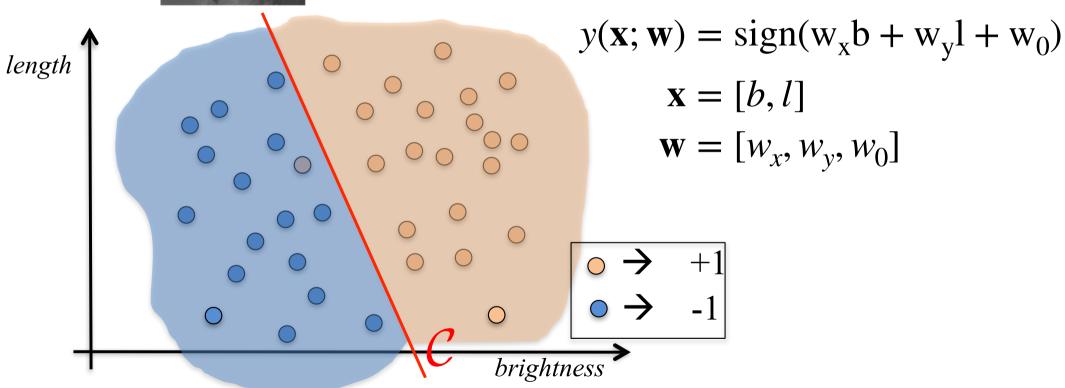
Linear Classification

Pascal Fua IC-CVLab



Reminder: Linear 2D Model





How do we find w?



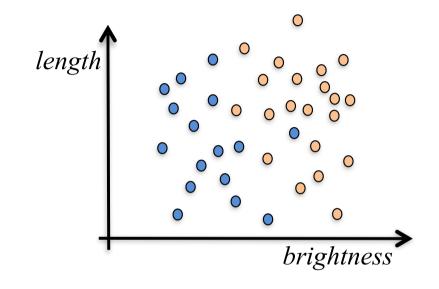
Reminder: Training vs Testing

Supervised training:

Given a **training** set $\{(\mathbf{x}_n, t_n)_{1 \le n \le N}\}$ minimize:

$$E(\mathbf{w}) = \sum_{n=1}^{N} L(y(\mathbf{x}_n; \mathbf{w}), t_n)$$

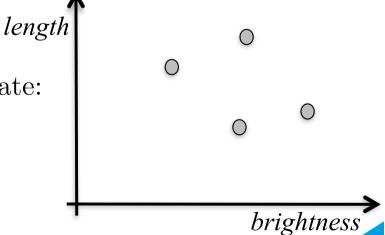
$$= \sum_{n=1}^{N} [y(\mathbf{x}_n; \mathbf{w}) \neq t_n]$$



Testing:

Given a **test** set $\{(\mathbf{x}_n, t_n)_{1 \leq n \leq N}\}$ compute the error rate:

$$1/N \sum_{n=1}^{N} [y(\mathbf{x}_n; \mathbf{w}) \neq \mathbf{t}_n]$$

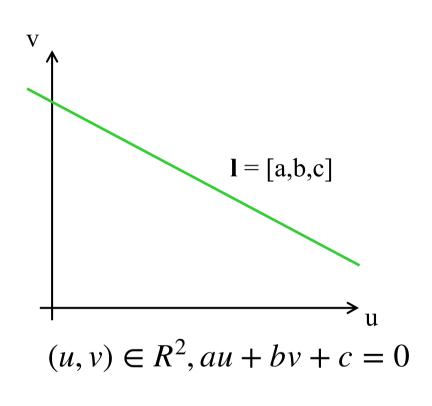


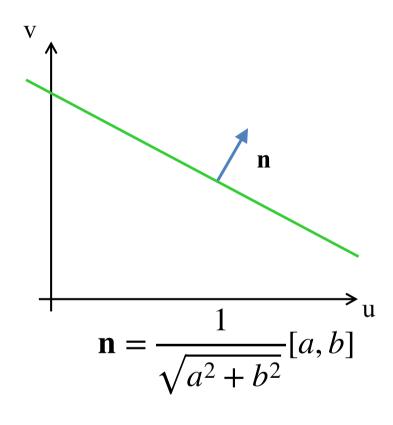


Parameterizing Lines

Equation of a line

Normal vector



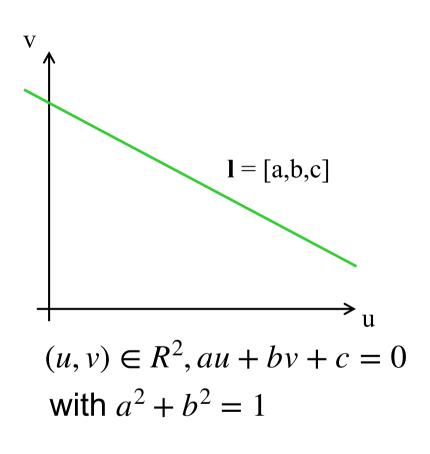


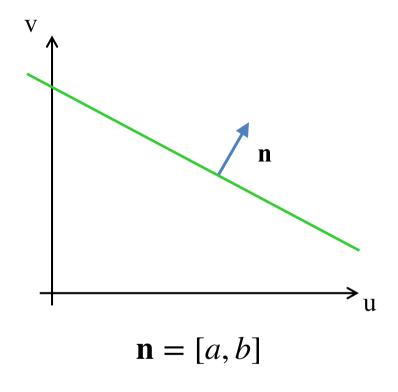
$$[a,b,c]$$
 and $\frac{1}{\sqrt{a^2+b^2}}[a,b,c]$ define the same line.

Normalized Parameterization

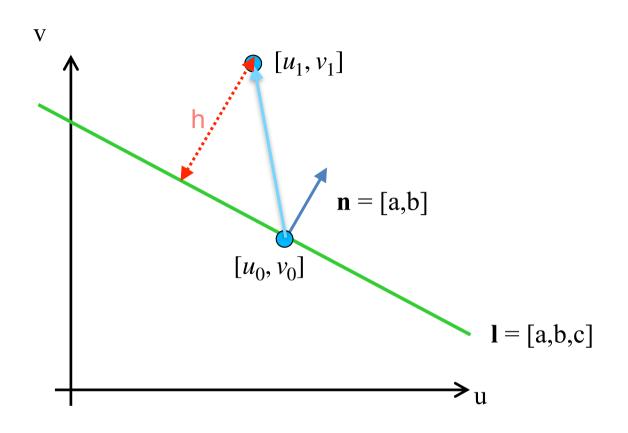
Equation of a line







Signed Distance to Line



Signed distance:
$$h = \mathbf{n} \cdot [u_1 - u_0, v_1 - v_0]$$

 $= a(u_1 - u_0) + b(v_1 - v_0)$
 $= au_1 + bv_1 - (au_0 - bv_0)$
 $= au_1 + bv_1 + c - (au_0 - bv_0 - c)$
 $= au_1 + bv_1 + c$

h=0: Point is on the line.

h>0: Point on one side.

h<0: Point on the other side.

