

Lecture 4: Classification and Logistic Regression

Part 1: Classification

So far, every supervised learning algorithm that we've seen has been regression

We will next look at classification. First, let's define what classification is.

Review: Regression vs. Classification

Consider a training dataset $\mathcal{D} = \{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})\}.$

We distinguish between supervised learning problems depending on the targets $y^{(i)}$:

- 1. **Regression**: The target space is continuous: $\mathcal{Y} \subseteq \mathbb{R}$.
- 2. Classification: The target space is discrete: $\mathcal{Y} = \{y_1, y_2, \dots y_K\}$.

Each of the *K* discrete values corresponds to a *class* that we want to predict

An Example: Classifying Iris Flowers

Our running example for classification problems will be the iris flower dataset.

This is a classical dataset originally published by R. A. Fisher in 1936.

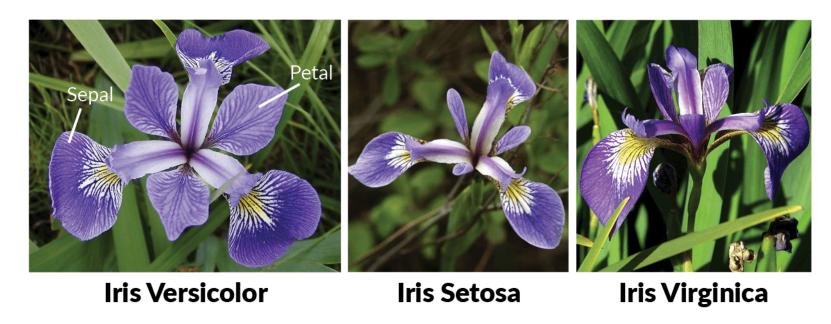


Image source: Machine Learning in R for Beginners

Let's import the dataset from sklearn.

```
import warnings; warnings.filterwarnings('ignore')
from sklearn import datasets
iris = datasets.load_iris(as_frame=True)
print(iris.DESCR)
.. _iris_dataset:
Iris plants dataset
**Data Set Characteristics:**
    :Number of Instances: 150 (50 in each of three classes)
    :Number of Attributes: 4 numeric, predictive attributes and the class
    :Attribute Information:
        sepal length in cm
        sepal width in cm
        - petal length in cm
        - petal width in cm
        - class:
               - Iris-Setosa
               - Iris-Versicolour
               - Iris-Virginica
    :Summary Statistics:
                                           Class Correlation
                   Min Max
                              Mean
                                             0.7826
    sepal length: 4.3 7.9 5.84 0.83
    sepal width:
                   2.0 4.4 3.05
                                     0.43
                                            -0.4194
```

Below, we print out five random rows of this dataset.

```
import pandas as pd
iris_X, iris_y = iris.data, iris.target
pd.concat([iris_X, iris_y], axis=1).sample(5) # sample 5 random rows from the data
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
1	4.9	3.0	1.4	0.2	0
104	6.5	3.0	5.8	2.2	2
46	5.1	3.8	1.6	0.2	0
13	4.3	3.0	1.1	0.1	0
112	6.8	3.0	5.5	2.1	2

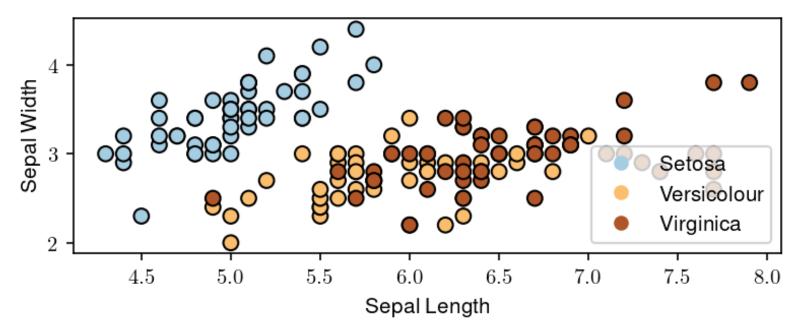
Understanding Classification

How is classification different from regression?

- **Regression**: fit a curve that passes close to the targets $y^{(i)}$.
- Classification: find boundaries separating the classes in the feature space
- Classification model outputs often have a simple **probabilistic interpretation**: they are probabilities that a data point belongs to a given class

Let's look at an example on the Iris dataset. We visualize this dataset using the first 2 attributes using matplotlib.

```
import matplotlib.pyplot as plt # Code from: https://scikit-learn.org/stable/auto_examples/datasets/plot_iris_dataset.html
plt.rcParams.update({ "figure.figsize": [6, 2], "figure.dpi": 150, "text.usetex": True, "font.family": "Helvetica"})
pl = plt.scatter(iris_X.iloc[:, 0], iris_X.iloc[:, 1], c=iris_y, edgecolor='k', s=50, cmap=plt.cm.Paired)
plt.xlabel('Sepal Length'); plt.ylabel('Sepal Width')
plt.legend(handles=p1.legend_elements()[0], labels=['Setosa', 'Versicolour', 'Virginica'], loc='lower right');
```



Let's train a classification algorithm called logistic regression.

```
from sklearn.linear_model import LogisticRegression
logreg = LogisticRegression(C=1e5)
X, Y = iris_X.to_numpy()[:,:2], iris_y.copy()
logreg.fit(X, Y);
```

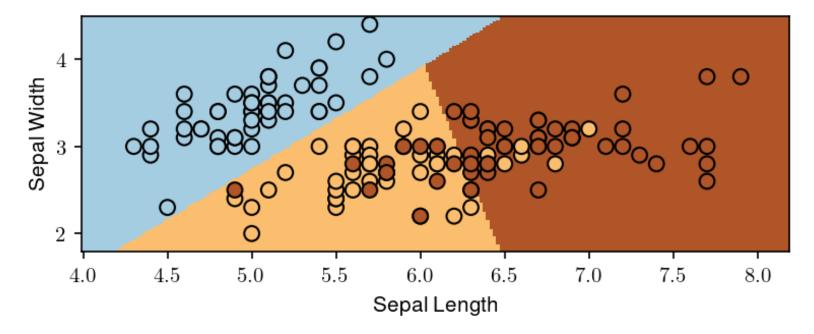
Note: logistic regression actually a classification algorithm.

The regions below are predicted to be associated with the blue, brown, and yellow classes. The lines between them are the decision boundaries.

```
import numpy as np
xx, yy = np.meshgrid(np.arange(4, 8.2, .02), np.arange(1.8, 4.5, .02))
Z = logreg.predict(np.c_[xx.ravel(), yy.ravel()])

# Put the result into a color plot
plt.pcolormesh(xx, yy, Z.reshape(xx.shape), cmap=plt.cm.Paired)

# Plot also the training points
plt.scatter(X[:, 0], X[:, 1], c=Y, edgecolors='k', cmap=plt.cm.Paired, s=50)
plt.xlabel('Sepal Length'); plt.ylabel('Sepal Width');
```





Part 2: Logistic Regression

We are now going to define our first classification algorithm: logistic regression.

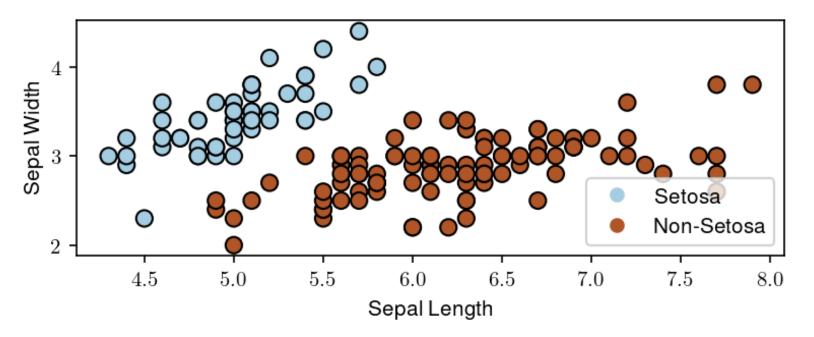
Binary Classification

- We will to start by looking at binary (two-class) classification.
- To keep things simple, we will use the Iris dataset.

We will attempt to distinguish class 0 (*Iris Setosa*) from the other two classes, using only the first two features in the dataset

```
iris_y2 = iris_y.copy()
iris_y2[iris_y2==2] = 1 # rename class two to class one

# plot also the training points
p1 = plt.scatter(iris_X.iloc[:, 0], iris_X.iloc[:, 1], c=iris_y2, edgecolor='k', s=60, cmap=plt.cm.Paired)
plt.xlabel('Sepal Length'); plt.ylabel('Sepal Width')
plt.legend(handles=p1.legend_elements()[0], labels=['Setosa', 'Non-Setosa'], loc='lower right');
```



Review: Ordinary Least Squares

Recall that in ordinary least squares, we fit a linear model of the form

$$f(x) = \sum_{j=0}^d heta_j \cdot x_j = heta^ op x.$$

It minimizes the mean squared error (MSE)

$$J(heta) = rac{1}{2n} \sum_{i=1}^n (y^{(i)} - heta^ op x^{(i)})^2$$

on a dataset $\{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})\}.$

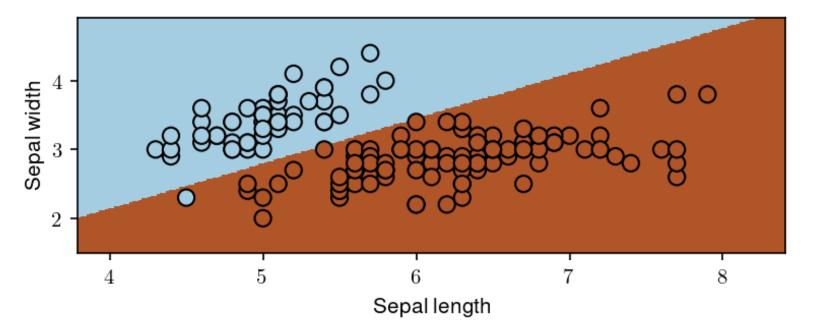
We could use least squares to solve our classification problem, setting $\mathcal{Y} = \{0, 1\}$.

```
from sklearn.linear_model import LinearRegression
linreg = LinearRegression()
X, Y = iris_X.to_numpy()[:,:2], iris_y2
linreg.fit(X, Y);
```

Ordinary least squares returns a decision boundary that is not unreasonable.

```
# Plot the decision boundary by assigning a color to each point in the mesh
x_min, x_max = X[:, 0].min() - .5, X[:, 0].max() + .5
y_min, y_max = X[:, 1].min() - .5, X[:, 1].max() + .5
h = .02 # step size in the mesh
xx, yy = np.meshgrid(np.arange(x_min, x_max, h), np.arange(y_min, y_max, h))
Z = linreg.predict(np.c_[xx.ravel(), yy.ravel()])
Z[Z>0.5] = 1.; Z[Z<0.5] = 0.

plt.pcolormesh(xx, yy, Z.reshape(xx.shape), cmap=plt.cm.Paired)
plt.scatter(X[:, 0], X[:, 1], c=Y, edgecolors='k', cmap=plt.cm.Paired, s=60)
plt.xlabel('Sepal length'); plt.ylabel('Sepal width');</pre>
```



However, applying OLS is problematic for a few reasons.

- Unbounded outputs: There is nothing to prevent outputs larger than one or smaller than zero, which is conceptually wrong
- Performance issues: At least one point is misclassified, and others are too close to the decision boundary.

Logistic Regression: The Model

Logistic regression is a classification algorithm where f_{θ} has the form

$$f_{ heta}(x) = \sigma(heta^ op x) = rac{1}{1 + \exp(- heta^ op x)}.$$

This is a composition of a linear model $\theta^{\top}x$ with

$$\sigma(z) = rac{1}{1 + \exp(-z)},$$

which is the sigmoid or logistic function.

The Logistic Function

To address the fact that the output of linear regression is not in [0,1], we will squeeze it into that range using

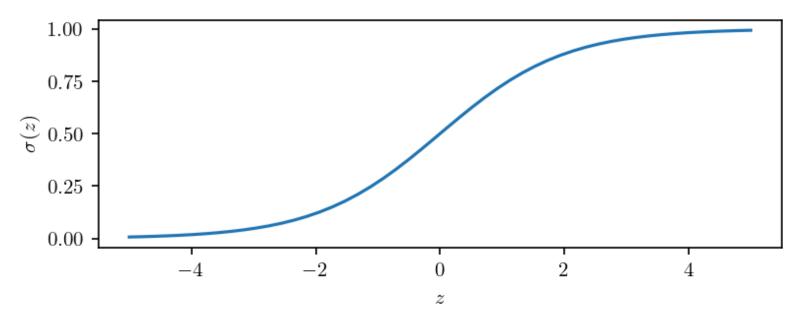
$$\sigma(z) = rac{1}{1 + \exp(-z)}.$$

This is known as the sigmoid or logistic function.

The logistic function $\sigma: \mathbb{R} \to [0,1]$ "squeezes" points from the real line into [0,1].

```
def sigmoid(z):
    return 1/(1+np.exp(-z))

z = np.linspace(-5, 5)
plt.plot(z, sigmoid(z))
plt.xlabel('$z$'); plt.ylabel(r'$\sigma(z)$');
```



The Logistic Function: Properties

The sigmoid function is defined as

$$\sigma(z) = rac{1}{1 + \exp(-z)}.$$

A few observations:

- The function tends to 1 as $z \to \infty$ and tends to 0 as $z \to -\infty$.
- Thus, models of the form $\sigma(\theta^{\top}x)$ output values between 0 and 1, which is suitable for binary classification.
- It is easy to show that the derivative of $\sigma(z)$ has a simple form: $\frac{d\sigma}{dz} = \sigma(z)(1 \sigma(z))$.

Probabilistic Interpretations

The logistic model can be interpreted to output a probability, and defines a conditional probability distribution as follows:

$$egin{aligned} P_{ heta}(y=1|x) &= \sigma(heta^ op x) \ P_{ heta}(y=0|x) &= 1 - \sigma(heta^ op x). \end{aligned}$$

Recall that a probability over $y \in \{0,1\}$ is called Bernoulli.



Part 3: Maximum Likelihood

In order to train a logistic regression model, we need to define an objective.

We derive this objective using the principle of maximum likelihood.

Recall: Supervised Learning Models

A model is a function f_{θ} that maps inputs $x \in \mathcal{X}$ to targets $y \in \mathcal{Y}$.

 $egin{aligned} \mathbf{Model \, Type} & \mathbf{How \, To \, Use} & \mathbf{Example} \ f_{ heta} : \underbrace{\mathcal{X}}_{ ext{input}}
ightarrow \underbrace{\mathcal{Y}}_{ ext{output}} & y_{ ext{pred}} = f_{ heta}(x) & y_{ ext{pred}} = heta^ op x \end{aligned}$

Probabilistic Supervised Learning Models

A probabilistic model parameterizes a probability over y_{pred} given x.

$$P_{ heta}(y=y_{ ext{pred}}|x)$$

Given $x \in \mathcal{X}$, the model outputs a *conditional probability* P(y|x) over targets $y \in \mathcal{Y}$.

For example, our logistic regression model f_{θ} defines ("parameterizes") a probability distribution $P_{\theta}(y|x)$ as follows:

$$P_{ heta}(y=1|x) = \sigma(heta^ op x) \ P_{ heta}(y=0|x) = 1 - \sigma(heta^ op x).$$

For each x, $P_{\theta}(y|x)$ is a valid probability over $y \in \{0,1\}$.

This is a more general recipe for a probabilistic model:

- Given an x, we have **a formula for** $P_{\theta}(y|x)$ as a function of x and θ .
- The formula assigns higher probabilities ("scores") to more likely y.
- These **scores sum to 1** over all y, hence they're valid probabilities.

We usually use the following notation to denote probabilistic models:

$$P_{ heta}(y|x): \mathcal{X}
ightarrow \underbrace{\left(\mathcal{Y}
ightarrow [0,1]
ight)}_{ ext{probability } P(y|x) ext{ over } \mathcal{Y}}.$$

Why Probabilistic Models?

There are two reasons why we will be interested in probabilistic models.

- Confidence Estimation: Oftentimes, we want to know if the model is certain in its prediction.
- Learning Objective: Probabilistic models can be used to define a principled learning objective.

Conditional Maximum Likelihood

We can train any model that defines a probability distribution $P_{\theta}(y|x)$ by optimizing the conditional maximum likelihood objective

$$\max_{ heta} \ell(heta) = \max_{ heta} rac{1}{n} \sum_{i=1}^n \log P_{ heta}(y^{(i)}|x^{(i)}).$$

defined over a dataset $\mathcal{D} = \{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})\}.$

We want to choose the weights such that the true label $y^{(i)}$ has the highest possible probability under $P_{\theta}(y|x^{(i)})$ for each $x^{(i)}$.

Example: Flipping a Random Coin

Consider a simple example in which we repeatedly toss a biased coin and record the outcomes.

- There are two possible outcomes: heads (H) and tails (T). A training dataset consists of tosses of the biased coin, e.g., $\mathcal{D} = \{H, H, T, H, T\}$
- **Assumption:** true probability distribution is $P_{\text{data}}(y)$, $y \in \{H, T\}$
- Our task is to determine the probability θ of seeing heads.

Example: Flipping a Random Coin

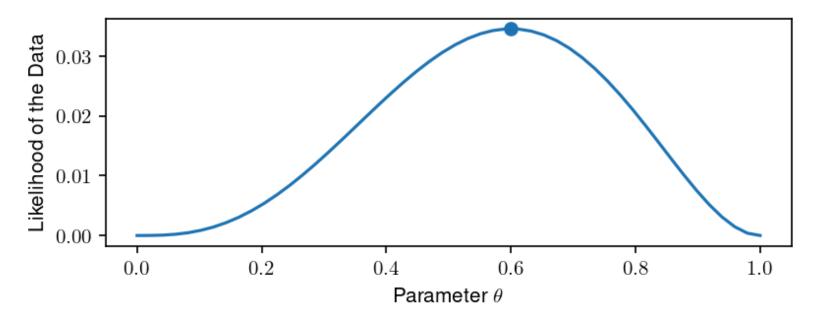
How should we choose θ if 3 out of 5 tosses are heads? Let's apply maximum likelihood:

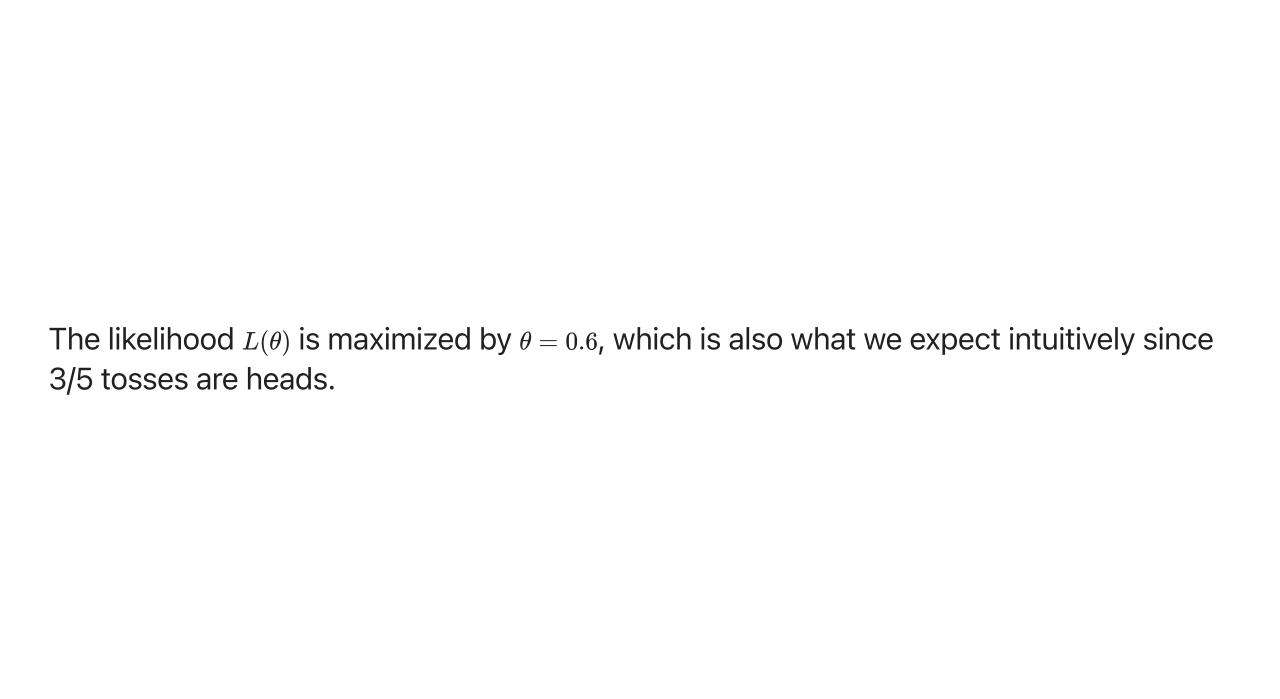
- Our dataset is $\mathcal{D} = \{y^{(1)}, y^{(2)}, y^{(3)}, y^{(4)}, y^{(5)}\} = \{H, H, T, H, T\}$
- Our model is $P_{\theta}(y) = \theta$ if y = H and $P_{\theta}(y) = 1 \theta$ if y = T, and there is a single parameter $\theta \in [0, 1]$
- The likelihood of the data is $L(\theta) = \prod_{i=1}^n P_{\theta}(y^{(i)}) = \theta \cdot \theta \cdot (1-\theta) \cdot \theta \cdot (1-\theta)$.

We optimize for θ which makes \mathcal{D} most likely. What is the solution?

```
# our dataset is {H, H, T, H, T}; if theta = P(x=H), we get:
coin_likelihood = lambda theta: theta*theta*(1-theta)*theta*(1-theta)

theta_vals = np.linspace(0,1)
plt.ylabel('Likelihood of the Data'); plt.xlabel(r'Parameter $\theta$')
plt.scatter([0.6], [coin_likelihood(0.6)])
plt.plot(theta_vals, coin_likelihood(theta_vals));
```







Part 4: Learning in Logistic Regression

We will use maximum likelihood to learn the parameters of a logistic regression model.

Logistic Regression

Recall that a logistic model defines ("parameterizes") a probability distribution $P_{\theta}(y|x): \mathcal{X} \times \mathcal{Y} \rightarrow [0,1]$ as follows:

$$P_{ heta}(y=1|x) = \sigma(heta^ op x) \ P_{ heta}(y=0|x) = 1 - \sigma(heta^ op x).$$

When $y \in \{0,1\}$, can write this more compactly as

$$P_{ heta}(y|x) = \sigma(heta^ op x)^y \cdot (1 - \sigma(heta^ op x))^{1-y}$$

Let's implement a logistic regression model in numpy.

```
def f(X, theta):
    """The sigmoid model we are trying to fit.

Parameters:
    theta (np.array): d-dimensional vector of parameters
    X (np.array): (n,d)-dimensional data matrix

Returns:
    y_pred (np.array): n-dimensional vector of predicted targets
    """
    return sigmoid(X.dot(theta))
```

Applying Maximum Likelihood

Following the principle of maximum likelihood, we want to optimize the following objective defined over a binary classification dataset

$$\mathcal{D} = \{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})\}.$$

$$egin{aligned} \ell(heta) &= rac{1}{n} \sum_{i=1}^n \log P_{ heta}(y^{(i)} \mid x^{(i)}) \ &= rac{1}{n} \sum_{i=1}^n \log \sigma(heta^ op x^{(i)})^{y^{(i)}} \cdot (1 - \sigma(heta^ op x^{(i)}))^{1 - y^{(i)}} \ &= rac{1}{n} \sum_{i=1}^n y^{(i)} \cdot \log \sigma(heta^ op x^{(i)}) + (1 - y^{(i)}) \cdot \log(1 - \sigma(heta^ op x^{(i)})). \end{aligned}$$

This objective is also often called the log-loss, or cross-entropy.

Observe that this objective asks the model to:

- Output a large score $\sigma(\theta^{\top}x^{(i)})$ (a score close to one) if $y^{(i)}=1$
- Output a score that's small (close to zero) if $y^{(i)} = 0$.

$$\ell(heta) = rac{1}{n} \sum_{i=1}^n y^{(i)} \cdot \log \sigma(heta^ op x^{(i)}) + (1-y^{(i)}) \cdot \log(1-\sigma(heta^ op x^{(i)})).$$

Let's implement the log-likelihood objective.

```
def log_likelihood(theta, X, y):
    """The cost function, J(theta0, theta1) describing the goodness of fit.

We added the 1e-6 term in order to avoid overflow (inf and -inf).

Parameters:
    theta (np.array): d-dimensional vector of parameters
    X (np.array): (n,d)-dimensional design matrix
    y (np.array): n-dimensional vector of targets
    """
    return (y*np.log(f(X, theta) + 1e-6) + (1-y)*np.log(1-f(X, theta) + 1e-6)).mean()
```

Review: Gradient Descent

If we want to minimize an objective $J(\theta)$, we may start with an initial guess θ_0 for the parameters and repeat the following update:

$$\theta_i := \theta_{i-1} - \alpha \cdot \nabla_{\theta} J(\theta_{i-1}).$$

Gradient of the Log-Likelihood

We want to use gradient descent to maximize the log-likelihood.

Hence our objective is $J(\theta) = -\ell(\theta)$

The gradient of the negative log-likelihood is

$$abla_{ heta} J(heta) =
abla_{ heta} \left[-\ell(heta)
ight] = rac{1}{n} \sum_{i=1}^n \left(\sigma(heta^ op x^{(i)}) - y^{(i)}
ight) \cdot x^{(i)}$$

Interestingly, this expression looks similar to the gradient of the mean squared error, which we derived in the previous lecture.

Let's implement the gradient.

```
def loglik_gradient(theta, X, y):
    """The cost function, J(theta0, theta1) describing the goodness of fit.

Parameters:
    theta (np.array): d-dimensional vector of parameters
    X (np.array): (n,d)-dimensional design matrix
    y (np.array): n-dimensional vector of targets

Returns:
    grad (np.array): d-dimensional gradient of the MSE
    """
    return np.mean((f(X, theta)-y) * X.T, axis=1)
```

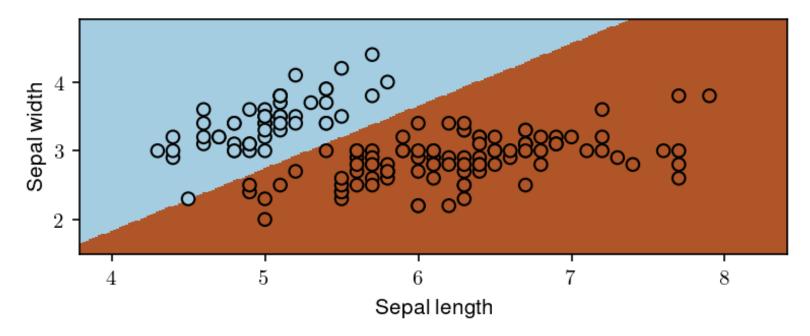
Let's now implement gradient descent.

```
threshold = 5e-5
step size = 1e-1
theta, theta_prev = np.zeros((3,)), np.ones((3,))
opt pts = [theta]
opt grads = []
iter = 0
iris_X['one'] = 1
X_{\text{train}} = iris_X.iloc[:,[0,1,-1]].to_numpy()
y_train = iris_y2.to_numpy()
while np.linalg.norm(theta - theta prev) > threshold:
    if iter % 50000 == 0:
        print('Iteration %d. Log-likelihood: %.6f' % (iter, log likelihood(theta, X train, y train)))
    theta prev = theta
    gradient = loglik_gradient(theta, X_train, y_train)
    theta = theta prev - step size * gradient
    opt_pts += [theta]
    opt grads += [gradient]
    iter += 1
```

```
Iteration 0. Log-likelihood: -0.693145
Iteration 50000. Log-likelihood: -0.021506
Iteration 100000. Log-likelihood: -0.015329
Iteration 150000. Log-likelihood: -0.012062
Iteration 200000. Log-likelihood: -0.010076
```

Let's now visualize the result.

```
xx, yy = np.meshgrid(np.arange(x_min, x_max, .02), np.arange(y_min, y_max, .02))
# predict using the learned model
Z = f(np.c_[xx.ravel(), yy.ravel(), np.ones(xx.ravel().shape)], theta)
Z[Z<0.5] = 0; Z[Z>=0.5] = 1
plt.pcolormesh(xx, yy, Z.reshape(xx.shape), cmap=plt.cm.Paired)
plt.scatter(X[:, 0], X[:, 1], c=Y, edgecolors='k', cmap=plt.cm.Paired)
plt.xlabel('Sepal length'); plt.ylabel('Sepal width');
```



Observations About Logistic Regression

Logistic regression finds a linear decision boundary.

This is the set of points for which P(y = 1|x) = P(y = 0|x).

Equivalently:

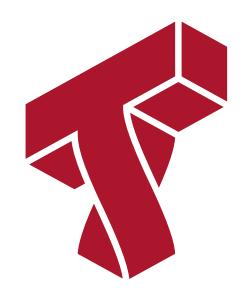
$$0 = \log rac{P(y=1|x)}{P(y=0|x)} = \log rac{rac{1}{1+\exp(- heta^ op x)}}{1-rac{1}{1+\exp(- heta^ op x)}} = heta^ op x$$

The set of x for which $0 = \theta^{T}x$ is a linear surface.

Unlike least squares, we don't have a closed form solution (a formula) for the optimal θ .
We can nonetheless find it numerically via gradient descent.

Algorithm: Logistic Regression

- Type: Supervised learning (binary classification)
- Model family: Linear decision boundaries.
- Objective function: Cross-entropy, a special case of log-likelihood.
- Optimizer: Gradient descent.
- Probabilistic interpretation: Parametrized Bernoulli distribution.



Part 5: Multi-Class Classification

Let's look at an extension of logistic regression to an arbitrary number of classes.

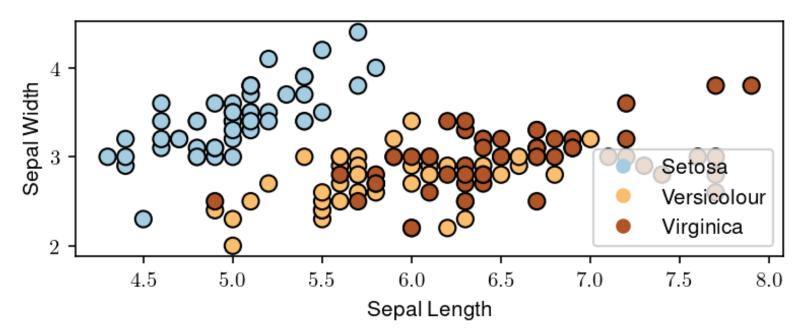
Multi-Class Classification

Logistic regression only applies to binary classification problems. What if we have an arbitrary number of classes K?

- The simplest approach is the "one vs. all" approach:

 Train one classifier for each class to distinguish that class from all the others.
- This works, but is not very elegant.
- Alternatively, we may fit a probabilistic model that outputs multi-class probabilities.

Let's load a fully multiclass version of the Iris dataset.



The Softmax Function

The logistic function $\sigma: \mathbb{R} \to [0,1]$ "squeezes" the score $z \in \mathbb{R}$ of a class into a probability in [0,1].

The softmax function $\vec{\sigma}: \mathbb{R}^K \to [0,1]^K$ is a multi-class version of σ

- It takes in a K-dimensional *vector* of class scores $\vec{z} \in \mathbb{R}$
- It "squeezes" \vec{z} into a length K vector of probabilities in $[0,1]^K$

The k-th component of the output of the softmax function $\vec{\sigma}$ is defined as

$$\sigma(ec{z})_k = rac{\exp(z_k)}{\sum_{l=1}^K \exp(z_l)}.$$

