

# Mathematical Background and the Perceptron

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#### ML Preliminaries

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One usually observes these variables for multiple "instances".

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One usually observes these variables for multiple "instances".

(Note) We use upper case to denote a random variable. To denote actual numbers we use lower case. One way to think about it: Y has not happened yet, and when it does, we see Y = Y.

#### ML Preliminaries

One may be interested in various things:

- What effects do the covariates have on the outcome?
- How well can we describe these effects?
- Can we predict the outcome using the covariates?, etc...

# Motivating Example: Credit Analysis

default	student	balance	income
No	No	729.5265	44361.625
No	Yes	817.1804	12106.135
No	No	1073.5492	31767.139
No	No	529.2506	35704.494
No	No	785.6559	38463.496
No	Yes	919.5885	7491.559

Task: predict account default What is the outcome Y? What are the predictors  $X_j$ ?

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We will denote predictors with x and outcomes with y (quantitative) and g (qualitative). Notice g are not numbers, so we cannot add or multiply them.

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Gender is a *qualitative measurement*. They are also called categorical or discrete. This is a particularly simple example because there are only two values. With two values we sometimes call it *binary*.

We will use G to denote the set of possible values. For gender it would be  $G = \{Male, Female\}$ .

A special case of qualitative variables are *ordered qualitative* where one can impose an order. With men/women this can't be done, but with, say,  $G = \{low, medium, high\}$  it can.

#### From data to feature vectors

The vast majority of ML algorithms we see in class treat instances as "feature vectors".

We can represent each instance as a *vector* in Euclidean space  $\langle x_1, \dots, x_p, y \rangle$ 

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#### This means:

- every measurement is represented as a continuous value
- in particular, categorical variables become numeric (e.g., one-hot encoding)

#### From data to feature vectors

Here is the same credit data represented as a matrix of feature vectors

default	student	balance	income
-1	1	685.5225	12361.86
-1	1	1647.0046	16614.80
1	1	2035.8616	14435.84
1	0	1903.6672	41173.50
1	1	1899.3906	20655.20
-1	0	1435.3601	33728.89

#### Technical notation

- Observed values will be denoted in lower case. So  $x_i$  means the ith observation of the random variable x.
- Matrices are represented with bold face upper case. For example  ${\bf x}$  will represent all observed predictors.

#### Technical notation

- N will usually mean the number of observations, or length of Y. i will be used to denote which observation and j to denote which covariate or predictor.
- Vectors will not be bold, for example  $x_i$  may mean all predictors for subject i, unless it is the vector of a particular predictor  $\mathbf{x}_j$ .
- All vectors are assumed to be column vectors, so the i-th row of  $\mathbf{x}$  will be  $x'_i$ , i.e., the transpose of  $x_i$ .

#### Geometry and Distances

Now that we think of instances as vectors we can do some interesting operations.

Let's try a first one: define a distance between two instances using Euclidean distance

$$d(x_1,x_2) = \sqrt{\sum_{j=1}^p (x_{1j} - x_{2j})^2}$$

# K-nearest neighbor classification

Now that we have a distance between instances we can create a classifier. Suppose we want to predict the class for an instance x.

K-nearest neighbors uses the closest points in predictor space predict *y*.

$$\hat{Y} = rac{1}{k} \sum_{x_k \in N_k(x)} y_k.$$

 $N_k(x)$  represents the k-nearest points to x. How would you use  $\hat{Y}$  to make a prediction?

```
function KNN-CLASSIFY(x, X, y, K)
     S \leftarrow []
                                                       \triangleright Compute distance to all points in X
     for all i = 1, \ldots, N do
         S \oplus \langle d(x,x_i),i \rangle
     end for
     S \leftarrow sort(S)
                                                                           \triangleright Find K nearest points
     \hat{y} \leftarrow 0
     for all k = 1, \dots, K do
          \langle d(x,x_i)\rangle \leftarrow S_k
         \hat{y} \leftarrow \hat{y} + y_i
                                                                                 ▶ Update prediction
     end for
     return sign(\hat{y})
                                                           \triangleright Return +1 if \hat{y} > 0, -1 otherwise
end function
```

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What assumptions we make about our data that allow us to make predictions.

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In KNN, our *inductive bias* is that points that are **nearby** will be of the same class.