Line Going Through 2 Points

$$\mathbf{x}_1 = (u_1, v_1)$$
 $\mathbf{n} = (a, b)$
 $\mathbf{1} = (a, b, c)$
 $\mathbf{x}_2 = (u_2, v_2)$

Points belong to line:

$$au_1 + bv_1 + c = 0$$

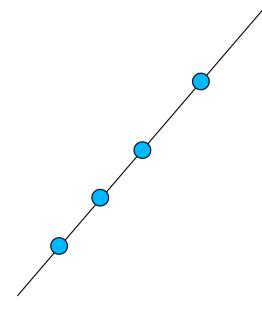
$$au_2 + bv_2 + c = 0$$

subject to:

$$a^2 + b^2 = 1$$

Line Going Through N Points

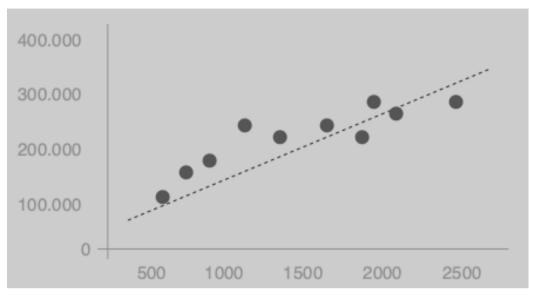
$$\mathbf{x}_i = (u_i, v_i)$$



In practice, this never happens due to noise.

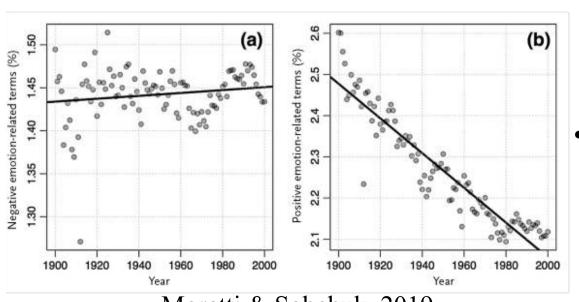


In Real Life



Price of a house a function of its size

https://www.internalpointers.com/post/linear-regression-one-variable



 Proportion of negative and positive emotions in anglophone fiction.

Moretti & Sobchuk, 2019

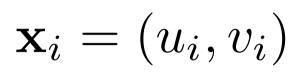


Fitting a Line to N Points

$$\mathbf{x}_i = (u_i, v_i)$$
Orthogonal distance to line:
$$h_i = au_i + bv_i + c \text{ if } a^2 + b^2 = 1$$

 \rightarrow We want to minimize $\sum_{i} (au_i + bv_i + c)^2$ w.r.t. a, b, and c, subject to $a^2 + b^2 = 1$.

Centering the Coordinates







Original points:

$$\mathbf{x}_{i} = (u_{i}, v_{i})_{1 \le i \le n}$$

$$\overline{\mathbf{x}} = \frac{1}{N} \sum_{i} \mathbf{x}_{i}$$

Centered points:

$$\mathbf{x}_{i}^{c} = \mathbf{x}_{i} - \overline{\mathbf{x}}$$

$$\overline{\mathbf{x}^{c}} = \mathbf{0}$$

—> The points can always be translated so that their center of gravity is at the origin.



Minimization using Centered Points

Minimize $\sum_{i} (au_i + bv_i + c)^2$ w.r.t. a, b, and c, subject to $a^2 + b^2 = 1$.

Minimize
$$\sum_{i} (au_i + bv_i)^2 + 2c(a\sum_{i} u_i + b\sum_{i} v_i) + \sum_{i} c^2$$
.

Zero if coordinates are centered. c=0

Minimize
$$\|\mathbf{M} \begin{bmatrix} a \\ b \end{bmatrix}\|^2$$
 subject to $a^2 + b^2 = 1$, with $\mathbf{M} = \begin{bmatrix} u_1 & v_1 \\ u_2 & v_2 \\ \vdots & \vdots \\ u_n & v_n \end{bmatrix}$.

 $\rightarrow \begin{bmatrix} a \\ b \end{bmatrix}$ is the eigenvector associated to the smallest eigenvalue of $\mathbf{M}^T \mathbf{M}$.



Optional: Proof Sketch

Let us consider the symmetric matrix $A = M^TM$ of size N x N:

- We can write $A = R^T DR$ where D is diagonal and $R^T R = I$, therefore $\forall X \in R^N$, $||AX||^2 = X^T A^T A X = X^T M X = (RX)^T D(RX)$.
- R is a rotation matrix and $\forall X \in R^N, ||RX|| = ||X||.$
- Let $X \in \mathbb{R}^N$, ||X|| = 1 and $Y = \mathbb{R}X$, we have $||AX||^2 = Y^T DY$ with ||Y|| = 1.
- D is a diagonal matrix, therefore we have

$$\begin{split} D &= \begin{bmatrix} \lambda_1 & 0 & 0 \\ 0 & \ddots & 0 \\ 0 & 0 & \lambda_N \end{bmatrix} \;, \\ &\Rightarrow \lambda_1 \leq \|DY\| \leq \lambda_N \text{ if } \|Y\| = 1 \;. \end{split}$$

This result applies in any dimension.



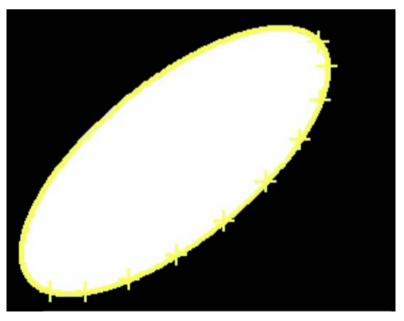
Optional: Fitting Ellipses

For each point:

$$d_a(\mathbf{X}_i, \mathbf{X}) = au_i^2 + bu_i v_i + cv_i^2 + du_i + ev_i + f$$
$$= \begin{bmatrix} u_i^2 & u_i v_i & v_i^2 & u_i & v_i & 1 \end{bmatrix} \bullet \mathbf{X}$$

Minimize:

$$\sum_{i} d_a(\mathbf{x}_i, \mathbf{X})^2 = \|\mathbf{A}\mathbf{X}\|^2 \text{ subject to } \|\mathbf{X}\| = 1$$



where
$$\mathbf{A} = \begin{bmatrix} u_1^2 & u_1 v_1 & v_1^2 & u_1 & v_1 & 1 \\ \dots & \dots & \dots & \dots & \dots \\ u_n^2 & u_n v_n & v_n^2 & u_n & v_n & 1 \end{bmatrix}$$

The line and ellipse fitting algorithms we showed are examples of an important technique known as the Direct Linear Transform.

Optional: Generalization

Line:

$$\mathbf{x}_i = \begin{bmatrix} u_i, v_i \end{bmatrix}$$
 $\mathbf{w} = \begin{bmatrix} a, b \end{bmatrix}$
 $\phi(\mathbf{x}) = \begin{bmatrix} u, v, 1 \end{bmatrix}$
 $t_i = 0$

Ellipse:

$$\mathbf{x}_{i} = [u_{i}, v_{i}]$$

$$\mathbf{w} = [a, b, c, d, e, f]$$

$$\phi(\mathbf{x}) = [u^{2}, uv, v^{2}, u, v, 1]$$

$$t_{i} = 0$$

In both case we minimize:

$$\sum_{i} d_{a}(\mathbf{x}_{i}, \mathbf{X})^{2} = \sum_{i} (\mathbf{w}^{T} \phi(\mathbf{x}_{i}) - t_{i})^{2}$$

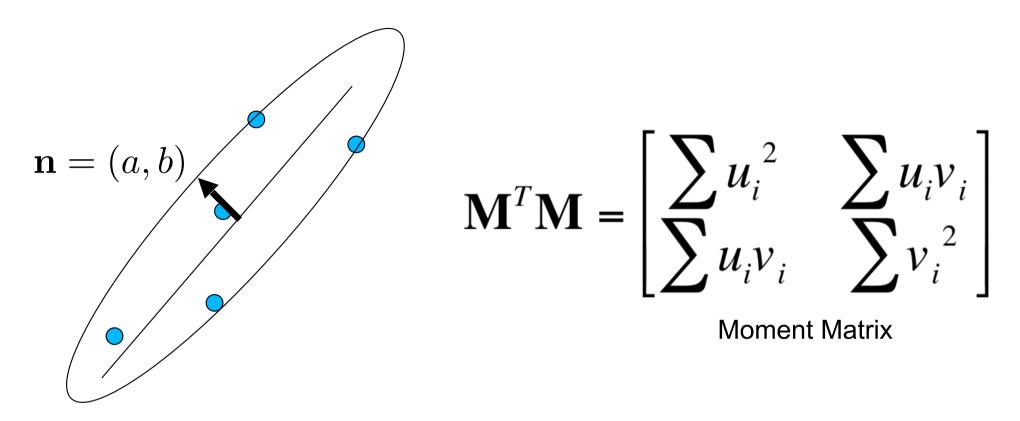
$$\Rightarrow \forall i, \mathbf{w}^{T} \phi(\mathbf{x}_{i}) \approx t_{i}$$

We will encounter this formulation again later in the class.