Softmax takes a vector of scores \vec{z} , exponentiates each score z_k , and normalizes the exponentiated scores such that they sum to one.

Softmax Regression: Model Class

Softmax regression is a multi-class classification algorithm which uses a model $f_{\theta}: \mathcal{X} \to [0,1]^K$ that generalizes logistic regression.

Softmax regression works as follows:

1. Given an input x, we compute K scores, one per class. The score

$$z_k = heta_k^ op x$$

of class k is a linear function of x and parameters θ_k for class k

1. We "squeeze" the vector of scores \vec{z} into $[0,1]^K$ using the softmax function $\vec{\sigma}$ and we output $\vec{\sigma}(\vec{z})$, a vector of K probabilities.

The parameters of this model are $\theta = (\theta_1, \theta_2, \dots, \theta_K)$, and the parameter space is $\Theta = \mathbb{R}^{K \times d}$.

The output of the model is a *vector* of class membership probabilities, whose k-th component $f_{\theta}(x)_k$ is

$$f_{ heta}(x)_k = \sigma(heta_k^ op x)_k = rac{\exp(heta_k^ op x)}{\sum_{l=1}^K \exp(heta_l^ op x)},$$

where each $\theta_l \in \mathbb{R}^d$ is the vector of parameters for class ℓ and $\theta = (\theta_1, \theta_2, \dots, \theta_K)$.

Softmax Regression: Probabilistic Interpretation

The softmax model outputs a vector of probabilities, and defines a conditional probability distribution as follows:

$$P_{ heta}(y=k|x) = ec{\sigma}(ec{z})_k = rac{\exp(heta_k^ op x)}{\sum_{l=1}^K \exp(heta_l^ op x)}.$$

Recall that a probability over $y \in \{1, 2, ..., K\}$ is called Categorical.

Softmax Regression: Learning Objective

We again maximize likelihood over a dataset \mathcal{D} .

$$egin{aligned} L(heta) &= \prod_{i=1}^n P_ heta(y^{(i)} \mid x^{(i)}) = \prod_{i=1}^n ec{\sigma}(ec{z}^{(i)})_{y^{(i)}} \ &= \prod_{i=1}^n \left(rac{\exp(heta_{y^{(i)}}^ op x^{(i)})}{\sum_{l=1}^K \exp(heta_l^ op x^{(i)})}
ight). \end{aligned}$$

We optimize this using gradient descent.

Let's now apply softmax regression to the Iris dataset by using the implementation from sklearn.

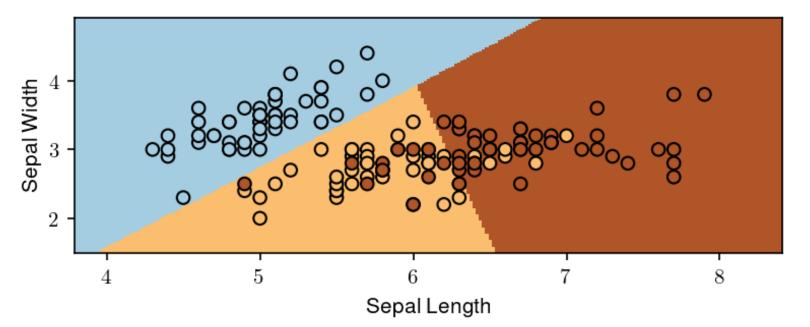
```
# https://scikit-learn.org/stable/auto_examples/linear_model/plot_iris_logistic.html
from sklearn.linear_model import LogisticRegression
logreg = LogisticRegression(C=1e5, multi_class='multinomial')

logreg.fit(X, iris_y) # Create an instance of Softmax and fit the data.
Z = logreg.predict(np.c_[xx.ravel(), yy.ravel()])

plt.pcolormesh(xx, yy, Z.reshape(xx.shape), cmap=plt.cm.Paired)

plt.scatter(X[:, 0], X[:, 1], c=iris_y, edgecolors='k', cmap=plt.cm.Paired)

plt.xlabel('Sepal Length'); plt.ylabel('Sepal Width');
```



Algorithm: Softmax Regression

- Type: Supervised learning (multi-class classification)
- Model family: Linear decision boundaries.
- Objective function: Softmax loss, a special case of log-likelihood.
- Optimizer: Gradient descent.
- Probabilistic interpretation: Parametrized categorical distribution.

Conclusion and Key Ideas

In summary, the key takeaways are:

- Do classification by taking a regression model and mapping the output into [0, 1]
 e.g., via sigmoid or softmax
- ML models usually have **probabilistic interpretations** as $P_{\theta}(y|x)$
- Any probabilistic model can be trained using maximum likelihood:

$$\max_{ heta} \sum_{i=1}^n P_{ heta}(y^{(i)}|x^{(i)})$$

Next class: regularization, further analysis