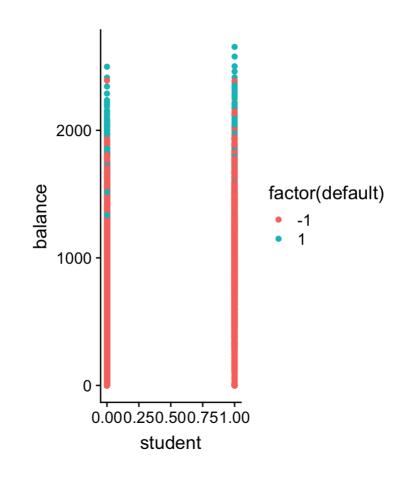
Parameter  $\kappa$  is a *hyper-parameter*, it's value may affect prediction accuracy significantly.

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Feature scaling is an important issue in distance-based methods.

Which of these two features will affect distance the most?

We will see in later lectures how to address this.



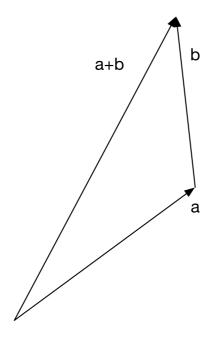
- A (real-valued) vector is just an array of real values, for instance  $x = \langle 1, 2.5, -6 \rangle$  is a three-dimensional vector.
- Vector sums are computed pointwise, and are only defined when dimensions match, so

$$\langle 1, 2.5, -6 
angle + \langle 2, -2.5, 3 
angle = \langle 3, 0, -3 
angle$$

.

In general, if c = a + b then cd = ad + bd for all vectors d.

Vector addition can be viewed geometrically as taking a vector a, then tacking on b to the end of it; the new end point is exactly c.



Scalar Multiplication: vectors can be scaled by real values;

$$2\langle 1,2.5,-6
angle = \langle 2,5,-12
angle$$

In general,  $ax = \langle ax_1, ax_2, \dots, ax_p 
angle$ 

The norm of a vector x, written ||x|| is its length.

Unless otherwise specified, this is its Euclidean length, namely:

$$\|x\| = \sqrt{\sum_{j=1}^p x_j^2}$$

Quiz

Write Euclidean distance of vectors u and v as a vector norm

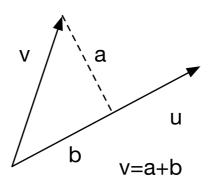
The *dot product*, or *inner product* of two vectors u and v is defined as

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$$u'v = \sum_{j=1}^p u_i v_i$$

A useful geometric interpretation of the inner product v'u is that it gives the projection of v onto u (when ||u|| = 1).



#### Back to KNN classification

The algorithm we saw scans the complete training set x to make a prediction for observation x.

This is not a good idea when the training set is massive.

#### Back to KNN classification

Suppose I give you function hash\_vector that does the following:

- Generate random unit vector u, ||u|| = 1
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- Round *a* to the nearest integer

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#### Quiz

Sketch a system that uses locality sensitive hashing (LSH) with hash\_vector to find candidate near neighbors of x to avoid computing distance to all vectors in dataset x.

# The curse of dimensionality

Distance-based methods like KNN can be problematic in highdimensional problems

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Consider the case where we have many covariates. We want to use k-nearest neighbor methods.

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Consider the case where we have many covariates. We want to use k-nearest neighbor methods.

Basically, we need to define distance and look for small multidimensional "balls" around the target points. With many covariates this becomes difficult.

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If we have p covariates and we are forming p-dimensional cubes, then each side of the cube must have size l determined by  $l \times l \times \cdots \times l = l^p = .10$ .

If the number of covariates is p=10, then  $l = .1^{1/10} = .8$ . So it really isn't local! If we reduce the percent of data we consider to 1%, l = 0.63. Still not very local.

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If we keep reducing the size of the neighborhoods we will end up with very small number of data points in each average and thus predictions with very large variance.

This is known as the curse of dimensionality.

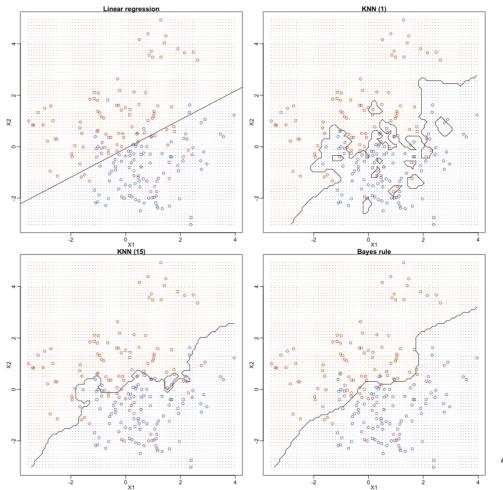
Because of this so-called curse, it is not always possible to use KNN. But other methods, like Decision Trees, thrive on multidimensional data.

