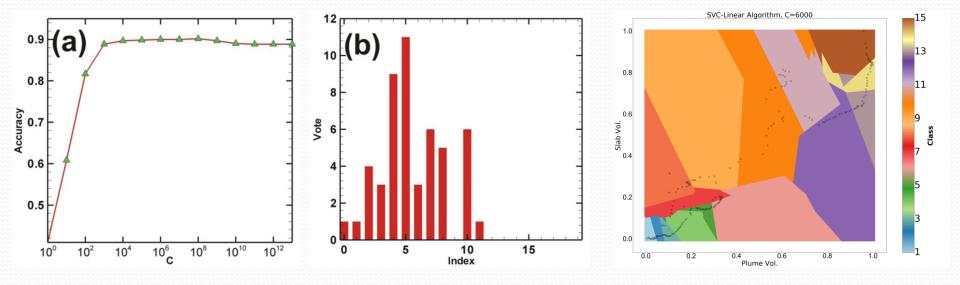
 $K(x_i, x) = (x_i, x)$ Linear $K(x_i, x) = (\gamma(x_i, x) + r)^d$ Polynomial $K(x_i, x) = Exp(-\gamma ||x_i, -x||^2)$ RBF $K(x_i, x) = tanh(\gamma(x_i, x) + r)$ Sgmiod

$$k(x^{i}, x^{j}) = \exp\left(-\frac{\|x^{i} - x^{j}\|^{2}}{2}\right) = \exp\left(-\frac{1}{2} < x^{i} - x^{j}, x^{i} - x^{j} > \right) = \exp\left(-\frac{1}{2}\|x^{i}\| - \frac{1}{2}\|x^{j}\|\right) \exp\left(\right) = C\sum_{k=0}^{\infty} \frac{^{k}}{k!},$$

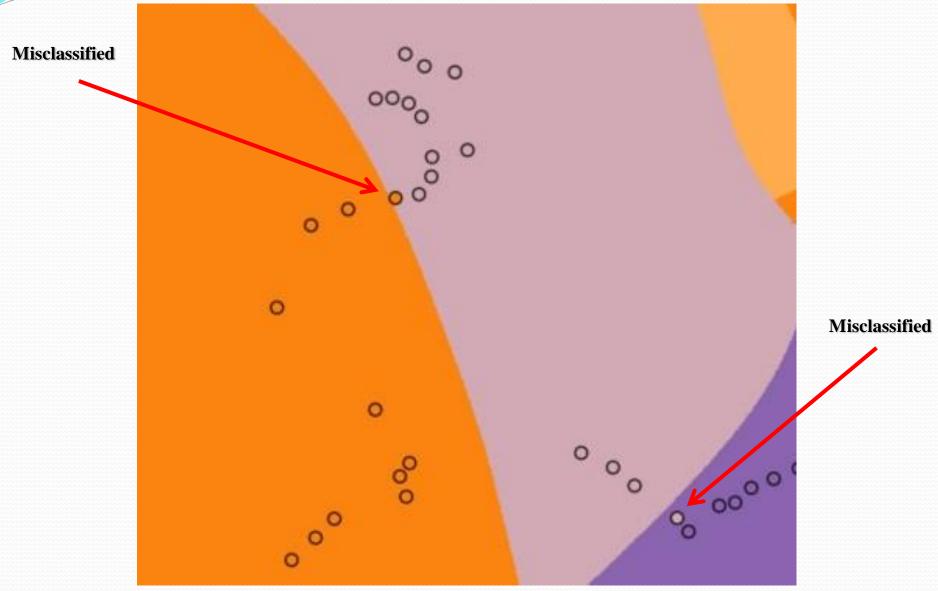
Support Vector Machine (SVM)



Assignment 1 - SVC-Linear



Assignment 1 - SVC-Linear



Coarse Grid Search – Related to Assignment 1

Cross validator

My_cv = StratifiedShuffleSplit(n_splits=10, test_size=0.2, train_size = None, random_state=19)

StratifiedShuffleSplit: Stratified ShuffleSplit cross-validator Stratification is the process of rearranging the data as to ensure each fold is a good representative of the whole (each fold comprises around the same fraction of classes).

My_cv: user defined cross-validator n_splits: number of splits (default = 10) test_size: default = 0.1 if train_size is unspecified, otherwise it will complement the specified train_size. It should be specified if train_size = None. random_state: seed for random shuffling

Grid search

grid = GridSearchCV(SVC(kernel='linear'), param_grid=grid_parameters, cv=My_cv, return_train_score=False)

GridSearchCV implements a "fit" and a "score" method. It also implements "predict".

SVC(kernel='linear'): estimator param_grid: grid parameters cv: cross validator return_train_score: if False, the cv_results_attribute will not include training scores

grid.fit(feature_data, class_labels)

Fine Grid Search – Related to Assignment 1

Cross validator

My_cv = KFold(n_splits=splits_num, shuffle=True, random_state=i)

My_svm = svm.SVC(kernel='linear') # linear kernel

Grid search

grid = GridSearchCV(estimator=My_svm, param_grid=p_grid, cv=My_cv, return_train_score = False)

My_svm = svm.SVC(kernel='linear')

grid.fit(feature_data, class_labels)

grid.best_score_ grid.best_index_

Learning Model – Related to Assignment 1

Cross validator

X = feature_train y = class_labels_train my_C = 6000.0 # SVM regularization parameter svm_SVC_lin = svm.SVC(kernel='linear', C=my_C) svm_SVC_lin.fit(X, y)

grid.best_score_ grid.best_index_