

## Scikit-learn and Tour of Classifiers

**Support Vector Machines** 

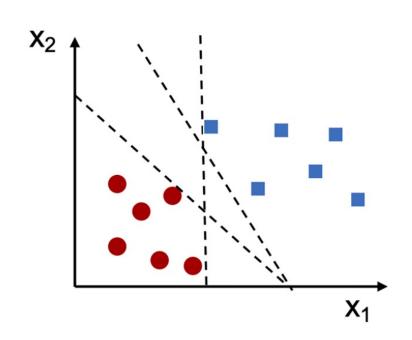


## **Support Vector Machines**

- Widely-used, powerful classifiers
- Can be considered an extension of the Perceptron
- Support vector classification (SCV) objective: maximize margin
- Margin: the distance between the separating hyperplane (decision boundary) and the training samples that are closest to this hyperplane
- The training samples closest to the hyperplane are called support vectors
- Kernel-SVM for nonlinear decision boundaries, resulting predictor is sparse

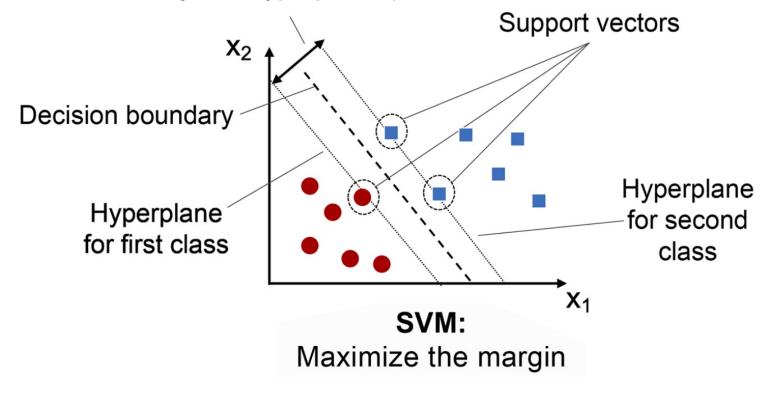


### Support Vector Machines



Which hyperplane?

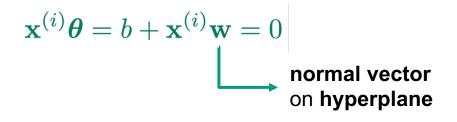
Margin (gap between decision boundary and hyperplanes)

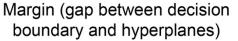


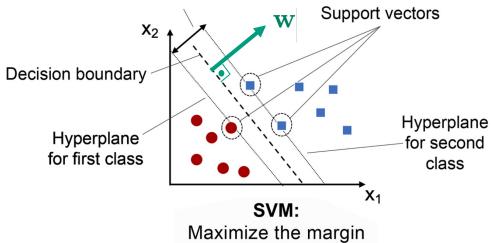


## Maximum margin classifier

- Decision boundaries with large margins tend to have a lower generalization error
- Decision boundaries with small margins are more prone to overfitting
- Linear decision boundary:







$$\mathbf{x}^{(i)} = \begin{bmatrix} 1 & x_1^{(i)} & x_2^{(i)} & \cdots & x_m^{(i)} \end{bmatrix} \qquad \boldsymbol{\theta} = \begin{bmatrix} \theta_0 \\ \theta_1 \\ \vdots \\ \theta_m \end{bmatrix} := \begin{bmatrix} b \\ \mathbf{w} \end{bmatrix} \qquad \mathbf{w} = \begin{bmatrix} \theta_1 \\ \theta_2 \\ \vdots \\ \theta_m \end{bmatrix}, \quad b = \theta_0$$

- Distance between positive and negative hyperplanes: 2 / ||w||
- Maximize distance while classifying samples correctly



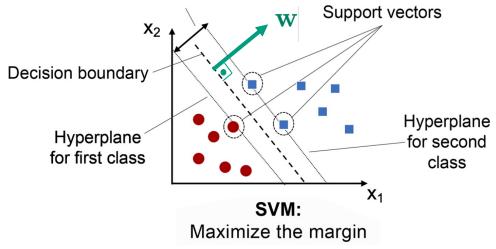
## Maximum margin (linearly separable) Margin (gap between decision boundary and hyperplanes)

- Derivation of maximum margin formula not syllabus; see e.g.
   <a href="https://www.cs.cornell.edu/courses/cs4780/2018fa/lectures/lecturenote09.html">https://www.cs.cornell.edu/courses/cs4780/2018fa/lectures/lecturenote09.html</a>
- Finding the **maximum margin** hyperplanes can be cast into the **quadratic optimization problem**:

$$\min_{\mathbf{w},b} \mathbf{w}^T \mathbf{w}$$
 such that for all  $i$   $y^{(i)} \left( \mathbf{x}^{(i)} \mathbf{w} + b \right) \geq 1$ 

