

Lecture 16: Unsupervised Learning

Applied Machine Learning

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Part 1: What is Unsupervised Learning?

Let's start by understanding what is unsupervised learning at a high level, starting with a dataset and an algorithm.

Unsupervised Learning

We have a dataset *without* labels. Our goal is to learn something interesting about the structure of the data:

- Clusters hidden in the dataset.
- Outliers: particularly unusual and/or interesting datapoints.
- Useful signal hidden in noise, e.g. human speech over a noisy phone.

Components of Unsupervised Learning

At a high level, an unsupervised machine learning problem has the following structure:

Dataset + Algorithm \rightarrow Unsupervised Model

The unsupervised model describes interesting structure in the data. For instance, it can identify interesting hidden clusters.

An Unsupervised Learning Dataset

As a first example of an unsupervised learning dataset, we will use our Iris flower example, but we will discard the labels.

We start by loading this dataset.

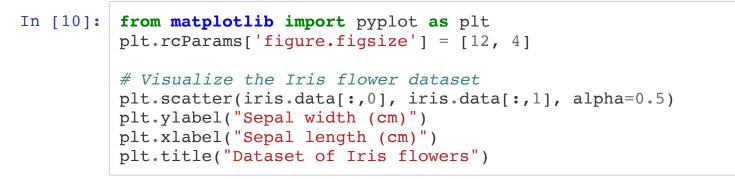
```
In [6]:
        # import standard machine learning libraries
        import numpy as np
        import pandas as pd
        from sklearn import datasets
        # Load the Iris dataset
        iris = datasets.load iris()
        print(iris.DESCR)
        .. iris dataset:
        Iris plants dataset
        **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:
```

Min Max Mean SD Class Correlation _____ ___ _____ ____ ____ 7.9 0.83 sepal length: 4.3 5.84 0.7826 3.05 sepal width: 2.0 4.4 0.43 -0.4194petal length: 0.9490 3.76 1.76 (high!) 1.0 6.9 petal width: 0.9565 0.1 2.5 1.20 0.76 (high!) :Missing Attribute Values: None :Class Distribution: 33.3% for each of 3 classes. :Creator: R.A. Fisher :Donor: Michael Marshall (MARSHALL%PLU@io.arc.nasa.gov) :Date: July, 1988 The famous Iris database, first used by Sir R.A. Fisher. The dataset is take n from Fisher's paper. Note that it's the same as in R, but not as in the UCI Machine Learning Repository, which has two wrong data points. This is perhaps the best known database to be found in the pattern recognition literature. Fisher's paper is a classic in the field an d is referenced frequently to this day. (See Duda & Hart, for example.) The data set contains 3 classes of 50 instances each, where each class refers to а type of iris plant. One class is linearly separable from the other 2; the latter are NOT linearly separable from each other. .. topic:: References - Fisher, R.A. "The use of multiple measurements in taxonomic problems" Annual Eugenics, 7, Part II, 179-188 (1936); also in "Contributions to Mathematical Statistics" (John Wiley, NY, 1950). - Duda, R.O., & Hart, P.E. (1973) Pattern Classification and Scene Analys is. (Q327.D83) John Wiley & Sons. ISBN 0-471-22361-1. See page 218. - Dasarathy, B.V. (1980) "Nosing Around the Neighborhood: A New System Structure and Classification Rule for Recognition in Partially Exposed

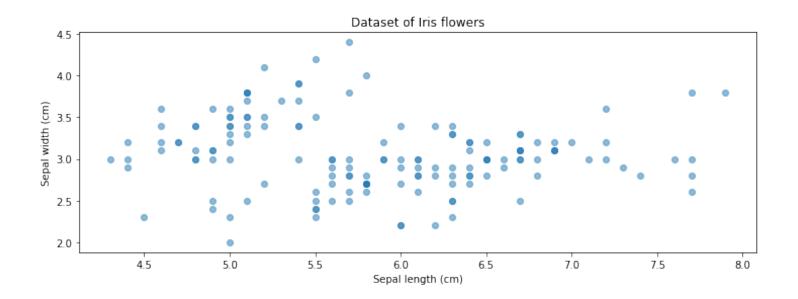
Environments". IEEE Transactions on Pattern Analysis and Machine Intelligence, Vol. PAMI-2, No. 1, 67-71.
Gates, G.W. (1972) "The Reduced Nearest Neighbor Rule". IEEE Transactions
on Information Theory, May 1972, 431-433.
See also: 1988 MLC Proceedings, 54-64. Cheeseman et al"s AUTOCLASS II conceptual clustering system finds 3 classes in the data.