Feature Vector

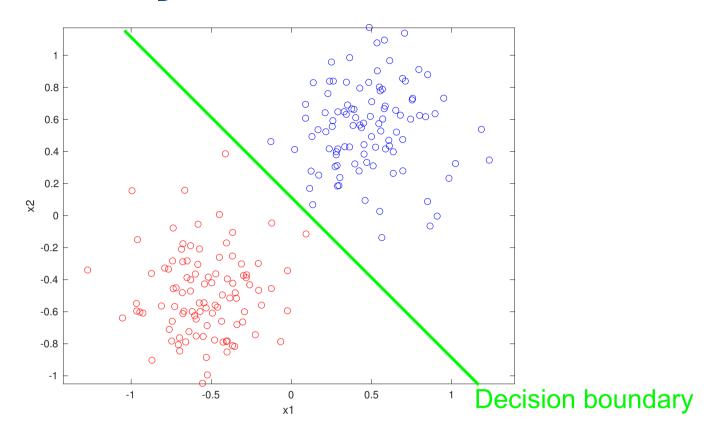
Back to 2D Lines



- The eigenvector corresponding to the largest eigenvalue of **M**^T**M** is the direction of the fitted line.
- The eigenvector corresponding to the smallest eigenvalue is its normal.
- The ratio of the two eigenvalues indicates how much noise there is:
 - zero without noise,
 - close to one if the points are randomly distributed.



Binary Classification

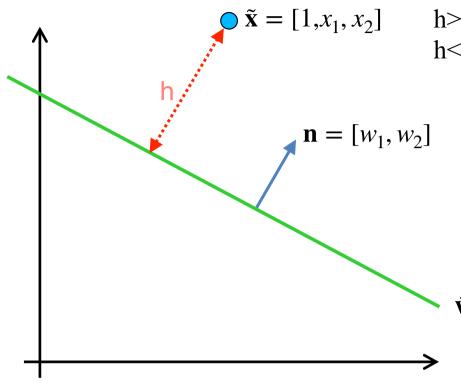


Two classes shown as different colors:

- The label $y \in \{-1,1\}$ or $y \in \{0,1\}$.
- The samples with label 1 are called positive samples.
- The samples with label -1 or 0 are called negative samples.



Signed Distance Reformulated



h=0: Point is on the line.

h>0: Point in the normal's direction.

h<0: Point in the other direction.

$$\tilde{\mathbf{w}} = [w_0, w_1, w_2] \text{ with } w_1^2 + w_2^2 = 1$$

Notation:

$$\mathbf{x} = [x_1, x_2]$$

$$\tilde{\mathbf{x}} = [1, x_1, x_2]$$

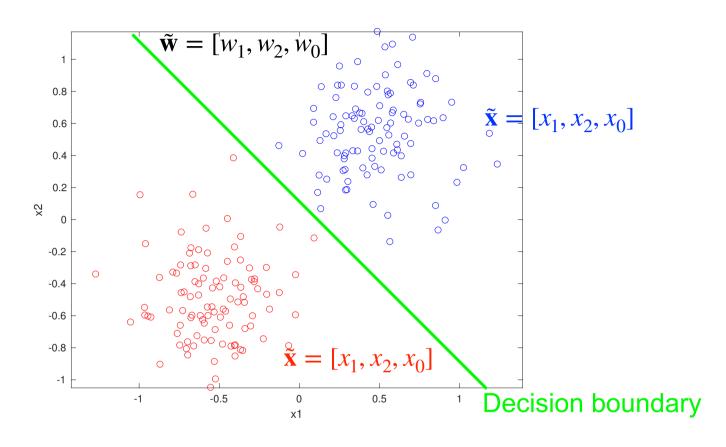
Signed distance:

$$h = w_0 + w_1 x_1 + w_2 x_2$$

$$= \tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}}$$



Problem Statement in 2D

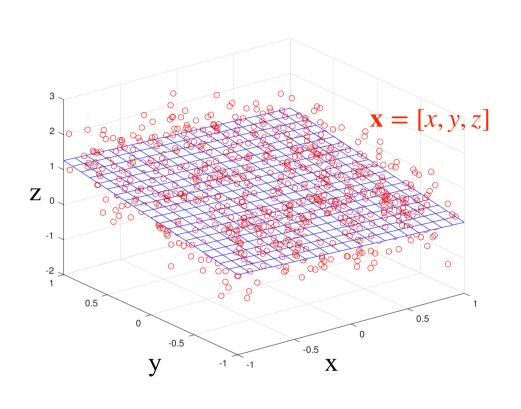


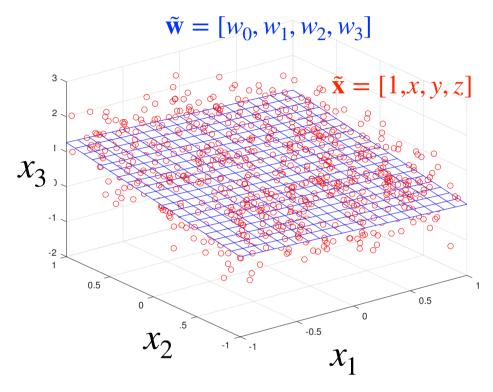
Find $\tilde{\mathbf{w}}$ such that:

- For all or most positive samples $\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} > 0$.
- For all or most negative samples $\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} < 0$.



Signed Distance in 3D





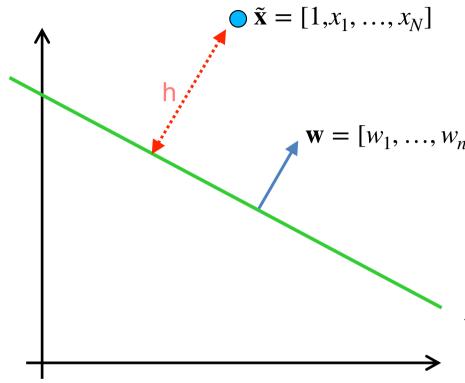
$$\mathbf{x} \in R^3, \, 0 = ax + by + cz + d$$

$$\tilde{\mathbf{x}} \in R^4, \, \tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} = 0$$

Signed distance $h = \tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}}$ if $w_1^2 + w_2^2 + w_3^2 = 1$.