• For support vectors (data on the hyperplanes), it holds

$$y^{(i)}\left(\mathbf{x}^{(i)}\mathbf{w} + b\right) = 1$$



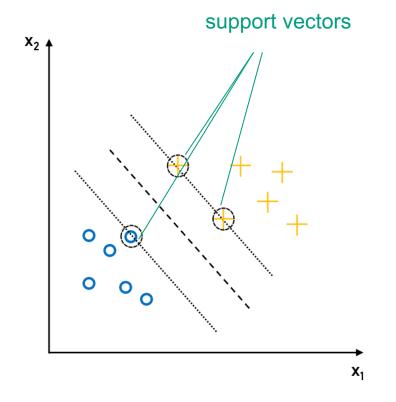
Classes:  $y^{(i)} \in \{-1,1\}$ 

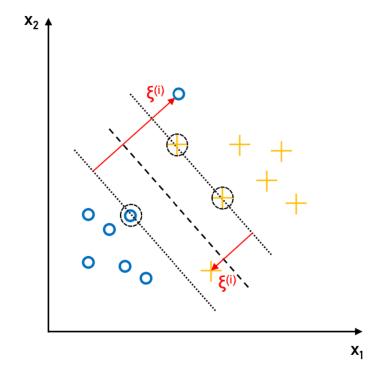
Recall that: 
$$\mathbf{w}^T\mathbf{w} = ||\mathbf{w}||_2^2 = \sum_{i=1}^m w_i^2$$



## Maximum margin – soft margin

• Often, data is not linearly separable. Idea: soft margin by introducing "slack variables"  $\xi^{(i)}$ 







## Maximum margin – soft margin

- Often, data is not linearly separable. Idea: soft margin by introducing "slack variables"  $\xi^{(i)}$
- Modified optimization problem:

$$\min_{\mathbf{w},b} \mathbf{w}^T \mathbf{w} + C \sum_{i=1}^n \xi^{(i)} \quad \text{ such that for all } i \quad y^{(i)} \left( \mathbf{x}^{(i)} \mathbf{w} + b \right) \ge 1 - \xi^{(i)} \quad \text{and} \quad \xi^{(i)} \ge 0$$

- The slack variables allow for some samples to be within the margin (or even on the wrong side = misclassified)
- For high C we penalize sample in or on the wrong side of the margin strongly
- The optimization problem with slack variables can be written in terms of the loss function:

$$L(\boldsymbol{\theta}) = L(\mathbf{w}, b) = \underbrace{\mathbf{w}^T \mathbf{w}}_{\ell_2 - \text{regul.}} + C \sum_{i=1}^n \underbrace{\max \left\{ 1 - y^{(i)} \left( \mathbf{x}^{(i)} \mathbf{w} + b \right), 0 \right\}}_{\text{hinge-loss}}$$

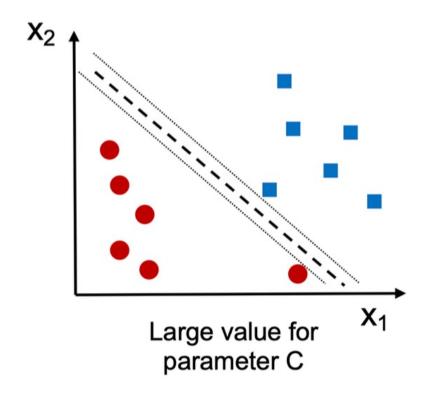
Recall that:

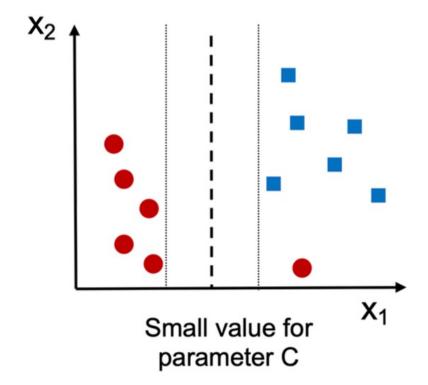
$$\mathbf{w}^T \mathbf{w} \equiv ||\mathbf{w}||_2^2 = \sum_{i=1}^m w_i^2$$



## Maximum margin – soft margin

• Effect of C: decreasing C increases the bias (underfitting) and lowers the variance (overfitting) of the model