- Many, many more ...

We can visualize this dataset in 2D. Note that we are no longer using label information.



Out[10]: Text(0.5, 1.0, 'Dataset of Iris flowers')



An Unsupervised Learning Algorithm

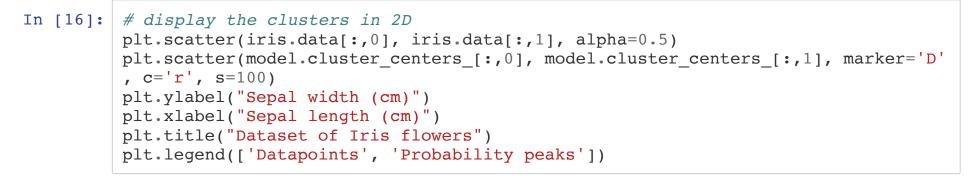
We can use this dataset as input to a popular unsupervised learning algorithm, K-means.

- The algorithm seeks to find *K* hidden clusters in the data.
- Each cluster is characterized by its centroid (its mean).
- The clusters reveal interesting structure in the data.

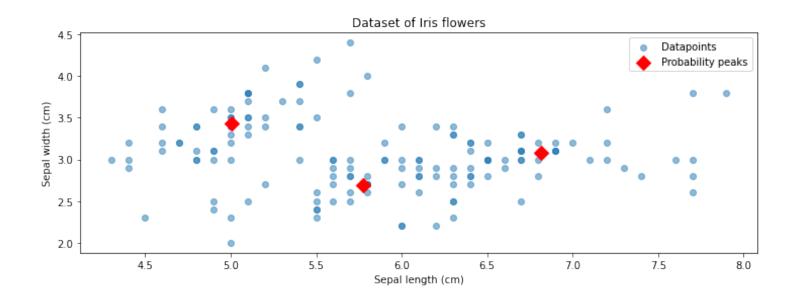
In [7]: # fit K-Means with K=3
from sklearn import cluster
model = cluster.KMeans(n_clusters=3)
model.fit(iris.data[:,[0,1]])

Out[7]: KMeans(n_clusters=3)

Running K-means on this dataset identifies three clusters.



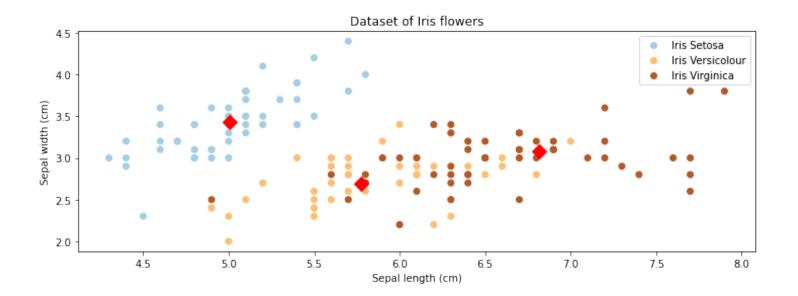
Out[16]: <matplotlib.legend.Legend at 0x1231df4a8>



These clusters correspond to the three types of flowers found in the dataset, which we obtain from the labels.

```
In [17]: p1 = plt.scatter(iris.data[:,0], iris.data[:,1], alpha=1, c=iris.target, cmap='P
aired')
plt.scatter(model.cluster_centers_[:,0], model.cluster_centers_[:,1], marker='D'
, c='r', s=100)
plt.ylabel("Sepal width (cm)")
plt.xlabel("Sepal length (cm)")
plt.title("Dataset of Iris flowers")
plt.legend(handles=p1.legend_elements()[0], labels=['Iris Setosa', 'Iris Versico
lour', 'Iris Virginica'])
```

Out[17]: <matplotlib.legend.Legend at 0x123247fd0>



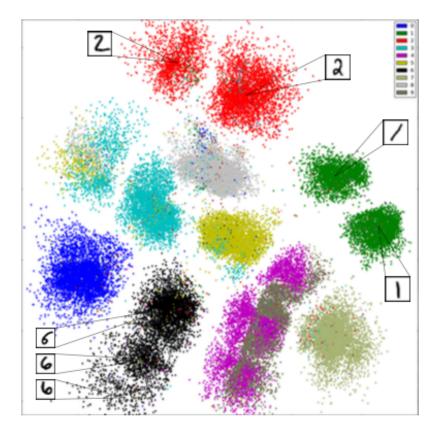
Applications of Unsupervised Learning

Unsupervised learning has numerous applications:

- Visualization: identifying and making accessible useful hidden structure in the data.
- Anomaly detection: identifying factory components that are likely to break soon.
- Signal denoising: extracting human speech from a noisy recording.