Signed Distance in N Dimensions



h=0: Point is on the decision boundary.

h>0: Point on one side.

h<0: Point on the other side.

$$\tilde{\mathbf{w}} = [w_0, w_1, ..., w_n] \text{ with } \sum_{i=1}^{N} w_i^2 = 1$$

Notation:

$$\mathbf{x} = [x_1, ..., x_n]$$

$$\tilde{\mathbf{x}} = [1, x_1, \dots, x_n]$$

Hyperplane:

$$\mathbf{x} \in R^n, \quad 0 = \tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} \\ = w_0 + w_1 x_1 + \dots w_n x_n$$

Signed distance: $h = \tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}}$

$$h = \tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}}$$



Problem Statement in N Dimensions

Hyperplane: $\mathbf{x} \in R^N$, $\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} = 0$, with $\tilde{\mathbf{x}} = [1 \mid \mathbf{x}]$.

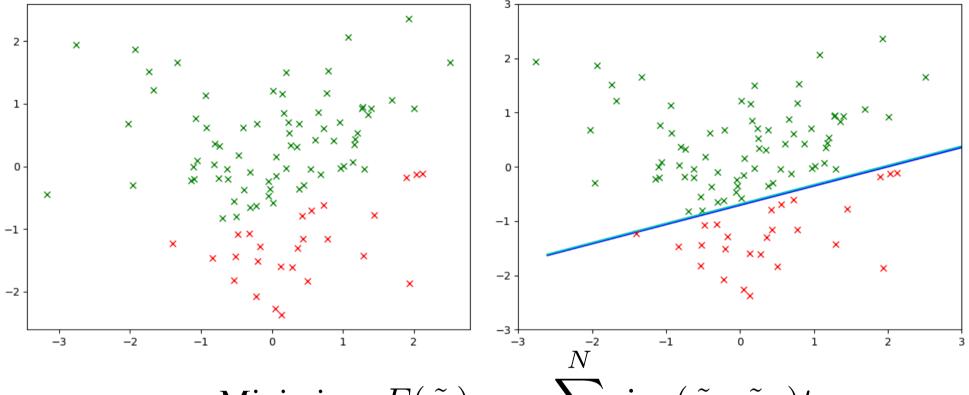
Signed distance: $\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}}$, with $\tilde{\mathbf{w}} = [w_0 | \mathbf{w}]$ and $||\mathbf{w}|| = 1$.

Find w such that

- for all or most positive samples $\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} > 0$,
- for all or most negative samples $\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} < 0$.



Perceptron

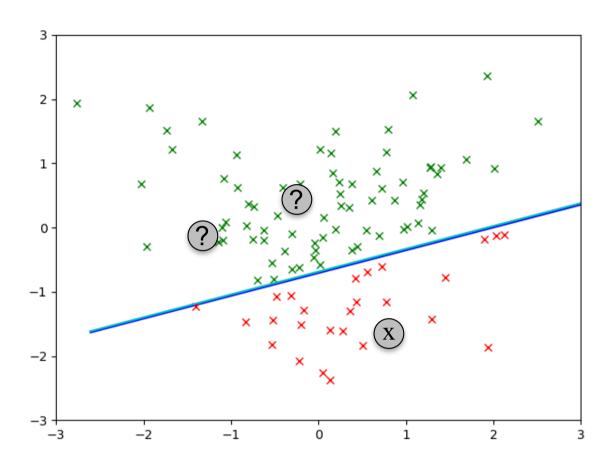


Minimize:
$$E(\tilde{\mathbf{w}}) = -\sum_{n=1} \operatorname{sign}(\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}}_n) t_n$$

- Set $\tilde{\mathbf{w}}_1$ to $\mathbf{0}$.
- Iteratively, pick a random index n.
 - If $\tilde{\mathbf{x}}_n$ is correctly classified, do nothing.
 - Otherwise, $\tilde{\mathbf{w}}_{t+1} = \tilde{\mathbf{w}}_t + t_n \tilde{\mathbf{x}}_n$.



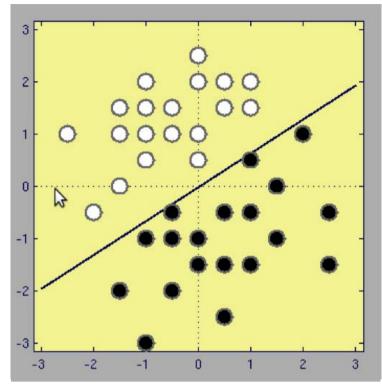
Test Time



$$y(\mathbf{x}; \tilde{\mathbf{w}}) = \begin{cases} 1 & \text{if } \tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} \ge 0, \\ -1 & \text{otherwise.} \end{cases}$$
$$\tilde{\mathbf{x}} = [1, x_1, ..., x_n]$$



Centered Perceptron



If the two populations are of the same size, the decision boundary can be assumed to go through the center of gravity.

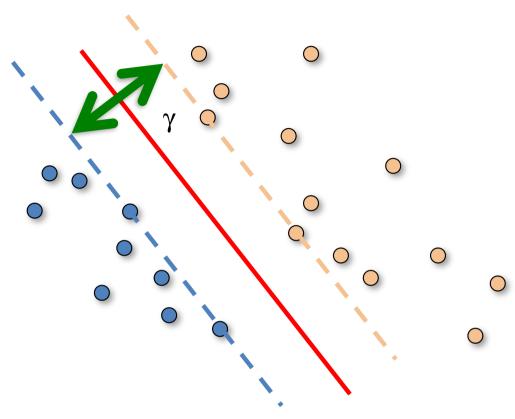
Given a **training** set $\{(\mathbf{x}_n, t_n)_{1 \le n \le N}\}$ minimize:

$$E(\mathbf{w}) = -\sum_{n=1}^{N} \operatorname{sign}(\mathbf{w} \cdot \mathbf{x}_n) t_n$$

- Center the \mathbf{x}_n s so that $w_0 = 0$.
- Set \mathbf{w}_1 to $\mathbf{0}$.
- Iteratively, pick a random index n.
 - If \mathbf{x}_n is correctly classified, do nothing.
 - Otherwise, $\mathbf{w}_{t+1} = \mathbf{w}_t + t_n \mathbf{x}_n$.



Convergence Theorem



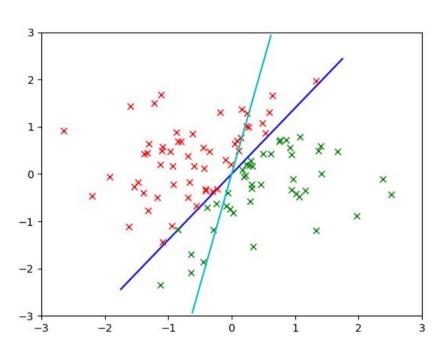
 γ is the margin

If there exists a number $\gamma > 0$ and a parameter vector \mathbf{w} ,* with $\|\mathbf{w}\| = 1$, such that

$$\forall n \quad t_n(\mathbf{x}_n \cdot \mathbf{w}^*) \ge \gamma ,$$

then the perceptron algorithm makes at most $\frac{R^2}{\gamma^2}$ errors, where $R = max_n ||\mathbf{x}_n||$.

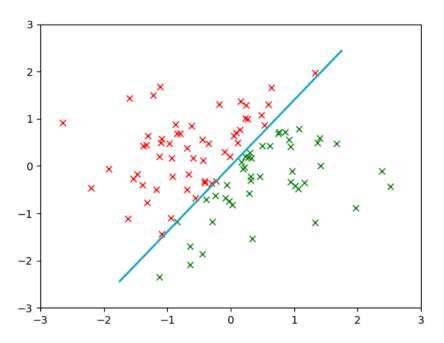
What if γ is Small?



for n in range(nIt): for i in range(ns):

- If \mathbf{x}_n is correctly classified, do nothing.
- Otherwise, $\mathbf{w}_{t+1} = \mathbf{w}_t + t_n \mathbf{x}_n$.

Randomizing helps!

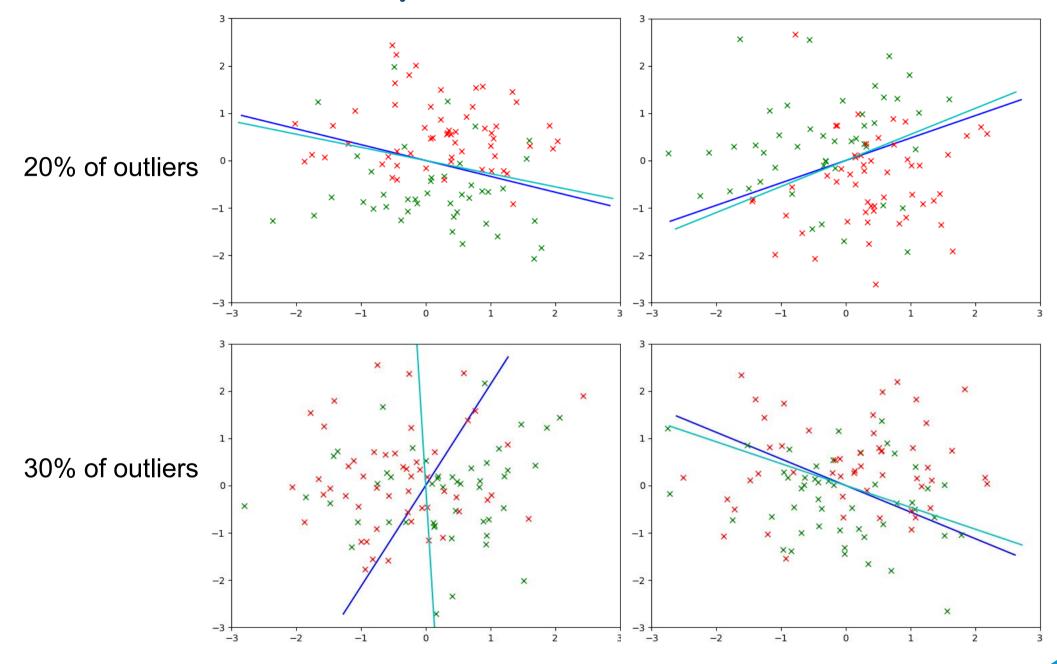


for n in range(nIt): inds=list(range(ns)) random.shuffle(inds) for i in range(inds):

- If \mathbf{x}_n is correctly classified, do nothing.
- Otherwise, $\mathbf{w}_{t+1} = \mathbf{w}_t + t_n \mathbf{x}_n$.



What if γ Does Not Exist?



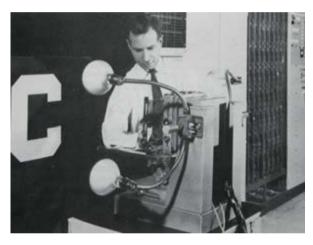


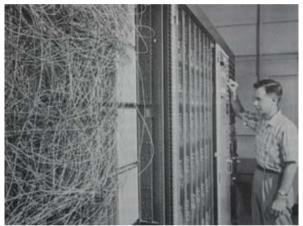
Optional: Ancient History

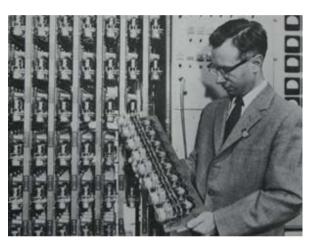
 The perceptron is a simple algorithm, but imagine coding it on this IBM 704, which Frank Rosenblatt used to implement it in 1957.

 There was much initial enthusiasm. But, it was later realized there were serious limitations, such as the linear separability requirement.

Optional: Dedicated Hardware (1960)







Imagine coding on that!



Optional: Python Implementation (1)

```
def perceptronRand(xs,ys,nIt=200,randP=True):
  N, D = xs.shape
                                 # Get data shape.
  w = np.zeros(D)
                                 # Init weights.
  for it in range(nIt):
                                 # Train.
    allCorrect = True
                                 # Generate indices.
    inds = np.random.permutation(N) if randP else np.arange(N)
    for i in inds:
                                  # Pick one sample.
      x = xs[i]
                                                                        Call to numpy. Mostly
                                                                        coded in C or Fortran.
       y = 2*(np.inner(x,w) > 0)-1 # Predict the label.
      if y != ys[i]:
                                  # Misclassified.
         w += vs[i] * x
                         # Update weights.
         w /= np.linalg.norm(w) # Normalize length.
         allCorrect = False
                                   # Something has changed.
    print('It {}: {}'.format(it + 1,linearAccuracy(xs, ys, w)))
    if allCorrect:
       break
                                   # Finish training.
  return w
def linearAccuracy(xs,ys,ws):
```

EPFL

return(sum(ys == (2 * (xs @ ws > 0)) - 1) * 100/len(ys))

Optional: Python Implementation (2)

```
def perceptronRand(xs,ys,nIt=200,randP=True):
  N, D = xs.shape
                                 # Get data shape.
                                  # Init weights.
  w = np.zeros(D)
  bestW = None
  bestA = 0.0
  for it in range(nIt):
                                 # Train.
    allCorrect = True
                                  # Generate indices.
    inds = np.random.permutation(N) if randP else np.arange(N)
    for i in inds:
                                                                         Record best solution.
       x = xs[i]
                                   # Pick one sample.
       y = 2*(np.inner(x,w) > 0)-1 # Predict the label.
                                   # Misclassified.
       if y != ys[i]:
         w += vs[i] * x
                                   # Update weights.
         w = np.linalg.norm(w)
                                   # Normalize length.
         allCorrect = False
                                    # Something has changed.
         acc = linearAccuracy(xs, ys, w)
         if(acc>bestA):
            bestW = w
            bestA = acc
    print('It {}: {}'.format(it + 1,bestA))
    if allCorrect:
       break
                                    # Finish training.
  return bestW
```



Optional: JAVA Implementation

```
import org.nd4j.linalg.api.ndarray.INDArray;
import org.nd4i.linalg.factory.Nd4i:
import java.lang.Float;
class Perceptron {
  public Perceptron() {}
  public static INDArray perceptronRand(INDArray xs, INDArray ys, int nlt, boolean randP){
     long[] shape = xs.shape();
                                                                     // Get data shape
     long N
                = shape[0];
     Iona D
                = shape[1]:
     INDArray w = Nd4j.zeros(D,1); // Init weights
     for (int it = 0: it < nlt: it++)\{
       boolean allCorrect = true;
       INDArray inds = Nd4j.arange(0,D);
                                                                      // Generate samples indices.
       if (randP)
          Nd4i.shuffle(inds);
       for (int i = 0; i < N; i++){
          INDArray x = xs.getRow(i);
                                                                       // Pick one sample.
          INDArray y = (x.mmul(w).gt(0)).mul(2).sub(1);
                                                                       // Predict the label.
          if (v.data().asFloat()[0] != vs.getRow(i).data().asFloat()[0]){ // Misclassified
            w = x.mul(ys.getRow(i)).add(w.transpose());
                                                                       // Update weights.
            w = w.div(w.norm2().add(1e-3)).transpose();
                                                                       // Unit normal length.
            allCorrect = false;
       System.out.println("It" + it + ":" + linearAccuracy(xs, vs, w));
       if (allCorrect){
          break;
     return w;
```

```
public static String linearAccuracy(INDArray xs,INDArray ys,INDArray w){
    INDArray y = (xs.mmul(w).gt(0)).mul(2).sub(1);
    return Nd4j.sum((y.eq(ys))).div(4).toString();
}

public class Main{
    public static void main (String[] args){

    INDArray xs = Nd4j.create(new float[][]{{1,0},{0,1},{1,1},{0,0}});
    INDArray ys = Nd4j.create(new float[][]{{1},{1},{1},{-1}});
    int nlt = 200;
    boolean randP = true;
    INDArray weights = Perceptron.perceptronRand(xs, ys, nlt, randP);
}
```

More verbose!



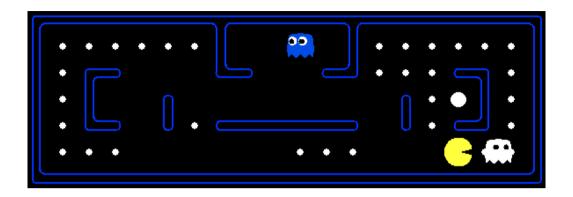
NumPy/SciPy

The time-critical loops are usually **implemented in C, C++ or Fortran**. Parts of SciPy are thin layers of code on top of the scientific routines that are freely available at http://www.netlib.org/. Netlib is a huge repository of incredibly valuable and robust scientific algorithms written in C and Fortran.

One of the design goals of NumPy was to make it buildable without a Fortran compiler, and if you don't have LAPACK available NumPy will use its own implementation. SciPy requires a Fortran compiler to be built, and **heavily depends on wrapped Fortran code**.



Optional: Pacman Apprenticeship



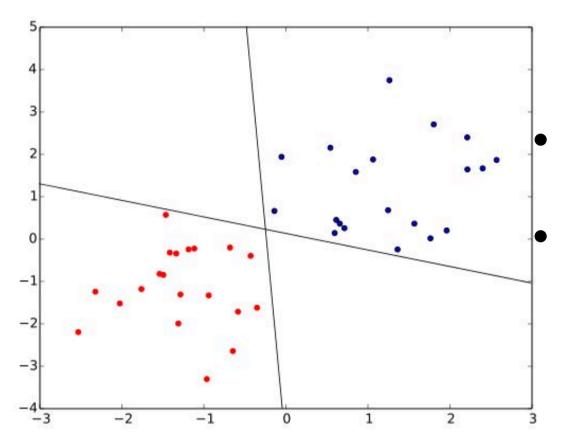
- Examples are state s.
- Correct actions are those taken by experts.
- Feature vectors defined over pairs φ(a,s).
- Score of a pair taken to be $\mathbf{w} \cdot \phi(a,s)$.
- Adjust w so that

$$\forall a, \mathbf{w} \cdot \phi(a^*, s) \ge \mathbf{w} \cdot \phi(a, s)$$

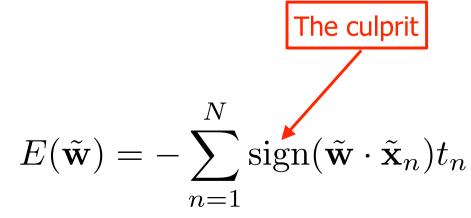
when a* is the correct action for state s.



The Problem with the Perceptron

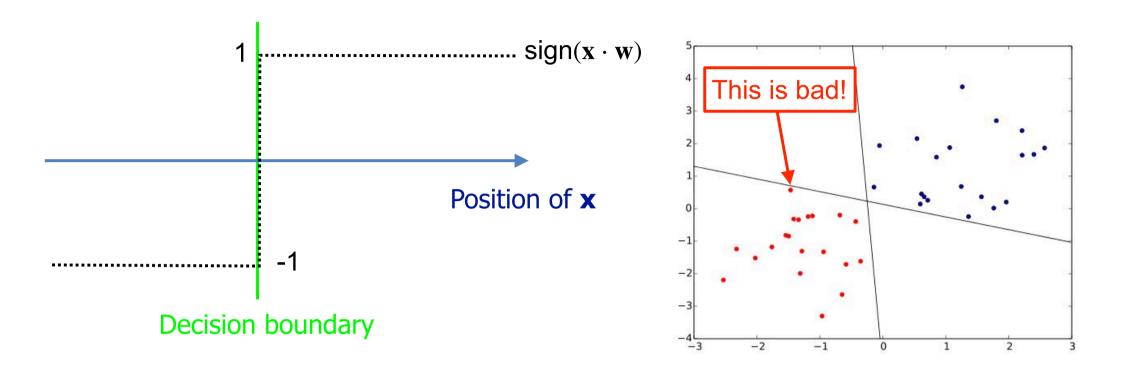


- Two different solutions among infinitely many.
- The perceptron has no way to favor one over the other.





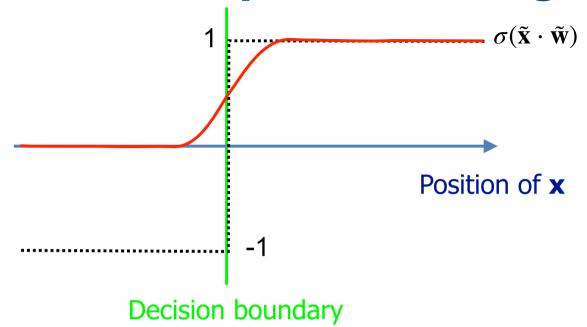
The Problem with the Perceptron



- There is no difference between close and far from the decision boundary.
- We want the positive and negative examples to be as far as possible from it.



From Perceptron to Logistic Regression



Replace the step function (black) by a smoother one (red).

- Replace the step function by a smooth function σ .
- The prediction becomes $y(\mathbf{x}; \widetilde{\mathbf{w}}) = \sigma(\widetilde{\mathbf{w}} \cdot \widetilde{\mathbf{x}})$.
- Given the training set $\{(\mathbf{x}_n, t_n)_{1 \leq n \leq N}\}$ where $t_n \in \{0, 1\}$, minimize the cross-entropy

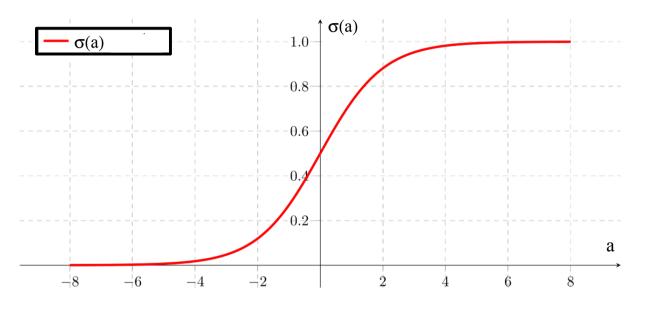
$$E(\widetilde{\mathbf{w}}) = -\sum_{n} \{t_n \ln y_n + (1 - t_n) \ln(1 - y_n)\}\$$
$$y_n = y(\mathbf{x}_n; \widetilde{\mathbf{w}})$$

This is a convex function of w!



with respect to $\tilde{\mathbf{w}}$.

Sigmoid Function



$$\sigma(a) = \frac{1}{1 + \exp(-a)}$$
$$\frac{\partial \sigma}{\partial a} = \sigma(1 - \sigma)$$

- It is infinitely differentiable.
- Its derivatives are easy to compute.
- It is asymptotically equal to zero or one.

—> Can be understood as a smoothed step function.



Cross Entropy

$$E(\tilde{\mathbf{w}}) = -\sum_{n} \{t_n \ln y_n + (1 - t_n) \ln(1 - y_n)\}$$

$$\nabla E(\tilde{\mathbf{w}}) = \sum_{n} (y_n - t_n) \tilde{\mathbf{x}}_n$$

$$y_n = \sigma(\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}}_n)$$

- $-(t_n \ln y_n + (1 t_n) \ln(1 y_n))$ is close to 0 if $t_n = 1$ and y_n is close to 1 or if $t_n = 0$ and y_n is close to zero. Minimizing $E(\mathbf{w})$ encourages that.
- $-(t_n \ln y_n + (1 t_n) \ln(1 y_n))$ is larger if $t_n = 1$ and $y_n < 0.5$ or $t_n = 0$ and $y_n > 0.5$. Minimizing E(w) discourages that.
- E(w) is a convex function whose gradient is easy to compute.
 - —> The global optimum can be found very effectively.



Probabilistic Interpretation

$$y(\mathbf{x}; \tilde{\mathbf{w}}) = \sigma(\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}})$$
$$= \frac{1}{1 + \exp(-\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}})}$$

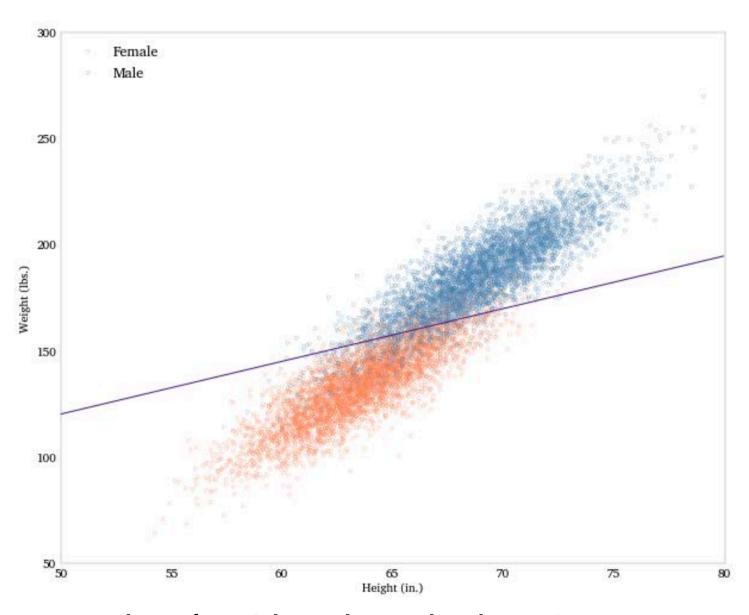
- $0 \le y(\mathbf{x}; \mathbf{w}) \le 1$
- $y(\mathbf{x}; \mathbf{w}) = 0.5$ if $\tilde{\mathbf{w}} \cdot \tilde{\mathbf{x}} = 0$, i.e. \mathbf{x} is on the decision boundary.
- $y(\mathbf{x}; \mathbf{w}) = 0.0$ or 1.0 if \mathbf{x} far from the decision boundary.

 \Rightarrow $y(\mathbf{x}; \tilde{\mathbf{w}})$ can be interpreted as the probability that x belongs to one class or the other.

Logistic regression finds what is called the **maximum likelihood solution** under the assumption that the noise is Gaussian.



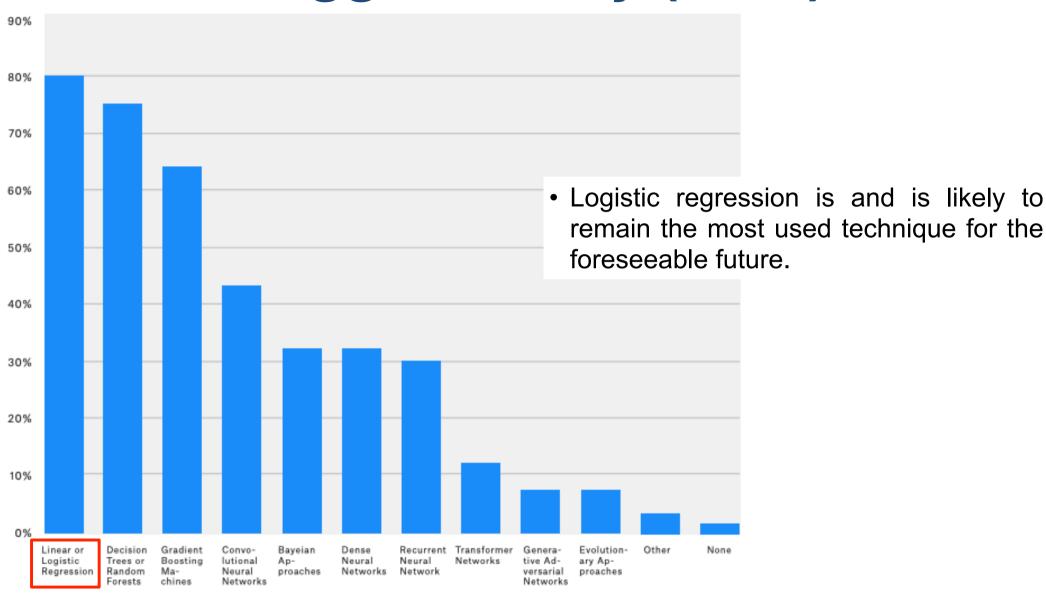
Example



- The algorithm does the best it can.
- Some samples can be misclassified.



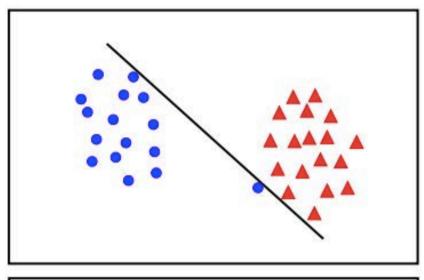
Kaggle Survey (2019)

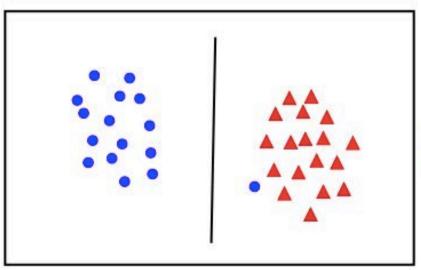


What data science methods do you use at work?



Outliers Can Cause Problems

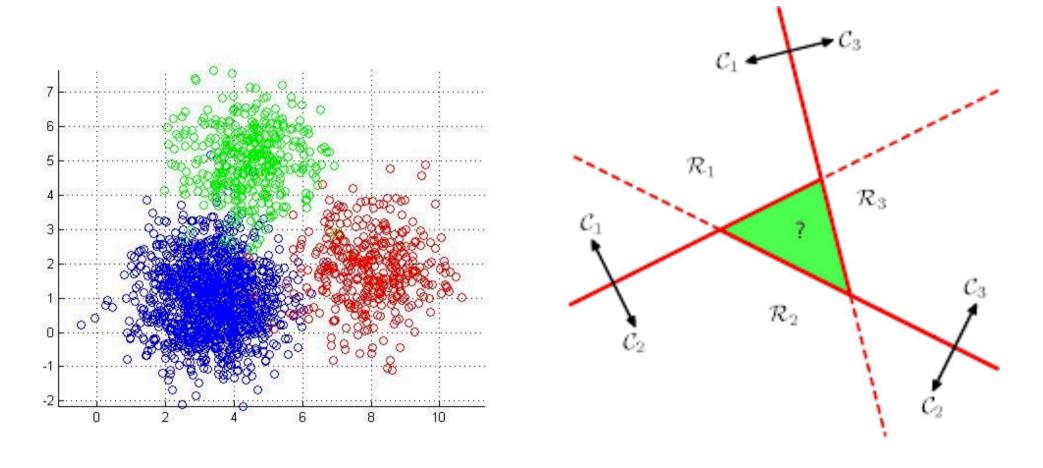




- Logistic regression tries to minimize the error-rate at training time.
- Can result in poor classification rates at test time.

—> We will talk about ways to prevent this in the next lecture.

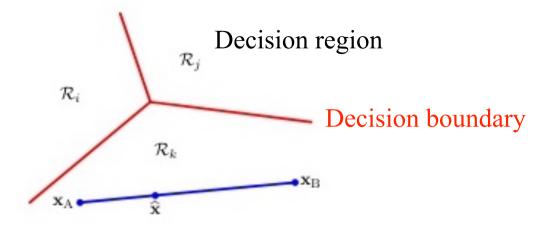
From Binary to Multi-Class



- k classes.
- Simply using k (k-1)/2 binary classifiers results in ambiguities.



Linear Discriminant



Given K linear classifiers of the form $y_k(\mathbf{x}) = \tilde{\mathbf{w}}_k \cdot \tilde{\mathbf{x}}$:

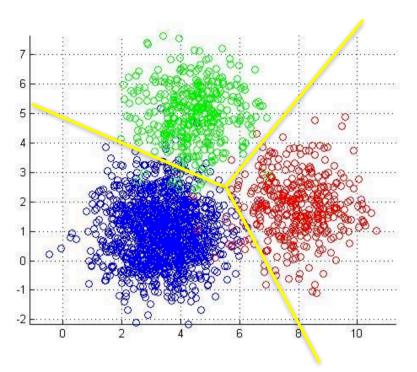
- Decision boundaries $y_k(\mathbf{x}) = y_l(\mathbf{x}) \Leftrightarrow (\tilde{\mathbf{w}}_k \tilde{\mathbf{w}}_l) \cdot \tilde{\mathbf{x}} = 0$.
- These boundaries define decision regions.
- Decision regions are convex:

$$\begin{split} (\tilde{\mathbf{w}}_k - \tilde{\mathbf{w}}_l) \cdot \tilde{\mathbf{x}}_A &> 0 \\ (\tilde{\mathbf{w}}_k - \tilde{\mathbf{w}}_l) \cdot \tilde{\mathbf{x}}_B &> 0 \\ \Rightarrow \forall \lambda \in [0,1], \text{ if } \mathbf{x} = \lambda \mathbf{x}_A + (1 - \lambda) \mathbf{x}_B, \text{ then} \\ (\tilde{\mathbf{w}}_k - \tilde{\mathbf{w}}_l) \cdot \tilde{\mathbf{x}} &> 0 \end{split}$$

In other words, if two points are on the same side of a decision boundary so are all point between them.



Multi-Class Linear Classification



$$k = \arg\max_{j} \tilde{\mathbf{w}}_{j}^{T} \tilde{\mathbf{x}}$$

$$\begin{bmatrix} y_1 \\ \vdots \\ y_K \end{bmatrix} = \begin{bmatrix} \tilde{\mathbf{w}}_1^T \\ \vdots \\ \tilde{\mathbf{w}}_K^T \end{bmatrix} \tilde{\mathbf{x}}$$
$$k = \arg\max y_i$$

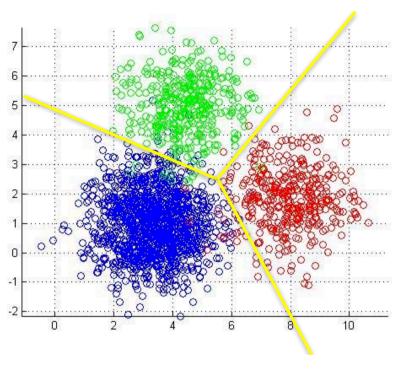
- K linear classifiers of the form $y^k(\mathbf{x}) = \tilde{\mathbf{w}}_k \cdot \tilde{\mathbf{x}} = \mathbf{w}_k^T \mathbf{x}$.
- Assign x to class k if $y^k(\mathbf{x}) > y^l(\mathbf{x}) \forall l \neq k$.

—> This still is a linear classification problem but in a space of dimension K times the dimension of the original one, 6 in this example.

Vector of dimension K times the dimension of $\tilde{\mathbf{w}}$.



Multi-Class Logistic Regression



$$k = \underset{j}{\operatorname{arg\,max}} \ y_k(\mathbf{x})$$

$$\begin{bmatrix} y_1 \\ \vdots \\ y_K \end{bmatrix} = \begin{bmatrix} \tilde{\mathbf{w}}_1^T \\ \vdots \\ \tilde{\mathbf{w}}_K^T \end{bmatrix} \tilde{\mathbf{x}}$$
$$k = \arg\max_j y_j$$

- K linear classifiers of the form $y^k(\mathbf{x}) = \sigma(\mathbf{w}_k^T \mathbf{x})$.
- Assign x to class k if $y^k(\mathbf{x}) > y^l(\mathbf{x}) \forall l \neq k$.

- Because the sigmoid function is monotonic, the formulation is almost unchanged.
- Only the objective function being minimized need to be reformulated.

Multi-Class Cross Entropy

Let the training set be $\{(\mathbf{x}_n, [t_n^1, ..., t_n^K])_{1 \le n \le N}\}$ where $t_n^k \in \{0, 1\}$ is the probability that sample \mathbf{x}_n belongs to class k.

Activation:

$$a^k(\mathbf{x}) = \boldsymbol{\sigma}(\mathbf{w}_k^T \mathbf{x})$$

Probability that **x** belongs to class k:

$$y^{k}(\mathbf{x}) = \frac{\exp(a^{k}(\mathbf{x}))}{\sum_{j} \exp(a^{j}(\mathbf{x}))}$$

Multi-class entropy:

$$E(\tilde{\mathbf{w}}_1, ..., \tilde{\mathbf{w}}_K) = -\sum_n \sum_k t_n^k \ln(y^k(\mathbf{x}_n))$$

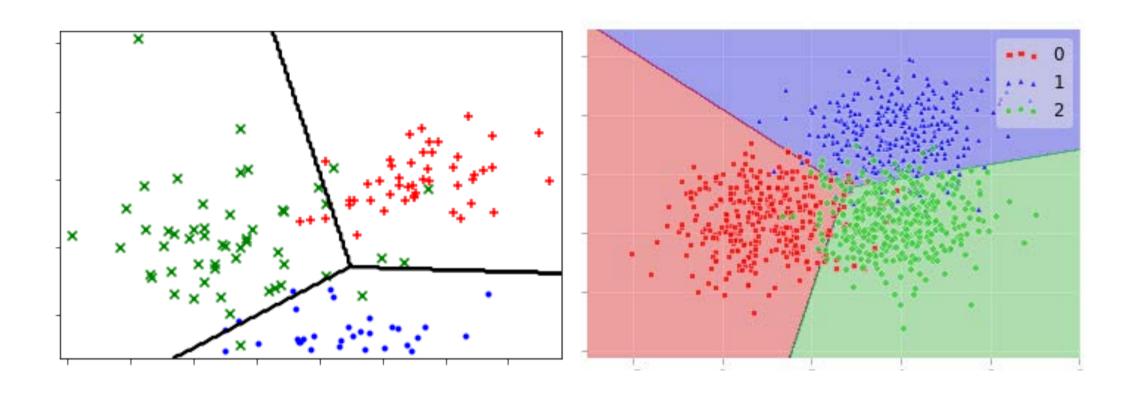
Gradient of the entropy:

$$\nabla E_{\mathbf{w_j}} = \sum_{n} (y^k(\mathbf{x}_n) - t_n^k) \mathbf{x}_n$$

- This is a natural extension of the binary case.
- The multi-class entropy is still convex and its gradient is easy to compute.



Multi-Class Results



Multiclass logistic regression is a very natural extension of binary logistic regression and has many of the same properties.

