# Linear regression

Much of machine learning is about fitting functions to data. That may not sound like an exciting activity that will give us artificial intelligence. However, representing and fitting functions is a core building block of most working machine learning or AI systems. We start with linear functions, both because this idea turns out to be surprisingly powerful, and because it's a useful starting point for more interesting models and methods.

A general linear function of a vector  $\mathbf{x}$  takes a weighted sum of each input and adds a constant. For example, for D = 3 inputs  $\mathbf{x} = [x_1 \ x_2 \ x_3]^\top$ , a general (scalar) linear function is:

$$f(\mathbf{x}; \mathbf{w}, b) = w_1 x_1 + w_2 x_2 + w_3 x_3 + b = \mathbf{w}^{\top} \mathbf{x} + b,$$

where **w** is a *D*-dimensional vector of "weights". The constant offset or "bias weight" b gives the value of the function at x = 0 (the origin).

We will fit the function to a training set of N input-output pairs  $\{\mathbf{x}^{(n)}, \mathbf{y}^{(n)}\}_{n=1}^{N}$ . Expressing the data and function values using linear algebra, makes the maths easier in the long run, and makes it easier to write fast array-based code.

We stack all of the observed outputs into a column vector  $\mathbf{y}$ , and the inputs into an  $N \times D$  "design matrix" X:

$$\mathbf{y} = \begin{bmatrix} y^{(1)} \\ y^{(2)} \\ \vdots \\ y^{(N)} \end{bmatrix}, \qquad X = \begin{bmatrix} \mathbf{x}^{(1)\top} \\ \mathbf{x}^{(2)\top} \\ \vdots \\ \mathbf{x}^{(N)\top} \end{bmatrix} = \begin{bmatrix} x_1^{(1)} & x_2^{(1)} & \cdots & x_D^{(1)} \\ x_1^{(2)} & x_2^{(2)} & \cdots & x_D^{(2)} \\ \vdots & \vdots & \ddots & \vdots \\ x_1^{(N)} & x_2^{(N)} & \cdots & x_D^{(N)} \end{bmatrix}.$$

Elements of these arrays are  $y_n = y^{(n)}$ , and  $X_{n,d} = x_d^{(n)}$ . Each row of the design matrix gives the features for one training input vector. We can simultaneously evaluate the linear function at every training input with one matrix-vector multiplication:

$$\mathbf{f} = X\mathbf{w} + b$$

where 
$$f_n = f(\mathbf{x}^{(n)}; \mathbf{w}, b) = \mathbf{w}^{\top} \mathbf{x}^{(n)} + b$$
.

We can compute the total square error of the function values above, compared to the observed training set values:

$$E(\mathbf{w}, b) = \sum_{n=1}^{N} \left[ y^{(n)} - f(\mathbf{x}^{(n)}; \mathbf{w}, b) \right]^{2}$$
$$= (\mathbf{y} - \mathbf{f})^{\top} (\mathbf{y} - \mathbf{f}).$$

The least-squares fitting problem is finding the parameters that minimize this error.

## Fitting weights with b = 0

To keep the maths simpler, we will temporarily assume that we know our function goes through the origin. That is, we'll assume b = 0. Thus we are fitting the relationship:

$$\mathbf{y} \approx \mathbf{f} = X\mathbf{w}.$$

Fitting **w** to this approximate relationship by least-squares is so common that Matlab/Octave makes fitting the parameters astonishingly easy:

With NumPy, fitting the weights is still one line of code:

## Fitting more general functions

We'll return to how least squares fitting routines (\ or lstsq) work later. For now, we'll assume they're available, and see what we can do with them.

In machine learning it's often the case that the same code can solve different tasks simply by using it on different representations of our data. In the rest of this note and the next, we will solve different tasks all with the same linear least-squares primitive operation.

We assumed that our function passed through the origin. We can remove that assumption simply by creating a new version of our design matrix. We add an extra column containing a one in every row:

$$X' = \begin{bmatrix} 1 & \mathbf{x}^{(1)\top} \\ 1 & \mathbf{x}^{(2)\top} \\ \vdots & \vdots \\ 1 & \mathbf{x}^{(N)\top} \end{bmatrix} = \begin{bmatrix} 1 & x_1^{(1)} & x_2^{(1)} & \cdots & x_D^{(1)} \\ 1 & x_1^{(2)} & x_2^{(2)} & \cdots & x_D^{(2)} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & x_1^{(N)} & x_2^{(N)} & \cdots & x_D^{(N)} \end{bmatrix}.$$