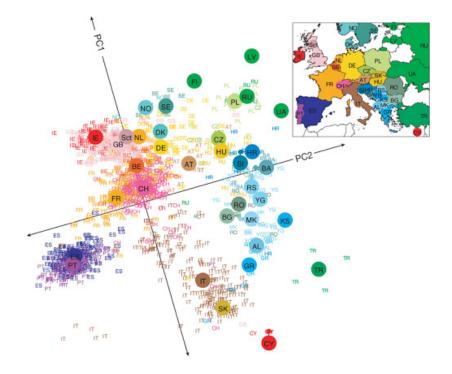
Application: Discovering Structure in Digits

Unsupervised learning can discover structure in digits without any labels.



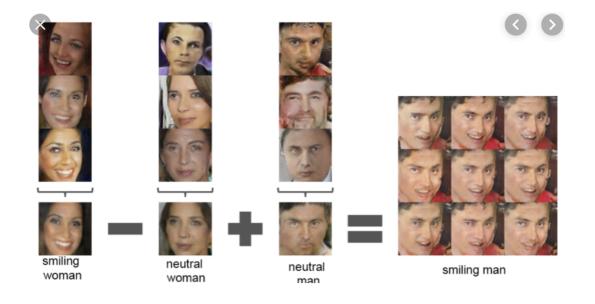
Application: DNA Analysis

Dimensionality reduction applied to DNA reveal the geography of European countries:



Application: Human Faces

Modern unsupervised algorithms based on deep learning uncover structure in human face datasets.



Unsupervised Learning in This Course

We will explore several types of unsupervised learning problems.

- Density estimation
- Anomaly and novelty detection
- Clustering
- Dimensionality Reduction

Next, we will start by setting up some notation.



Part 2: The Language of Unsupervised Learning

Next, let's look at how to define an unsupervised learning problem more formally.

Components of an Unsupervised Learning Problem

At a high level, an unsupervised machine learning problem has the following structure:

 $\underbrace{\text{Dataset}}_{\text{Attributes}} + \underbrace{\text{Learning Algorithm}}_{\text{Model Class + Objective + Optimizer}} \rightarrow \text{Unsupervised Model}$

The unsupervised model describes interesting structure in the data. For instance, it can identify interesting hidden clusters.

Unsupervised Dataset: Notation

We define of size n a dataset for unsupervised learning as $\mathcal{D} = \{x^{(i)} \mid i = 1, 2, \dots, n\}$

Each $x^{(i)} \in \mathbb{R}^d$ denotes an input, a vector of d attributes or features.

Data Distribution

We will assume that the dataset is sampled from a probability distribution \mathbb{P} , which we will call the *data distribution*. We will denote this as

 $x \sim \mathbb{P}$.

The dataset $\mathcal{D} = \{x^{(i)} \mid i = 1, 2, ..., n\}$ consists of independent and identicaly distributed (IID) samples from \mathbb{P} .

Data Distribution: IID Sampling

The key assumption in that the training examples are *independent and identicaly distributed* (IID).

- Each training example is from the same distribution.
- This distribution doesn't depend on previous training examples.

Example: Flipping a coin. Each flip has same probability of heads & tails and doesn't depend on previous flips.

Counter-Example: Yearly census data. The population in each year will be close to that of the previous year.

Components of an Unsupervised Learning Algorithm

We can think of a unsupervised learning algorithm as consisting of three components:

- A model class: the set of possible unsupervised models we consider.
- An **objective** function, which defines how good a model is.
- An **optimizer**, which finds the best predictive model in the model class according to the objective function

Model: Notation

We'll say that a model is a function

 $f: \mathcal{X} \to \mathcal{S}$ that maps inputs $x \in \mathcal{X}$ to some notion of structure $s \in \mathcal{S}$.

Structure can have many definitions (clusters, low-dimensional representations, etc.), and we will see many examples.

Often, models have parameters $\theta \in \Theta$ living in a set Θ . We will then write the model as $f_{\theta}: \mathcal{X} \to \mathcal{S}$

to denote that it's parametrized by θ .

