Lecture 12: Classification: SVM

Course: Biomedical Data Science
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Fall 2018

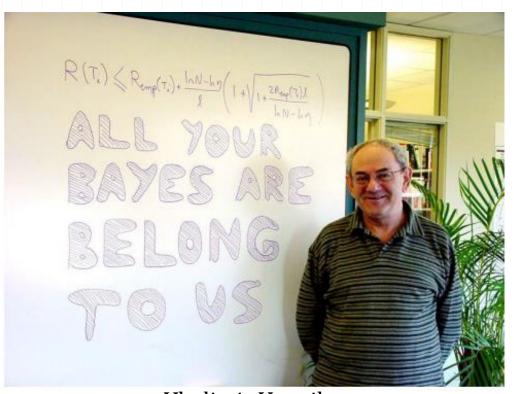
Administrative Items

- Grad Survey proposal topic due in two weeks
 - Provide a link to the survey paper on canvas
 - More comprehensive understanding is expected

Agenda

- k-NN
- Decision Tree based Methods
- Support Vector Machines
- Neural Networks
- Deep learning
- NLP

Support Vector Machines (SVM)

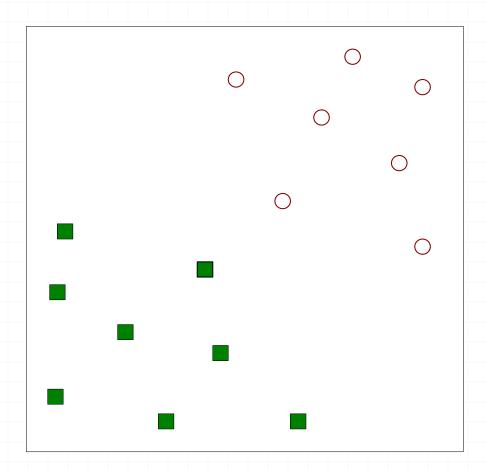


Vladimir Vapnik

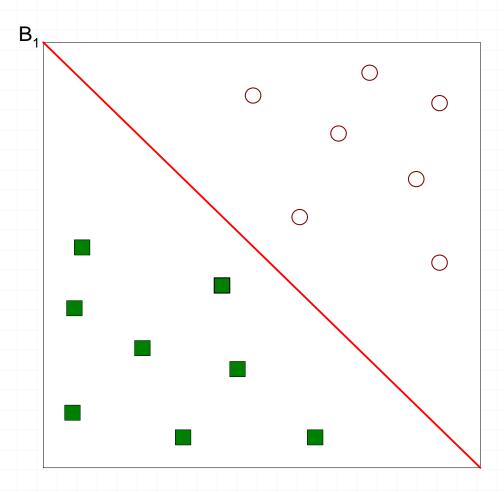
Support Vector Machine

- Represents the decision boundary using only a subset of training examples known as support vectors
- Convex optimization problems with a unique solution
- Normally, a linear classifier
 - But, we can also use non-linear kernels
 - Kernels: application-specific measures of similarity
 - Leading to "kernel machines"

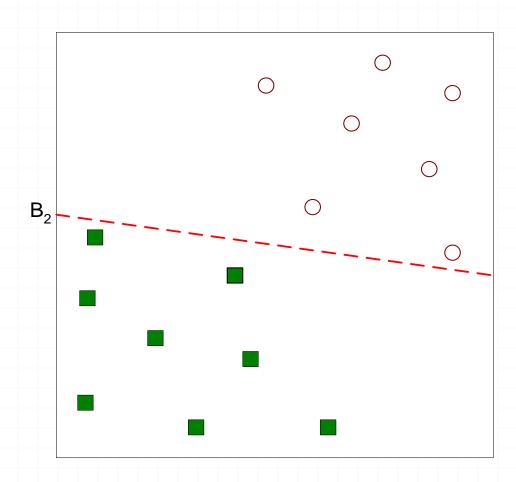
- Question
 - Find a linear hyperplane to separate the data