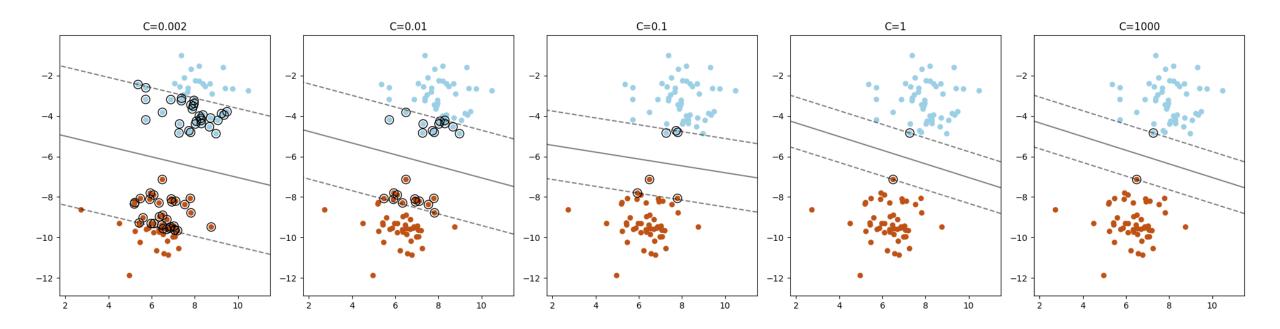






## SVM code example

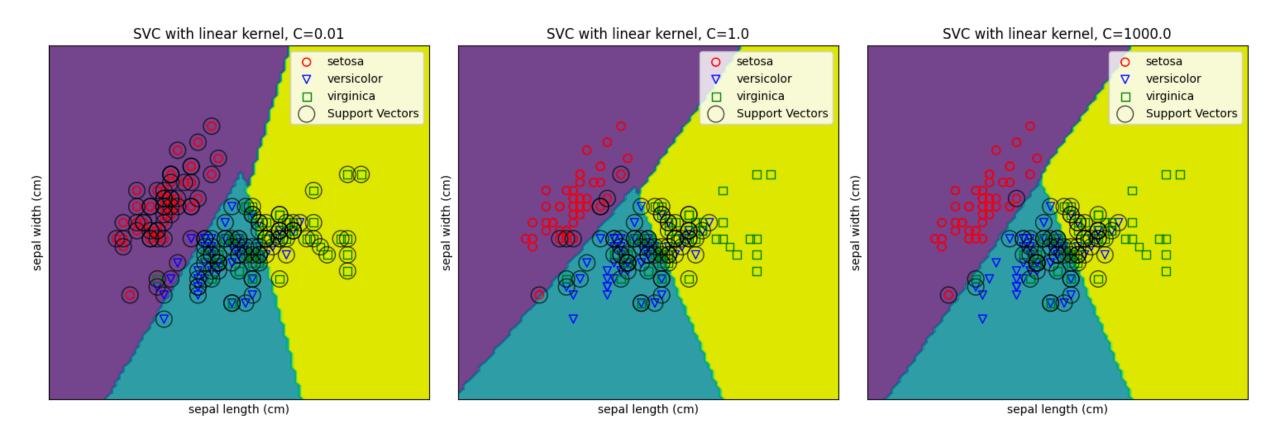
03\_svm\_linear\_blobs.ipynb





## SVM code example

03\_svm\_linear\_iris.ipynb





## Logistic regression versus Linear SVM

- LR and linear SVM often yield very similar results
- Logistic regression tries to maximize the conditional likelihoods for all of the training data, which makes it more prone to outliers than SVMs
- SVMs mostly care about the samples that are closest to the decision boundary (support vectors)
- Logistic regression has the advantage that it is a simpler model and can be implemented more easily



## Linear model (linear decision boundaries)

- Linear model classifiers may appear to be very restrictive in lowdimensional spaces (very few features in X) because of their straight line or plane boundaries
- For high dimensional data (many features in X) linear model classifiers may act as a guard against overfitting



## Scikit-learn implementation

- Perceptron, LogisticRegression and LinearSVC classes in scikit-learn
   make use of the LIBLINEAR library (highly optimized C/C++ library developed at the National Taiwan University)
- SVC and SVM classes in scikit-learn
   makes use of LIBSVM library (an equivalent C/C++ library specialized for SVMs developed at the National Taiwan University)
- LIBLINEAR and LIBSVM are faster than native Python implementations



## Scikit-learn implementation

- Perceptron, LogisticRegression and LinearSVC classes in scikit-learn
   make use of the LIBLINEAR library (highly optimized C/C++ library developed at the National Taiwan University)
- SVC and SVM classes in scikit-learn
   makes use of LIBSVM library (an equivalent C/C++ library specialized for SVMs developed at the National Taiwan University)
- LIBLINEAR and LIBSVM are faster than native Python implementations



## Scikit-learn alternative implementation

- When datasets are too large to fit into computer memory
  - Alternative implementations using the SGDClassifier class
  - SGDClassifier class supports online learning via the partial\_fit method
  - Concept behind the SGDClassifier class is the stochastic gradient descend algorithm
- Initialize the stochastic gradient descent version of the perceptron (ppn), logistic regression
  (lr), and a support vector machine (svm) with default parameters as follows

```
from sklearn.linear_model import SGDClassifier
ppn = SGDClassifier(loss='perceptron')
lr = SGDClassifier(loss='log')
svm = SGDClassifier(loss='hinge')
```

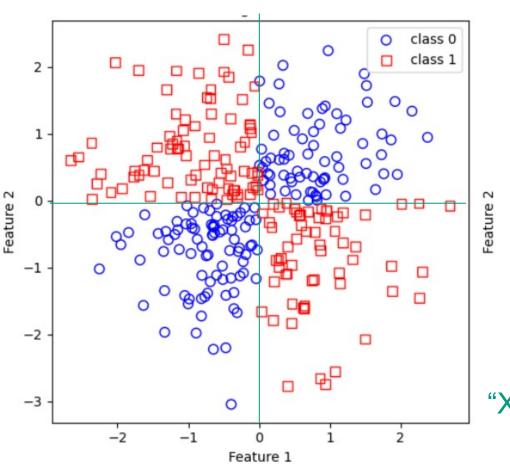


## Nonlinear decision boundaries

Kernel Support Vector Machines



### Nonlinear decision boundaries – Motivation

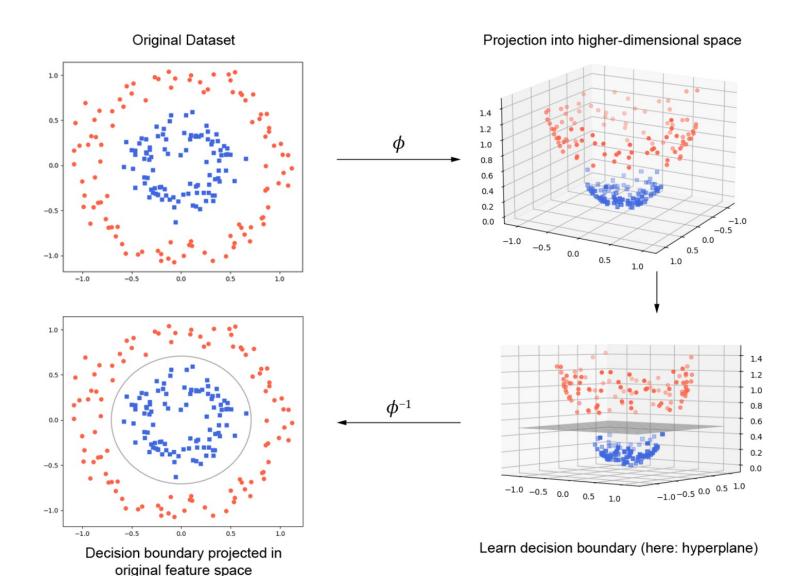


Data not linearly separable!

"XOR" dataset

### Idea behind kernel methods



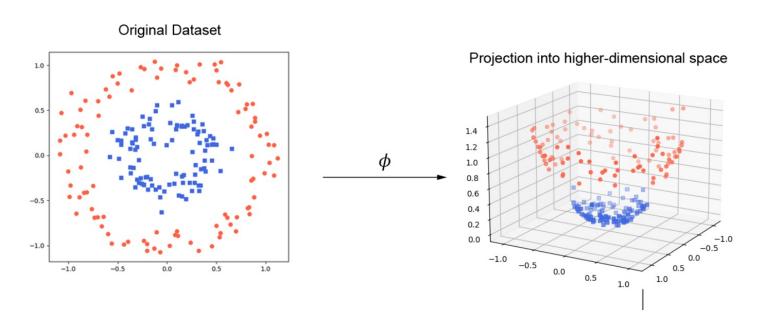


 $\phi$  feature mapping

Data may be linearly separable in a higher dimensional feature space

#### Idea behind kernel methods





 $\phi$  feature mapping, here:

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

### Kernel trick

sample: 
$$\mathbf{x}^{(i)} = \begin{bmatrix} 1 & x_1^{(i)} & x_2^{(i)} & \cdots & x_m^{(i)} \end{bmatrix}$$



Many algorithms can be formulated in terms of inner products of the samples

$$\mathbf{x}^{(i)}\mathbf{x}^{(j)^T} = \sum_{k=0}^{m} x_k^{(i)} x_k^{(j)}$$

- After feature mapping, we have an inner product in a high-dimensional space  $\phi(\mathbf{x}^{(i)})\phi(\mathbf{x}^{(j)})^T$
- Often, there is an easier way to compute the resulting number in the original space

$$\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \phi(\mathbf{x}^{(i)})\phi(\mathbf{x}^{(j)})^T$$

Kernel:  $\kappa(\mathbf{x}, \mathbf{x}') : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ 

### without ever computing $\phi(\mathbf{x}^{(i)})$

•  $\phi(\mathbf{x}^{(i)})$  is never needed. There is even kernels for which no explicit finite dimensional feature mapping exists

#### Kernel trick



Example: A quadratic polynomial kernel, m=3 (number of features)

$$\begin{split} \kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) &= \left(\mathbf{x}^{(i)}^T \mathbf{x}^{(j)}\right)^2 \\ &= \left(\sum_{k=1}^m x_k^{(i)} x_k^{(j)}\right) \left(\sum_{k=1}^m x_k^{(i)} x_k^{(j)}\right) \\ &= \sum_{k=1}^m \sum_{l=1}^m x_k^{(i)} x_l^{(i)} x_k^{(j)} x_l^{(j)} \\ &= \sum_{k=1}^m \sum_{l=1}^m (x_k^{(i)} x_l^{(i)}) (x_k^{(j)} x_l^{(j)}) \end{split}$$

Computation in original space via kernel: O(m)

Therefore, corresponding feature mapping is:

$$\begin{split} \phi(\mathbf{x}^{(i)}) &= \left[ x_1^{(i)} x_1^{(i)}, \quad x_1^{(i)} x_2^{(i)}, \quad x_1^{(i)} x_3^{(i)}, \quad x_2^{(i)} x_1^{(i)}, \quad x_2^{(i)} x_2^{(i)}, \quad x_2^{(i)} x_3^{(i)}, \quad x_3^{(i)} x_1^{(i)}, \quad x_3^{(i)} x_2^{(i)}, \quad x_3^{(i)} x_3^{(i)} \right] \\ \phi(\mathbf{x}^{(i)}) \phi(\mathbf{x}^{(i)})^T &= \sum_{p=1}^3 \phi(\mathbf{x}^{(i)})_p \phi(\mathbf{x}^{(j)})_p = \sum_{k=1}^{m=3} \sum_{l=1}^{m=3} (x_k^{(i)} x_l^{(i)}) (x_k^{(j)} x_l^{(j)}) & \text{Computation in feature} \\ &\text{space: O(m}^2) \end{split}$$

#### Kernel trick



- If the data only occurs in form of inner products in an ML algorithm, we can replace these by evaluations of a kernel function to compute inner products in a high-dimensional feature space
- Evaluation with data in the original space is cheaper
- Evaluation in the feature space is expensive
- We might not even know what the feature mapping looks like!



#### Kernel functions

- Kernel can be seen to measure similarity between samples
- "Valid" kernel (Mercer kernel): if kernel matrix is positive semi-definite

$$K_{ij} = \kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \begin{bmatrix} \kappa(\mathbf{x}^{(1)}, \mathbf{x}^{(1)}) & \cdots & \kappa(\mathbf{x}^{(1)}, \mathbf{x}^{(n)}) \\ \vdots & \ddots & \vdots \\ \kappa(\mathbf{x}^{(n)}, \mathbf{x}^{(1)}) & \cdots & \kappa(\mathbf{x}^{(n)}, \mathbf{x}^{(n)}) \end{bmatrix} \qquad \kappa(\mathbf{x}, \mathbf{x}') : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$$

a matrix **A** is positive semi-definite if  $\mathbf{v}^T \mathbf{A} \mathbf{v} \geq 0$  for all non-zero  $\mathbf{v}$ 

 For valid kernels, it is guaranteed that a corresponding feature mapping exists (although, it might be infinite dimensional)



## Typical kernel functions

Linear kernel

$$\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \boldsymbol{\phi}(\mathbf{x}^{(i)}) \boldsymbol{\phi}(\mathbf{x}^{(j)})^T = \mathbf{x}^{(i)} \mathbf{x}^{(j)}^T$$

Polynomial kernel (tuning parameter p)

$$\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = (\mathbf{x}^{(i)}\mathbf{x}^{(j)}^T + 1)^p$$
, for polynomial degree  $p$ 

• Isotropic Gaussian kernel / Radial basis function (RBF) kernel (tuning parameter  $\gamma$ )

$$\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \exp\left(-\frac{||\mathbf{x}^{(i)} - \mathbf{x}^{(j)}||_2^2}{2\sigma^2}\right) := \exp\left(-\gamma||\mathbf{x}^{(i)} - \mathbf{x}^{(j)}||_2^2\right)$$



## More kernel functions (not syllabus)

- There is also kernel functions for strings, text, etc.
- Kernelized algorithm can then operator directly on that data without explicit transformation
- Choice of kernel function should be tailored to the data



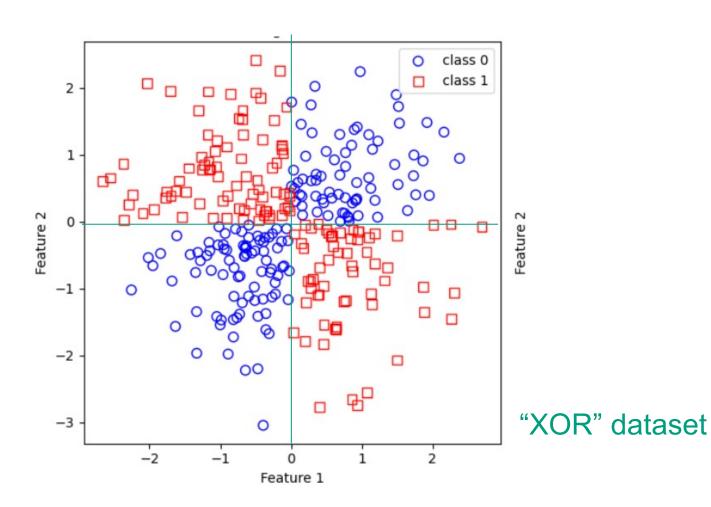
#### Kernel SVM

In prediction and optimization algorithm (we didn't look at it in detail) data only occurs
as inner products → replace inner products by kernel function calls

- In scikit-learn the type of kernel is a parameter of the SVC (Support Vector Classifier)
- The kernels have hyperparameters (e.g. degree for polynomial kernel)

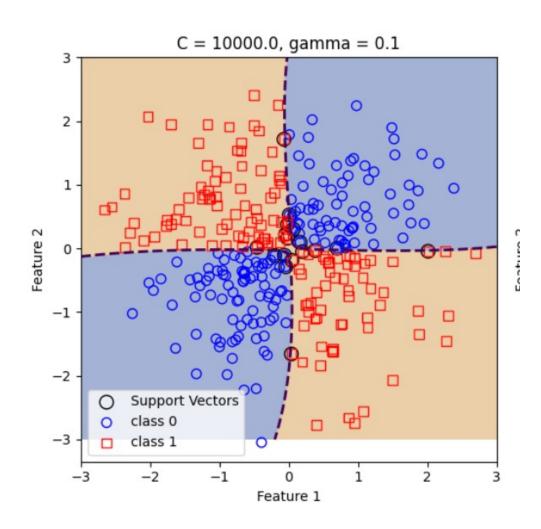


## Kernel SVM Code example





## Kernel SVM Code example (RBF kernel)



#### Chosen kernel function:

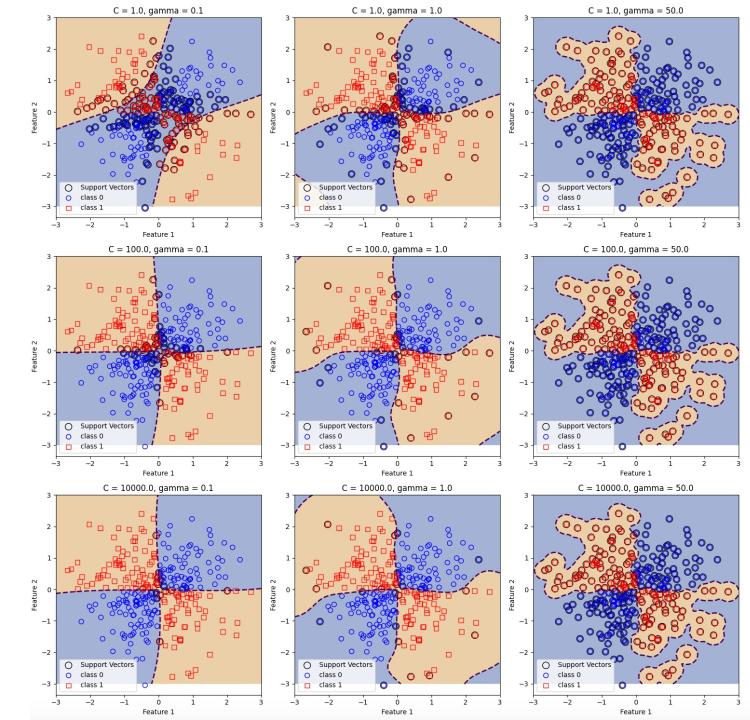
$$\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) := \exp\left(-\gamma ||\mathbf{x}^{(i)} - \mathbf{x}^{(j)}||_2^2\right)$$

### Kernel SVM

- We need to tune both C and the kernel parameter(s)
- For example: grid search
  - C in [1.0, 100.0, 10000.0]
  - $\gamma$  in [0.1, 1.0, 50.0]

03\_svm\_kernel\_xor.ipynb

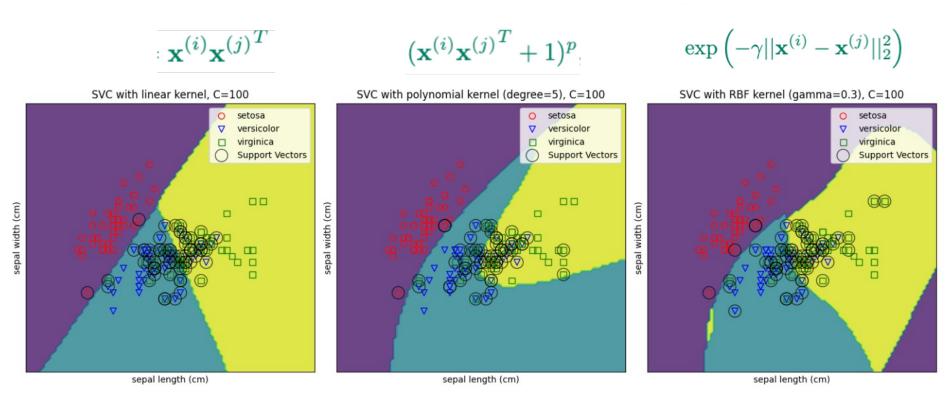
03\_svm\_gridsearch\_iris.ipynb





#### Kernel SVM

- We can also try different kernels
- For example:
  - Linear
  - Polynomial
  - RBF



03\_svm\_different\_kernels\_iris.ipynb



# Other kernelized ML algorithms

Nonlinear decision boundaries



## Kernelized ML algorithms

- Many of the presented algorithms can be kernelized by replacing inner products with kernel evaluations (sometimes after reformulation of the original algorithm)
- Examples:
  - Kernel Perceptron
  - Kernel Logistic Regression
  - Kernel Ridge Regression (linear regression + L2 regularizer)
  - Kernel SVM
  - Kernel k-nearest neighbors (next lecture)



# Nonlinear decision boundaries

Kernel perceptron

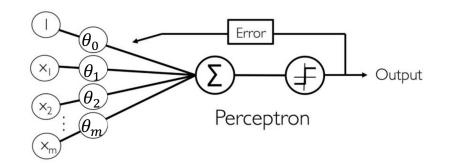


- To gain intuition of how to kernelize linear models
- Recall that for the perceptron  $y^{(i)} \in \{1, -1\}$ 
  - we predict as

$$\hat{y}^{(i)} = \operatorname{sign}\{\mathbf{x}^{(i)}\boldsymbol{\theta}\}$$

we train as

$$oldsymbol{ heta} \leftarrow \mathbf{0}$$
 for  $i=1,\cdots,n$  do  $\hat{y}^{(i)} \leftarrow \mathrm{sign}\{\mathbf{x}^{(i)}oldsymbol{ heta}\}$  if  $\hat{y} 
eq y^{(i)}$  then  $oldsymbol{ heta} \leftarrow oldsymbol{ heta} + \eta y^{(i)} \mathbf{x}^{(i)}$  end if end for



this actually means: 
$$\boldsymbol{\theta} = \sum_{j=1}^{n} \alpha^{(j)} \eta y^{(j)} \mathbf{x}^{(j)^T}$$

 $\alpha^{(j)}$ : number of times sample j is misclassified



- Parameters are linear combination of the samples:  $\theta = \sum_{i=1}^{\infty} \alpha^{(j)} \eta y^{(j)} \mathbf{x}^{(j)}$
- Recall that for the perceptron  $y^{(i)} \in \{1, -1\}$ 
  - we predict as

$$\hat{y}^{(i)} = \mathrm{sign}\{\mathbf{x}^{(i)}oldsymbol{ heta}\}$$

we train as

$$oldsymbol{ heta} \leftarrow \mathbf{0}$$
 for  $i=1,\cdots,n$  do  $\hat{y}^{(i)} \leftarrow \mathrm{sign}\{\mathbf{x}^{(i)}oldsymbol{ heta}\}$  if  $\hat{y} 
eq y^{(i)}$  then  $oldsymbol{ heta} \leftarrow oldsymbol{ heta} + \eta y^{(i)} \mathbf{x}^{(i)}^T$  end if end for

"dual" formulation

$$\hat{y}^{(i)} = \operatorname{sign} \left\{ \sum_{j=1}^{n} \alpha^{(j)} \eta y^{(j)} \mathbf{x}^{(i)} \mathbf{x}^{(j)^{T}} \right\}$$

$$\alpha \leftarrow \mathbf{0}$$

$$\text{for } i = 1, \cdots, n \text{ do}$$

$$\hat{y}^{(i)} \leftarrow \operatorname{sign} \left\{ \sum_{j=1}^{n} \alpha^{(j)} \eta y^{(j)} \mathbf{x}^{(i)} \mathbf{x}^{(j)^{T}} \right\}$$

$$\text{inner product}$$

$$\text{if } \hat{y} \neq y^{(i)} \text{ then}$$

$$\alpha^{(i)} \leftarrow \alpha^{(i)} + 1$$

$$\text{end if}$$

$$\text{end for}$$



Parameters are linear combination of the samples:

$$oldsymbol{ heta} = \sum_{j=1}^n lpha^{(j)} \eta y^{(j)} \mathbf{x}^{(j)}^T$$

Dual formulation

$$y^{(i)} \in \{1, -1\}$$

we predict as

$$\hat{y}^{(i)} = \operatorname{sign} \left\{ \sum_{j=1}^{n} \alpha^{(j)} \eta y^{(j)} \mathbf{x}^{(i)} \mathbf{x}^{(j)}^{T} \right\}$$

we train as

$$\begin{aligned} & \boldsymbol{\alpha} \leftarrow \mathbf{0} \\ & \mathbf{for} \ i = 1, \cdots, n \ \mathbf{do} \\ & \hat{y}^{(i)} \leftarrow \mathrm{sign} \left\{ \sum_{i=1}^n \alpha^{(j)} \eta y^{(j)} \mathbf{x}^{(i)} \mathbf{x}^{(j)^T} \right\} \\ & \mathbf{if} \ \hat{y} \neq y^{(i)} \ \mathbf{then} \\ & \alpha^{(i)} \leftarrow \alpha^{(i)} + 1 \\ & \mathbf{end} \ \mathbf{if} \\ & \mathbf{end} \ \mathbf{for} \end{aligned}$$

- Kernel perceptron
  - we predict as

$$\hat{y}^{(i)} = \operatorname{sign} \left\{ \sum_{j=1}^{n} \alpha^{(j)} y^{(j)} \overline{\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)})} \right\}$$

— we train as

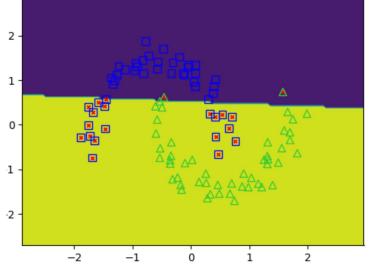
#### Implementation from scratch in Python:

03\_kernel\_perceptron.ipynb

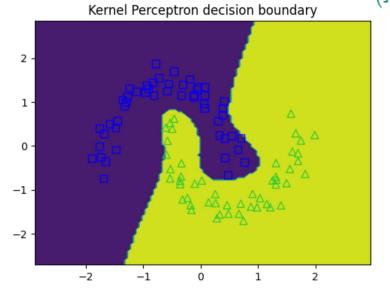


(p=1)





$$(\mathbf{x}^{(i)}\mathbf{x}^{(j)}^T + 1)^p$$





## Nonlinear decision boundaries

Kernel logistic regression (extra, not syllabus)



## Kernel logistic regression

- Logistic regression can be kernelized analogously to the perceptron by switching to a dual formulation
- Generalized linear models (like logistic regression) can be turned into kernel machines by considering a feature mapping of the form:

$$\boldsymbol{\phi}(\mathbf{x}^{(i)}) = [\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(i)}), \dots, \kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(n)})]$$

Number of features is number of samples (or somewhat less for sparse kernel machines)

- This corresponds to the realization that the weights can be written as a linear combination of the samples
- When only a subset of the samples is used (**support vectors**), we call the resulting model a **sparse kernel machine**



## Kernel logistic regression versus Kernel SVM

- Kernel SVMs yields sparse kernel machines which are faster in prediction and have lower memory consumption
- Kernelized Logistic regression uses all data points in kernel matrix, resulting in a dense kernel machine
- Kernelized Logistic regression with sparsity-promoting regularization like L1regularization yields a sparse predictor (samples with zero weights can be dropped)