Model Class: Notation

Formally, the model class is a set

$$\mathcal{M} \subseteq \{ f \mid f : \mathcal{X} \to \mathcal{S} \}$$

of possible models that map input features to structural elements.

When the models f_{θ} are paremetrized by *parameters* $\theta \in \Theta$ living in some set Θ . Thus we can also write

$$\mathcal{M} = \{ f_{\theta} \mid f : \mathcal{X} \to \mathcal{S}; \ \theta \in \Theta \}.$$

Objective: Notation

To capture this intuition, we define an *objective function* (also called a *loss function*) $J(f) : \mathcal{M} \to [0, \infty),$ which describes the extent to which f "fits" the data $\mathcal{D} = \{x^{(i)} \mid i = 1, 2, ..., n\}.$

When f is parametrized by $\theta \in \Theta$, the objective becomes a function $J(\theta) : \Theta \to [0, \infty)$.

Optimizer: Notation

An optimizer finds a model $f\in \mathcal{M}$ with the smallest value of the objective J. $\min_{f\in \mathcal{M}}J(f)$

Intuitively, this is the function that bests "fits" the data on the training dataset.

When f is parametrized by $\theta \in \Theta$, the optimizer minimizes a function $J(\theta)$ over all $\theta \in \Theta$.

An Example: *K*-Means

As an example, let's use the K-Means algorithm that we saw earlier.

Recall that:

- The algorithm seeks to find *K* hidden clusters in the data.
- Each cluster is characterized by its centroid (its mean).
- The clusters reveal interesting structure in the data.

The *K*-Means Model

We can think of the model returned by K-Means as a function $f_{\theta} : \mathcal{X} \to S$ that assigns each input x to a cluster $s \in S = \{1, 2, ..., K\}$.

The parameters θ of the model are *K* centroids $c_1, c_2, \ldots c_K \in \mathcal{X}$. The class of *x* is *k* if c_k is the closest centroid to *x*.

The *K*-Means Objective

How do we determine whether f_{θ} is a good clustering of the dataset \mathcal{D} ?

We seek centroids c_k such that the distance between the points and their closest centroid is minimized:

$$J(\theta) = \sum_{i=1}^{n} ||x^{(i)} - \text{centroid}(f_{\theta}(x^{(i)}))||,$$

where centroid(k) = c_k denotes the centroid for cluster k.

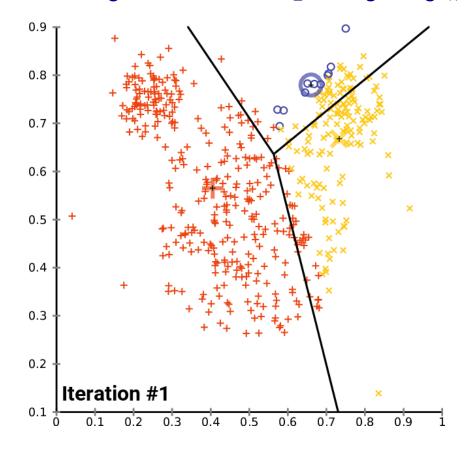
The *K*-Means Optimizer

We can optimize this in a two stop process, starting with an initial random cluster assignment f(x).

Repeat until convergence:

- 1. Set each c_k to be the center of the its cluster $\{x^{(i)} \mid f(x^{(i)}) = k\}$.
- 2. Update clustering f(x) such that $x^{(i)}$ is in the cluster of its closest centroid.

This is best illustrated visually (from <u>Wikipedia</u> (<u>https://commons.wikimedia.org/wiki/File:K-means_convergence.gif</u>)):



Algorithm: K-Means

- **Type**: Unsupervised learning (clustering)
- Model family: *k* centroids
- Objective function: Sum of distances to closest centroid
- **Optimizer**: Iterative optimization procedure.



Part 3: Unsupervised Learning in Practice

We will now look at some practical considerations to keep in mind when applying supervised learning.

Review: Data Distribution

We will assume that the dataset is sampled from a probability distribution \mathbb{P} , which we will call the *data distribution*. We will denote this as

$$x \sim \mathbb{P}$$
.

The dataset $\mathcal{D} = \{x^{(i)} \mid i = 1, 2, ..., n\}$ consists of independent and identicaly distributed (IID) samples from \mathbb{P} .

Review: Generalization

In machine learning, **generalization** is the property of predictive models to achieve good performance on new, heldout data that is distinct from the training set.

How does generalization apply to unsupervised learning?