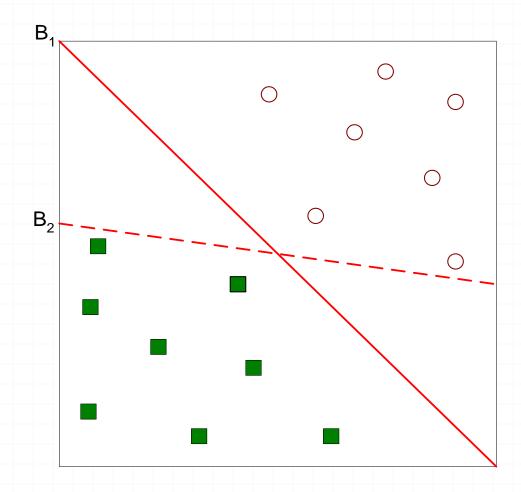
One possible solution



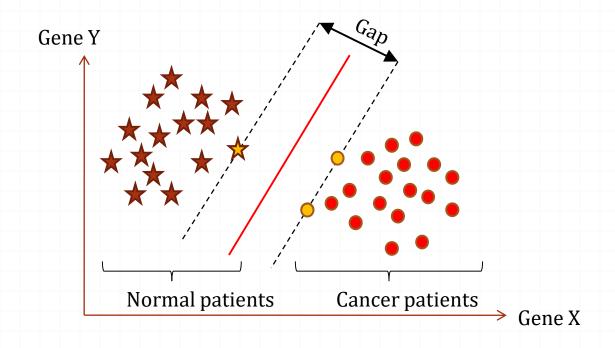
Another possible solution



• Which one is better?



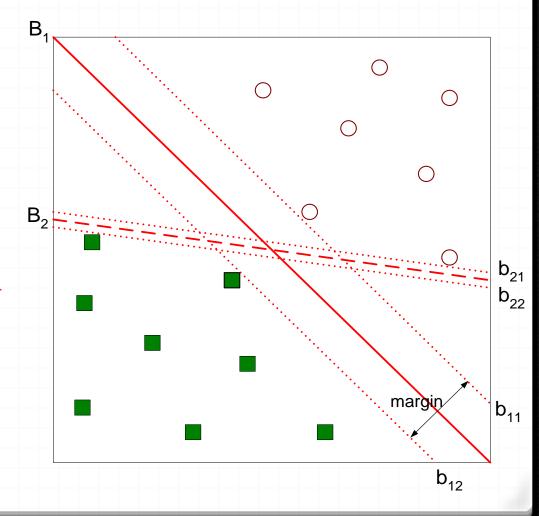
Main ideas of SVMs



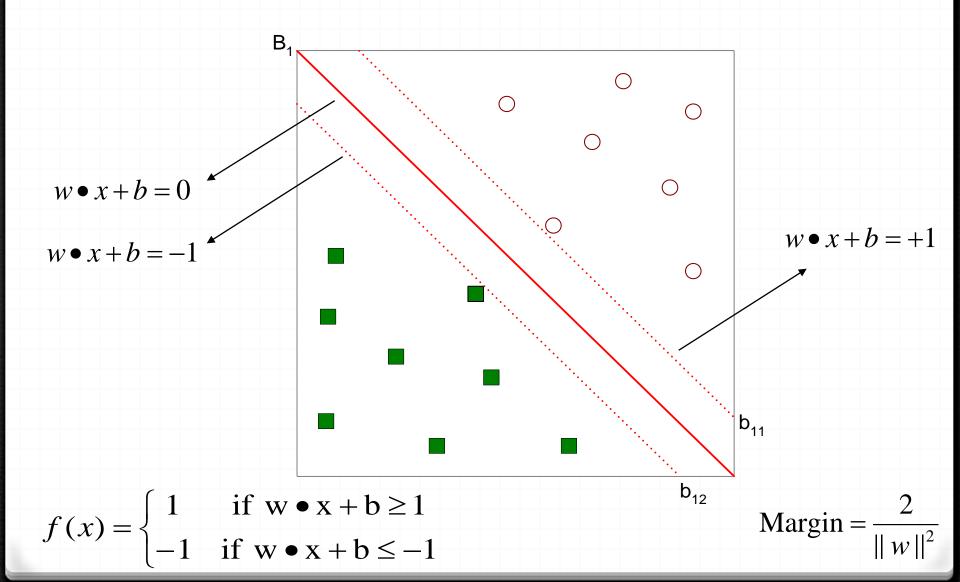
- Represent patients geometrically (by "vectors");
- Find a linear decision surface ("hyperplane") that can separate patient classes <u>and</u> has the largest distance (i.e., largest "gap" or "margin") between border-line patients (i.e., "support vectors");

Better Decision Boundary

- Margin definition:
 - Distance from the hyperplane to the closest instances on either side
- Find hyperplane that maximizes the margin
- B1 is better than B2
 - Wider margin, and better generalization



Support Vector Machine



Let's Formulate this ...