In Matlab/Octave:

 $X_{bias} = [ones(size(X,1),1) X]; % Nx(D+1)$ 

In Python

$$X_{bias} = np.hstack([np.ones((X.shape[0],1)), X]) # (N,D+1)$$

Then we fit least squares weights exactly as before, just using X' or  $X_b$ ias, instead of the original design matrix X. If our input was D-dimensional before, we will now fit D+1 weights,  $\mathbf{w}'$ . The first of these is always multiplied by one, and so is actually the bias weight b, while the remaining weights give the regression weights for our original design matrix:

$$X'\mathbf{w}' = X\mathbf{w}'_{2:D+1} + w'_1 = X\mathbf{w} + b.$$

## **POLYNOMIALS**

We can go further, replacing the design matrix with a new matrix  $\Phi$ . Each row of this matrix is an arbitrary vector-valued function of the original input:  $\Phi_{n,:} = \phi(\mathbf{x}^{(n)})^{\top}$ . If the function is non-linear, then our function  $f(\mathbf{x}) = \mathbf{w}^{\top} \phi(\mathbf{x})$  will be non-linear in  $\mathbf{x}$ . However, we can still use linear-regression code to fit the model, as the model is still linear in the parameters.

The introductory example you'll see in most textbooks is fitting a polynomial curve to one-dimensional data. Each column of the new design matrix  $\Phi$  is a monomial of the original feature:

$$\Phi = \begin{bmatrix} 1 & x^{(1)} & (x^{(1)})^2 & \cdots & (x^{(1)})^{K-1} \\ 1 & x^{(2)} & (x^{(2)})^2 & \cdots & (x^{(2)})^{K-1} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & x^{(N)} & (x^{(N)})^2 & \cdots & (x^{(N)})^{K-1} \end{bmatrix}.$$

Using  $\Phi$  as our design or data matrix we can then fit the model

$$y \approx f = \mathbf{w}^{\top} \phi(x) = w_1 + w_2 x + w_3 x^2 + \dots + w_K x^{K-1}$$

which is a general polynomial of degree K-1.

We could generalize the transformation for multivariate inputs,

$$\phi(\mathbf{x}) = [1 \ x_1 \ x_2 \ x_3 \ x_1x_2 \ x_1x_3 \ x_2x_3 \ x_1^2 \ \dots]^\top,$$

and hence fit a multivariate polynomial function of our original features.

Polynomials are usually taught in introductions to regression first, because the idea of fitting a polynomial curve may already be familiar. However, polynomial fits are not actually used very often in machine learning. They're probably avoided for two reasons. 1) The feature space quickly explodes if your original features are high-dimensional; 2) Polynomials rapidly take on extreme values as the input x moves away from the origin.

## Basis functions

Instead of creating monomial features, we can transform our data with any other vectorvalued function:

$$\boldsymbol{\phi}(\mathbf{x}) = [\phi_1(\mathbf{x}) \ \phi_2(\mathbf{x}) \ \dots \ \phi_K(\mathbf{x})]^{\top}.$$

By convention, each  $\phi_k$  is called a *basis function*. The function we fit is a linear combination of these basis functions:

$$f(\mathbf{x}) = \mathbf{w}^{\top} \boldsymbol{\phi}(\mathbf{x}) = \sum_{k} w_{k} \phi_{k}(\mathbf{x}).$$

We don't have to include a constant or bias term in the mathematics, because we can always set one of the  $\phi_k$  functions to a constant.

One possible choice for a basis function  $\phi_k$  is a *Radial Basis Function* (RBF):

$$\exp(-(\mathbf{x}-\mathbf{c})^{\top}(\mathbf{x}-\mathbf{c})/h^2),$$

where different basis functions can have different parameters  $\mathbf{c}$  and h. The function is proportional to a Gaussian probability density function (although is not a probability density in this context). The function has a bell-curve shape centred at  $\mathbf{c}$ , with 'bandwidth' h. The bell-curve shape is radially symmetric: the function value only depends on the radial distance from the centre  $\mathbf{c}$ .

Another possible choice is a *logistic-sigmoid* function:

$$\sigma(\mathbf{v}^{\top}\mathbf{x} + b) = \frac{1}{1 + \exp(-\mathbf{v}^{\top}\mathbf{x} - b)}.$$

This sigmoidal or "s-shaped" curve saturates at zero and one for extreme values of  $\mathbf{x}$ . The parameters  $\mathbf{v}$  and b determine the steepness and position of the curve.

Using just linear regression, one line of fitting code, we can fit flexible regression models with many parameters. The trick is to construct a large design matrix, where each column corresponds to a basis function evaluated on each of the original inputs. Each basis function should have a different position and/or shape. Models built with radial-basis functions and sigmoidal functions extrapolate more conservatively than polynomials for extreme inputs. Radial-basis functions tend to zero, and sigmoidal functions tend to a constant. Neither of these families of basis functions has fundamental status however, and other basis functions are also used.

# Checking your understanding

A question to check your mathematical understanding:

• An RBF basis function is centred at some position **c**. We can create multiple columns of  $\Phi$  by evaluating RBFs with different centres. Why does it not make sense to create a family of quadratic functions  $\phi_k(x) = (x - c_k)^2$ , and include several features with different centres  $c_k$ ?

You should also understand how the maths would translate to code, and actually *doing* regression. I give here a quick review of how to plot functions in Matlab/Octave or Python, and demonstrate how to plot different basis functions, and linear regression fits.

## One dimensional basis functions

We can plot one-dimensional functions by evaluating them on a fine grid. For example, the cosine function:

In Matlab/Octave:

```
grid_size = 0.1;
     x_grid = -10:grid_size:10;
     f_{vals} = cos(x_{grid});
     clf; hold on;
     plot(x_grid, f_vals, 'b-');
     plot(x_grid, f_vals, 'r.');
In Python:
     import numpy as np
     import matplotlib.pyplot as plt
     grid_size = 0.1
     x_grid = np.arange(-10, 10, grid_size)
     f_{vals} = np.cos(x_{grid})
     plt.clf(); plt.hold('on')
     plt.plot(x_grid, f_vals, 'b-')
     plt.plot(x_grid, f_vals, 'r.')
     plt.show()
```

We're not really evaluating the whole function. We just evaluated it at the points shown by red dots. However, the blue curve joining these dots closely approximates our function. If we decrease the grid\_size further, the blue line will become as accurate as it's possible to plot on a computer screen.

We could use the cosine function as a basis function in our model. We can also create and plot other basis functions.

% Matlab/Octave functions usually need to be defined in a file.

Matlab/Octave:

```
% To make a simple RBF function, create a file rbf_1d.m containing:
     %function vals = rbf_1d(xx, cc, hh)
     \text{%vals} = \exp(-(xx-cc).^2 / hh.^2);
     % But one-line functions can be created anywhere like this:
     rbf_1d = @(xx, cc, hh) exp(-(xx-cc).^2 / hh.^2);
     clf(); hold on;
     grid_size = 0.01;
     x_grid = -10:grid_size:10;
     plot(x_grid, rbf_1d(x_grid, 5, 1), '-b');
     plot(x_grid, rbf_1d(x_grid, -2, 2), '-r');
Python:
     # Unlike Matlab, function definitions can be made anywhere
     # in Python. They don't have to go in a file.
     def rbf_1d(xx, cc, hh):
         return np.exp(-(xx-cc)**2 / hh**2)
     # An alternative that can be used for one line functions:
     \# rbf_1d = lambda xx, cc, hh: np.exp(-(xx-cc)**2 / hh**2)
     plt.clf(); plt.hold('on')
     grid_size = 0.01
     x_grid = np.arange(-10, 10, grid_size)
     plt.plot(x_grid, rbf_1d(x_grid, cc=5, hh=1), '-b')
     plt.plot(x_grid, rbf_1d(x_grid, cc=-2, hh=2), '-r')
     plt.show() # I may forget sometimes. Not necessary in python --pylab
```

The code above plots a radial basis function centred at x = 5 in blue, and one in red that's twice as wide and centred at x = -2.

**Exercise:** You should plot logistic-sigmoid functions  $\sigma(vx+b) = 1/(1+e^{-vx-b})$  in one-dimension for different v and b. What is the effect of the parameters v and b?

#### Combinations of basis functions

You might think of the function  $f(\mathbf{x}) = \sum_k w_k \phi_k(\mathbf{x})$  as being made up of separate parts  $\phi_k$ . For example, plot the function

$$f(x) = 2\phi_1(x) - \phi_2(x),$$

where  $\phi_1$  and  $\phi_2$  are unit bandwidth RBFs centred at -5 and +5. You can identify two separate bumps, each a scaled version of a basis function.

However, it's frequently not obvious by eye what the underlying basis functions are. For example, we can create a function with monomial basis functions

$$\phi_1(x) = 1$$

$$\phi_2(x) = x$$

$$\phi_3(x) = x^2$$

and weights  $\mathbf{w} = \begin{bmatrix} 5 & 10 & -1 \end{bmatrix}^{\mathsf{T}}$ . The resulting function has a simple form,

$$f(x) = \mathbf{w}^{\top} \phi(x) = 5 + 10x - x^2 = 30 - (x - 5)^2$$

which, unlike any of the basis functions, peaks at x = 5. Similarly, a function made by combining several overlapping RBFs can have peaks in between the peaks of any of the underlying basis functions.

## Linear regression

Consider a dataset with 3 datapoints:

$$\mathbf{y} = \begin{bmatrix} 1.1 \\ 2.3 \\ 2.9 \end{bmatrix}, \qquad X = \begin{bmatrix} 0.8 \\ 1.9 \\ 3.1 \end{bmatrix}.$$

By working through the following Matlab/Octave demo or otherwise, you should make sure you know how to fit and plot a variety of linear regression models.

```
% Set up and plot the dataset
yy = [1.1 2.3 2.9]';
X = [0.8 1.9 3.1]';
clf(); hold all;
plot(X, yy, 'x', 'MarkerSize', 20, 'LineWidth', 2);

% phi-functions to create various matrices of new features
% from an original matrix of 1D inputs.
phi_linear = @(Xin) [ones(size(Xin,1),1) Xin];
phi_quadratic = @(Xorig) [ones(size(Xorig,1),1) Xorig Xorig.^2];
fw_rbf = @(xx, cc) exp(-(xx-cc).^2 / 2);
phi_rbf = @(Xin) [fw_rbf(Xin, 1) fw_rbf(Xin, 2) fw_rbf(Xin, 3)];

fit_and_plot(phi_linear, X, yy); % Helper routine (see below)
fit_and_plot(phi_rbf, X, yy);
legend({'data', 'linear fit', 'quadratic fit', 'rbf fit'});
```

Using the following helper routine saved as fit\_and\_plot.m:

```
function fit_and_plot(phi_fn, X, yy)
% phi_fn takes Nx1 inputs and returns NxD basis function values
w_fit = phi_fn(X) \ yy; % Dx1
X_grid = (0:0.01:4)'; % Nx1
f_grid = phi_fn(X_grid)*w_fit;
plot(X_grid, f_grid, 'LineWidth', 2);
```

Different regression models give different fitted functions. Any model with three independent basis functions can fit the three training points exactly. However, the models' predictions in other locations will usually be different.

# Aside on terminology and notation

[This section is non-examinable. It's intended to help you read other resources.]

Textbooks and code documentation use various different names and notations for the things covered in this note. While learning a subject these differences can be confusing, but dealing with different notations is a necessary research skill.

The "weights" **w** are the parameters of the model, or "regression coefficients". In statistics textbooks and papers they are often given the symbol  $\beta$ . The constant offset or intercept b, when included, could have various symbols, including  $\beta_0$ ,  $w_0$ , or c. In the neural network literature this parameter is called the "bias weight" or simply the "bias". There's an unfortunate clash in terminology: a "bias weight" does not set the statistical "bias" of the model (its expected test error, averaged over different training sets).

These notes try where possible to use lower-case letters for indexing, with the corresponding capital letter for a number of settings. For example training data-points usually use the index n running from n=1...N, feature dimensions use d=1...D, and components of a model might be k=1...K. As a result, the notation won't match some textbooks. In statistics it's common to index data-items with i=1...n, and parameters with j=1...p.

While most textbooks use subscripts for both features and items, these notes usually identify the nth vector of a dataset with a bracketed super-script  $\mathbf{x}^{(n)}$ . Those superscripts occasionally get in the way, but make a clear distinction from feature dimensions, which are always subscripts. The dth feature of the nth datapoint is thus  $x_d^{(n)}$ . When we cover Gaussian processes later in the course, this notational baggage may help avoid confusion.

In statistics it's standard for the design matrix X to be an  $N \times D$  matrix, containing datapoints as row vectors. However machine learning code occasionally expects your data in a  $D \times N$  array. It is worth double-checking that your matrices are the correct way around when calling library routines, and leave comments in your own code to document the intended sizes of arrays. Sometimes I leave a comment at the end of a line that simply gives the size of the result: for example, "% NxD" in Matlab, or "# (N,D)" in Python.