Generalization in Unsupervised Learning

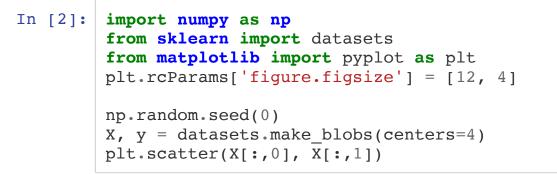
We can think of the data distribution as being the sum of two distinct components $\mathbb{P} = F + E$

- 1. A signal component F (hidden clusters, speech, low-dimensional data space, etc.)
- 2. A random noise component E

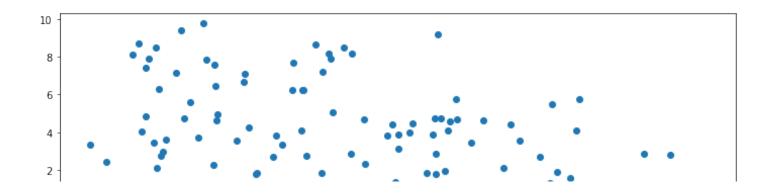
A machine learning model generalizes if it fits the true signal F; it overfits if it learns the noise E.

An Unsupervised Learning Dataset

Consider the following dataset, consisting of a mixture of Gaussians.



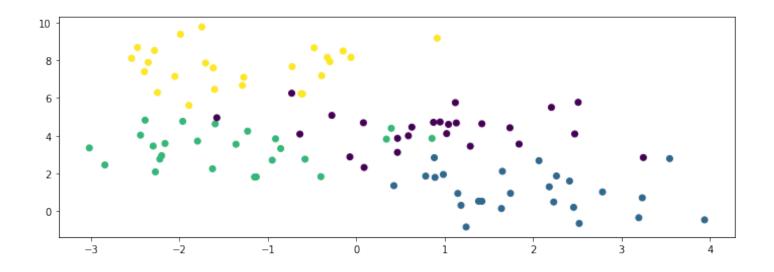
Out[2]: <matplotlib.collections.PathCollection at 0x120073b38>



We know the true labels of these clusers, and we can visualize them.



Out[30]: <matplotlib.collections.PathCollection at 0x122d99b00>



Underfitting in Unsupervised Learning

Underfitting happens when we are not able to fully learn the signal hidden in the data.

In the context of K-Means, this means not capturing all the clusters in the data.

Let's run *K*-Means on our toy dataset.

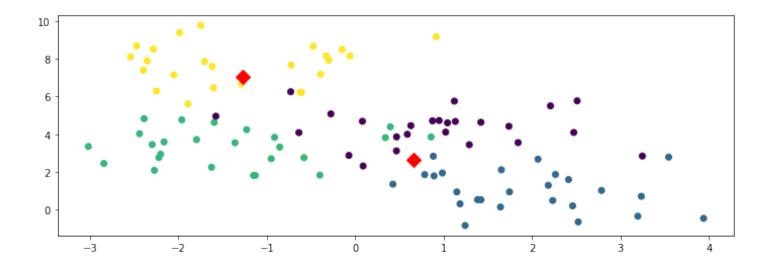
In [4]: # fit a K-Means
from sklearn import cluster
model = cluster.KMeans(n_clusters=2)
model.fit(X)

Out[4]: KMeans(n_clusters=2)

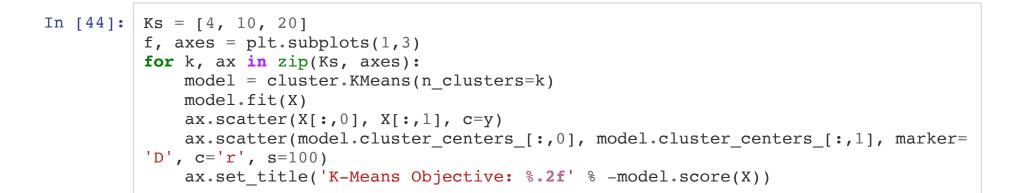
The centroids find two distinct components in the data, but they fail to capture the true structure.