Our simple rule

It can be re-written

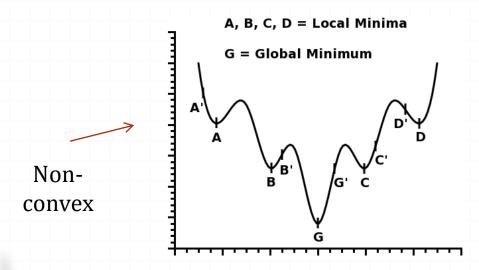
$$f(x) = \begin{cases} 1 & \text{if } \mathbf{w} \bullet \mathbf{x} + \mathbf{b} \ge 1 \\ -1 & \text{if } \mathbf{w} \bullet \mathbf{x} + \mathbf{b} \le -1 \end{cases} \qquad \mathbf{y}_i(\mathbf{w}. \mathbf{x}_i + \mathbf{b}) \ge 1$$

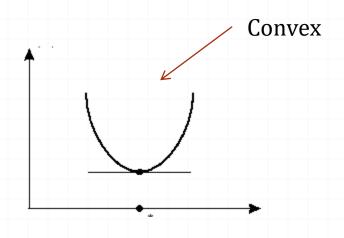
 But, we also want to maximize the margin (or minimize the inverse of the margin), so we get:

$$\min(\left|\frac{||\boldsymbol{w}||^2}{2}\right)$$
 Subject to $y_i(\boldsymbol{w}, \boldsymbol{x}_i + b) \ge 1$ $i = 1, 2, ... N$

A Convex Optimization Problem

- This is a convex optimization problem (quadratic programming or QP)
 - Therefore, it has only a single optimum (great!)
 - Therefore no worries about convergence, learning rates, etc.





SVM Training

- Involves estimating parameters w and b
- This is a quadratic optimization programming (QP) problem
 - Can be solved using Lagrange multiplier technique

$$\min(\left|\frac{||\boldsymbol{w}||^2}{2}\right)$$
 Subject to $y_i(\boldsymbol{w}, \boldsymbol{x}_i + b) \ge 1$ $i = 1, 2, ... N$

Lagrange Multiplier

Write its Lagrangian:

(0)
$$L_p = \frac{||\mathbf{w}||^2}{2} - \sum_{i=1}^N \lambda_i (y_i(\mathbf{w}. \mathbf{x}_i + b) - 1)$$
 where $\lambda_i \ge 0$

- Here λ_i are called the Lagrange multipliers
- We will minimize L_p with respect to w and b:

(1)
$$\frac{\partial L_p}{\partial \mathbf{w}} = 0 \quad \Rightarrow \mathbf{w} = \sum_{i=1}^{N} \lambda_i y_i x_i$$

(2) $\frac{\partial L_p}{\partial b} = 0 \quad \Rightarrow \sum_{i=1}^{N} \lambda_i y_i = 0$

$$\frac{\partial L_p}{\partial b} = 0 \quad \Rightarrow \sum_{i=1}^{N} \lambda_i y_i = 0$$

KKT Conditions

Additionally, the following should hold (KKT condition, Karush-Kuhn-Tucker):

$$(3) \lambda_i \geq 0$$

(4)
$$\lambda_i[y_i(\mathbf{w}.\mathbf{x}_i + b) - 1] = 0$$

Support Vectors

In equation (4) we had,

(4)
$$\lambda_i[y_i(\mathbf{w}.\mathbf{x}_i + b) - 1] = 0$$

To be zero, either λ_i must be zero, or

$$y_i(\mathbf{w}.\mathbf{x}_i + b) = 1$$

which means that such a training instance \mathbf{x}_i lies along the hyperplanes b_1 or b_2

Such a training instance is known as a support vector

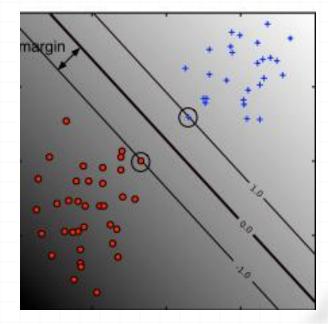
Support Vectors

- A subset of training set, which are on the decision boundary
 - i.e. those for which $\lambda_i > 0$

• For the other training instances (i.e. those far away

from the boundary), $\lambda_i = 0$

So, they have no effect on the hyperplane.



Final Solution

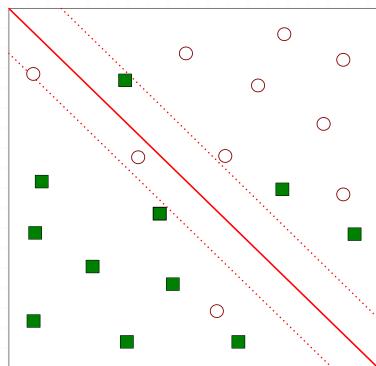
- We can obtain b and λ using numerical solutions from equations (0)-(4) (we won't go into details)
- The final solution will be:

We use this equation in order to classify a new data point z.
$$f(z) = sign(\mathbf{w}.\mathbf{z} + b) = sign\left(\sum_{i=1}^{n} \lambda_i y_i \mathbf{x}_i.\mathbf{z} + b\right)$$

Non-separable case

• How decision boundary can be modified to tolerate small training error?

- Instance may lie on the wrong side
- 2. Instance may be inside the margin



Soft Margin

- We should consider a decision boundary that is tolerable to small training errors
 - This approach is known as soft margin
 - But SVM must consider the trade-off between the width of the margin and the number of training errors committed by the boundary

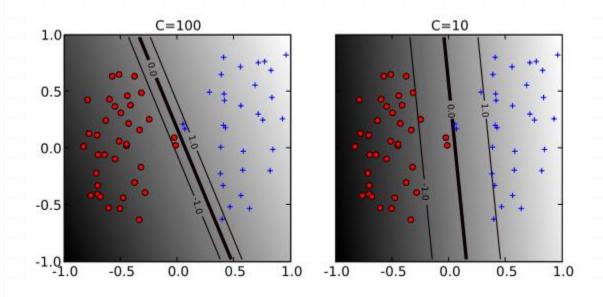
Soft Margin

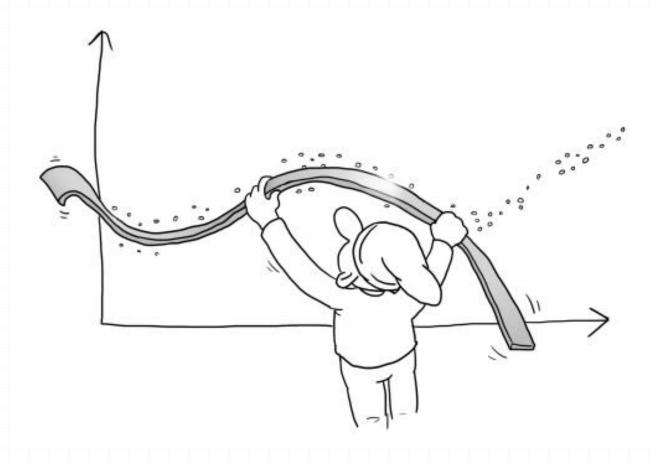
- The inequality constrained should be relaxed
 - Introduce the slack variables ξ
- The objective function must be modified to penalize a decision boundary with large values of slack variables

$$\min(\frac{\left||w|\right|^2}{2} + C(\sum_{i=1}^N \xi_i))$$

$$f(x_i) = \begin{cases} 1 & \text{if } \mathbf{w} \bullet \mathbf{x}_i + \mathbf{b} \ge 1 - \xi_i \\ -1 & \text{if } \mathbf{w} \bullet \mathbf{x}_i + \mathbf{b} \le -1 + \xi_i \end{cases}$$

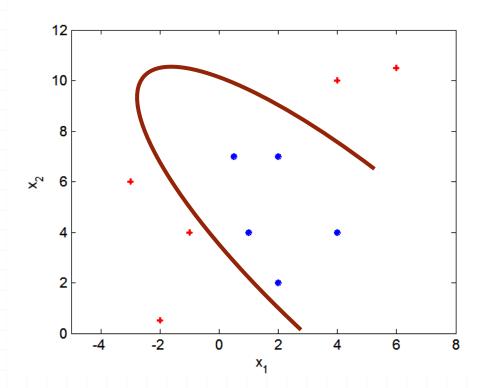
Effect of C



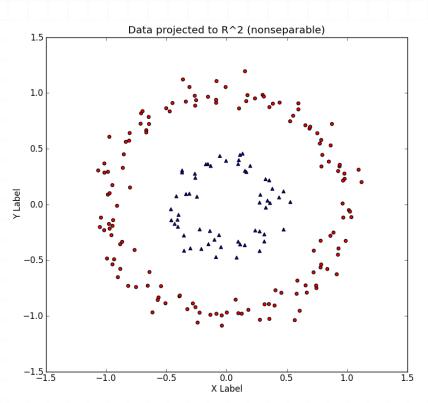


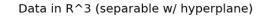
Nonlinear Decision Boundary

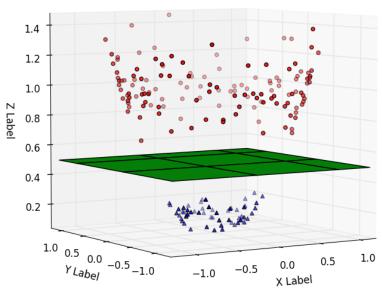
• What if the decision boundary is not linear?



Separable in higher-dimension







Kernels in Higher Dimensions Animated

- 1. https://www.youtube.com/watch?t=42&v=3liCbRZ
 PrZA
- 2. https://www.youtube.com/watch?v=9NrALgHFwTo

Nonlinear Decision Boundary

The trick is to transform data form its original coordinates space x into a new space $(\phi(x))$

$$\min(|\frac{||w||^2}{2})$$

Subject to
$$y_i(\mathbf{w}. \phi(\mathbf{x}_i) + b) \ge 1$$
 $i = 1, 2, ... N$

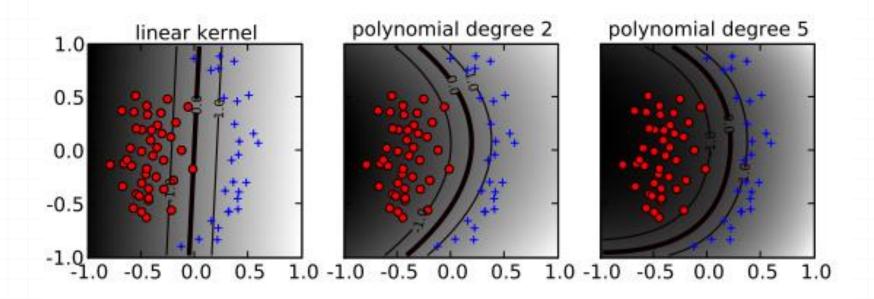
Solution



We use this data point z.

We use this equation in order to classify a new data point z
$$f(z) = sign\left(\sum_{i=1}^{n} \lambda_i y_i \phi(x_i). \phi(z) + b\right)$$

Example



Popular kernels

A kernel is a dot product in some feature space:

$$K(x_i, x_j) = \Phi(x_i) \cdot \Phi(x_j)$$

Examples:

$$K(x_i, x_j) = x_i \cdot x_j$$

$$K(x_i, x_j) = \exp(-\gamma \|x_i - x_j\|^2)$$

$$K(x_i, x_j) = \exp(-\gamma \|x_i - x_j\|)$$

$$K(x_i, x_j) = (p + x_i \cdot x_j)^q$$

$$K(x_i, x_j) = (p + x_i \cdot x_j)^q \exp(-\gamma \|x_i - x_j\|^2)$$

$$K(x_i, x_j) = \tanh(kx_i \cdot x_j - \delta)$$

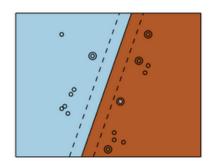
Linear kernel
Gaussian kernel
Exponential kernel
Polynomial kernel
Hybrid kernel
Sigmoidal

SVM Characteristics

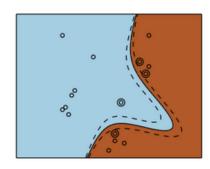
- Convex optimization problem
 - Finds global minimum
 - Other methods such as neural network find local minimum
- Parameters should be tuned
 - Kernel type, C, ..
- High computational demands (both test and training)
 - Training stage
 - Naïve QP implementation
 - $0(n^3)$
 - More efficient techniques
 - Between O(n) and O(n²)

SVM in Scikit-learn (Python)

Linear kernel



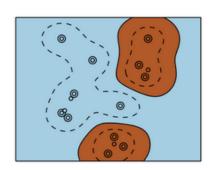
Polynomial kernel



```
>>> svc = svm.SVC(kernel='linear
```

```
>>> svc = svm.SVC(kernel='poly',
... degree=3)
>>> # degree: polynomial degree
```

RBF kernel (Radial Basis Function)



```
>>> svc = svm.SVC(kernel='rbf')
>>> # gamma: inverse of size of
>>> # radial kernel
```

Neuroimaging Data (fMRI)

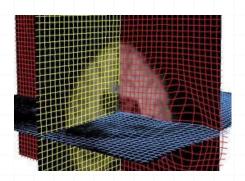
Neuroimaging Data Analysis

- Classify structural MR images
 - identify disease precursors as early as possible
 - e.g. MCI; Alzheimer's disorder (AD)
 - identify structural trajectories in neurodevelopment disorders
- Classify functional MR images
 - identify brain state associated with a stimulus



Neuroimaging Data

- Each image has lots and lots of voxels.
 - Of those, ~30,000 are actually brain voxels
 - grey matter ≈ 23,000 voxels
 - white matter $\approx 8,000$ voxels
- To reduce the feature-space, we can:
 - resample data
 - do some latent variable extraction
 - e.g. using Principal Components Analysis (PCA) or some other automated feature extraction
- Then run SVM using a smaller number of "features"
 - e.g. 3-8 PCs rather than thousands of voxels

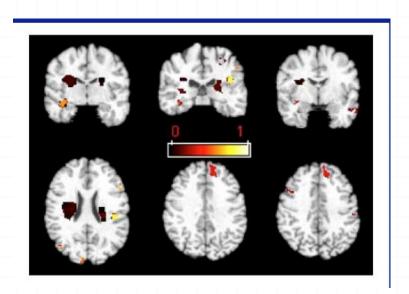


Example Features

- Average. For each ROI, calculate the mean activity over all voxels in the ROI. Use these ROI means as the input features.
- ActiveAvg(n). For each ROI, select the most active voxels, then calculate the mean of their values. Again, use these ROI means as the input features.
- "most active" voxels are those whose activity while performing the task varies the most from their activity when the subject is at rest.
- Active(n). Select the most active voxels over the entire brain. Use only these voxels as input features.

Example: Schizophrenia

• Motivation: Which brain areas best distinguish brains of female schizophrenic patients from typical?



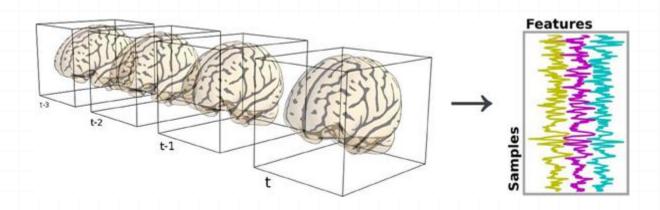
Analysis with scikit-learn

#First we need to read data (and its mask)

#nibabel is a reader of common neuroimaging formats

>>import nibabel as ni

>>> X = ni.load("bold.nii").get_data()



Preprocessing

- De-trending: remove linear trend (scipy.signal.detrend)
- Normalization: all features will have the same range (sklearn.preprocessing.normalize)
- Frequency filtering: e.g. low frequencies due to physiological mechanisms, use Fourier Transform (scipy.fftpack.fft)

Prediction

```
from sklearn.linear_model import
  LogisticRegression as LR
from sklearn.cross_validation import
   cross val score
pipeline_LR = Pipeline([('selection',
   SelectKBest(f_classif, 500)),
   ('clf', LR(penalty='11', C=0.05)])
scores lr = []
for pixel in y_train.T:
    score = cross_val_score(pipeline_LR,
       X_train, pixel, cv=5)
    scores_lr.append(score)
```