Recall that our setting is that we observe for subject i predictors (covariates)  $x_i$ , and *qualitative* outcomes (or classes)  $g_i$ , which can takes values from a discrete set  $g_i$ .

Recall that our setting is that we observe for subject i predictors (covariates)  $x_i$ , and *qualitative* outcomes (or classes)  $g_i$ , which can takes values from a discrete set G.

Since our prediction  $\hat{g}(x)$  will always take values in the discrete set G, we can always divide the input space into a collection of regions taking the same predicted values.

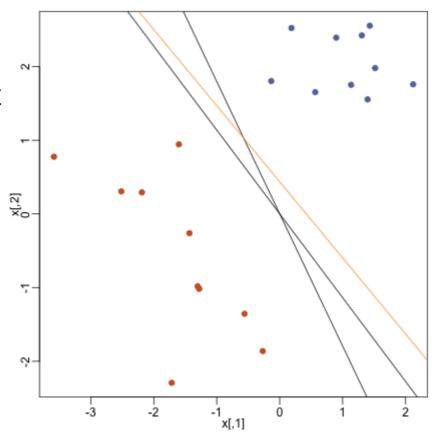
Boundaries can be smooth or rough depending on the prediction function.



For an important class of procedures, these *decision boundaries* are linear.

This is what we will refer to as linear methods for classification.

The *perceptron algorithm* is a way of finding discriminant functions that are *linear* with respect to the covariates  $X_1, ..., X_p$ .



In p-dimensional space  $\mathbb{R}^p$  these are described by vectors w. The *decision* boundary is thus

$$L=\{x:w'x=0\}.$$

Notice that this boundary partitions the input space into two sets on each side of the line.

If we restrict estimates to those for which |w| = 1

Then the *signed* distance of any point x to the decision boundary L is w'x.

With this we can easily describe the two partitions as

$$L^+ = \{x: w'x > 0\},$$

$$L^-=\{x:w'x<0\}$$

Intuitively, the w we want as an estimate is one that separates the training data as perfectly as possible.

If we code our classes as y = -1 if g = 1 and y = +1 if g = 2, we can describe our intuitive requirement for estimate w as:

$$y_i(w'x_i)>0, i=1,\dots,N$$

The *Perceptron* algorithm is one way of finding a vector w that satisfies the separation requirement as much as possible.

Penalize *w* by how far into the wrong side misclassified points are:

$$D(w) = -\sum_{i \in \mathcal{M}} y_i(w'x_i)$$

 $\mathcal{M}$ : set of points misclassified by w (on the wrong side of the hyper-plane).

The perceptron algorithm estimates w by *minimizing* D.

The perceptron algorithm estimates w by minimizing D. We'll see details how next class, but this introduces an important point:

We will write down learning algorithms as *optimization problems* where we *minimize* some *cost function* determined by prediction error.

```
function Perceptron(X,y,MaxIter)
    w \leftarrow \langle 0, \dots, 0 \rangle
    for all iter = 1, ..., MaxIter do
        for all i = 1, \ldots, N do
                                                               ▶ For every observation
            if y_i(w'x_i) < 0 then
                                                \triangleright Check if x_i is incorrectly classified
                w \leftarrow w + y_i x_i
            end if
        end for
    end for
    return w
end function
```

Why does this work?

Quiz

Show that if  $y_i(w'x_i) < 0$  then after updating w,  $w'x_i$  is changed in the proper direction. That is, it is made larger if  $y_i = +1$  and made smaller if  $y_i = -1$ .

#### Some Notes

- MaxIter is a hyper-parameter.
- You can interpret the size of  $w_j$  as the *importance* of feature j for classification

There are a few problems with this algorithm:

If there exists w that separates the training points perfectly,

There are a few problems with this algorithm:

If there exists w that separates the training points perfectly,

then there are an infinite number of ws that also separate the data perfectly

Algorithm will converge in a finite number of steps if the training data is separable

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However, the number of finite steps can be *very* large (see CIML for a bound on the number of steps)

When the training data is *not* separable, the algorithm will not converge.

### Summary

- We will represent many ML algorithms geometrically as vectors
- Vector math review
- K-nearest neighbors and perceptron algorithms