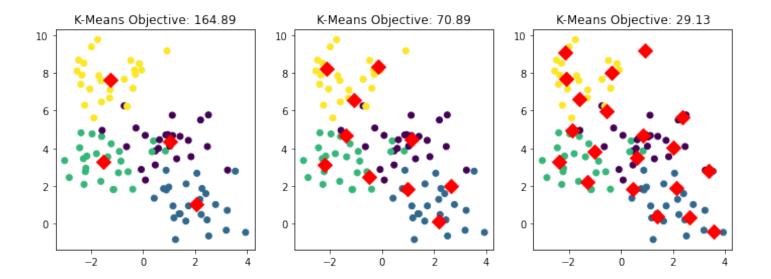
In [52]: plt.scatter(X[:,0], X[:,1], c=y)
plt.scatter(model.cluster_centers_[:,0], model.cluster_centers_[:,1], marker='D'
, c='r', s=100)
print('K-Means Objective: %.2f' % -model.score(X))

K-Means Objective: 462.03



Consider now what happens if we further increase the number of clusters.





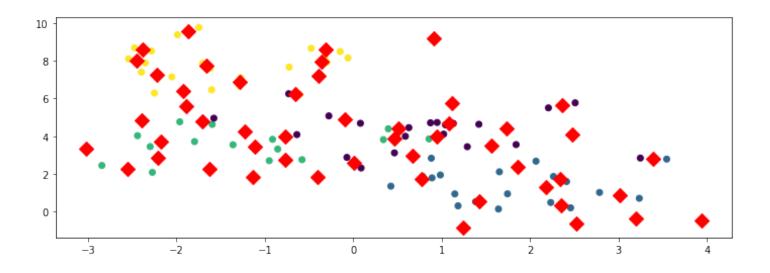
Overfitting in Unsupervised Learning

Overfitting happens when we fit the noise, but not the signal.

In our example, this means fitting small, local noise clusters rather than the true global clusters.

```
In [53]: model = cluster.KMeans(n_clusters=50)
model.fit(X)
plt.scatter(X[:,0], X[:,1], c=y)
plt.scatter(model.cluster_centers_[:,0], model.cluster_centers_[:,1], marker='D'
, c='r', s=100)
print('K-Means Objective: %.2f' % -model.score(X))
```

K-Means Objective: 4.94

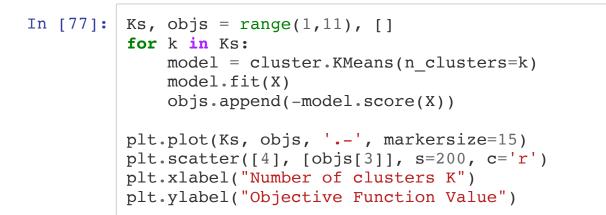


The Elbow Method

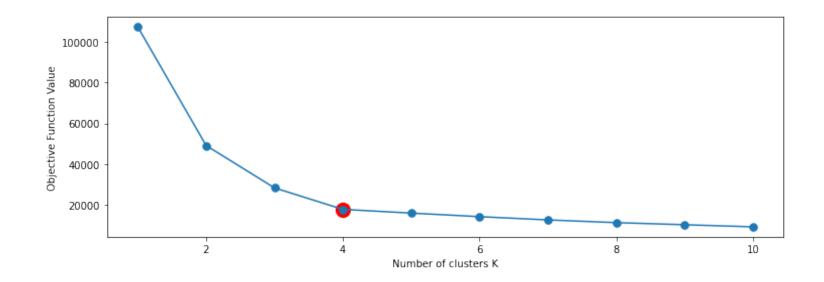
The Elbow method is a way of tuning hyper-parameters in unsupervised learning.

- We plot the objective function as a function of the hyper-parameter K.
- The "elbow" of the curve happens when its rate of decrease substantially slows down.
- The "elbow' is a good guess for the hyperparameter.

In our example, the decrease in objective values slows down after K = 4, and after that the curve becomes just a line.



Out[77]: Text(0, 0.5, 'Objective Function Value')



Detecting Overfitting and Underfitting

In unsupervised learning, overfitting and underfitting are more difficult to quantify than in supervised learning.

- Performance may depend on our intuition and require human evaluation
- If we know the true labels, we can measure the accuracy of the clustering

If our model is probabilistic, we can detect overfitting without labels by comparing the loglikelihood between the training set and a holdout set (next lecture!).

Reducing Overfitting

There are multiple ways to control for overfitting:

- 1. Reduce model complexity (e.g., reduce K in K-Means)
- 2. Penalize complexity in objective (e.g., penalize large K)
- 3. Use a probabilistic model and regularize it.

Summary

The concept of generalization applies to both supervised and unsupervised learning.

- In supervised learning, it is easier to quantify via the accuracy.
- In unsupervised learning, we may not be able to easily detect overfitting, but it still happens.

In []: