#### k-NN

CSC 461: Machine Learning

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## Instance-based learning

- ▶ Class of <u>learning methods</u>
  - ✓ also called **lazy learning**
- ▶ No need to learn any explicit hypothesis
- **Training** 
  - ✓ trivial, just need to store instances
- **Inference** 
  - ✓ time consuming, this is where computation happens

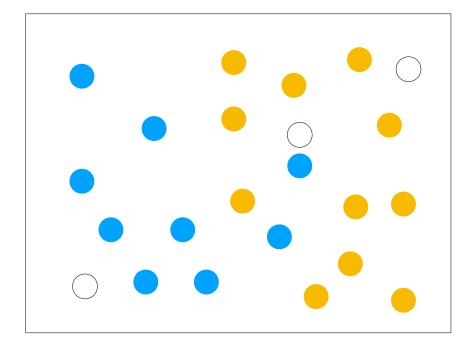
## Nearest neighbor classification

• Training examples are vectors with a class label

$$\mathbf{x_i} \in \mathbb{R}^d$$
  $y_i \in \{1, ..., C\}$ 

- ▶ Learning
  - **✓ store** all training examples
- Prediction
  - ✓ predict the label of the new example as the label of its **closest point** in the training set

what is the computational complexity of predicting a new label?



# k-Nearest Neighbors

## k-nearest neighbors

• Prediction for a new point x

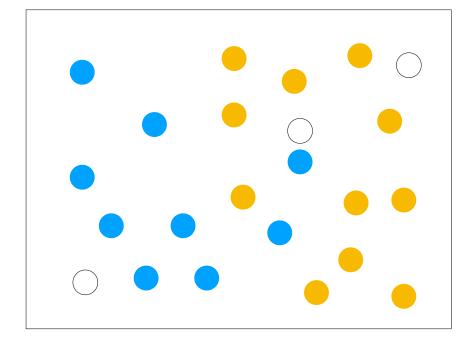
✓ recover a subset  $S_x$  (k nearest neighbors to x)

$$S_x \subseteq \mathcal{D}$$
 s.t.  $|S_x| = k$ 

$$\forall (\mathbf{x}', y') \in \mathcal{D} \setminus S_x$$
$$D(\mathbf{x}, \mathbf{x}') \ge \max_{(\mathbf{x}'', y'') \in S_x} D(\mathbf{x}, \mathbf{x}'')$$

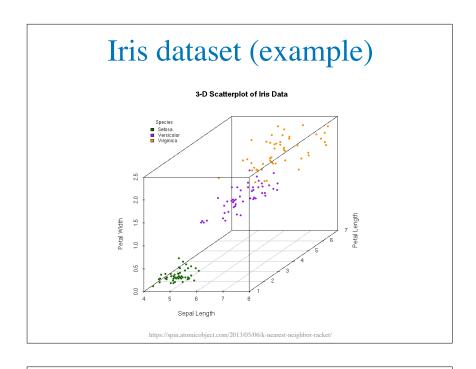
✓ take a majority vote (mode) (classification)

✓ calculate the **average** (<u>regression</u>)



## Iris dataset (example)

```
In [25]: from sklearn.datasets import load_iris
In [26]: data = load_iris()
                                                   sepal length, sepal width, petal
In [30]: data.data.shape
                                                   length, petal width and species
Out[30]: (150, 4)
In [27]: idx = np.random.choice(np.arange(len(data.target)), 10)
In [28]: data.target[idx]
Out[28]: array([1, 0, 0, 1, 1, 1, 2, 0, 0, 2])
In [29]: data.data[idx]
Out[29]:
array([[6.6, 3., 4.4, 1.4],
        [5. , 3.3, 1.4, 0.2], [4.7, 3.2, 1.3, 0.2],
        [6.3, 2.7, 4.9, 1.8]
        [4.6, 3.6, 1., 0.2],
                              https://en.wikipedia.org/wiki/Iris flower data set
```



#### Distance metrics

$$D(\mathbf{a}, \mathbf{b}) = \left(\sum_{i=1}^{d} |a_i - b_i|^p\right)^{1/p}$$
minkowski

 $\mathbf{a} \in \mathbb{R}^d, \mathbf{b} \in \mathbb{R}^d$ 

$$p = 1?$$
 manhattan

$$p = 2$$
? euclidean

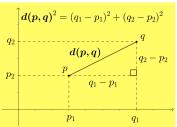
$$p = \infty$$
? chebyshev

$$\max_{i}(|a_{i}-b_{i}|)$$

#### More on the euclidean distance

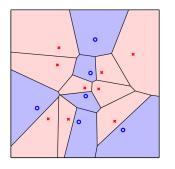
$$d(\mathbf{p}, \mathbf{q}) = \left(\sum_{i=1}^{d} |p_i - q_i|^2\right)^{1/2}$$
$$= \sqrt{\sum_{i=1}^{d} |p_i - q_i|^2}$$
$$= ||\mathbf{p} - \mathbf{q}||_2$$
$$= \sqrt{(\mathbf{p} - \mathbf{q}) \cdot (\mathbf{p} - \mathbf{q})}$$





## What is the decision boundary?

Is k-NN building an explicit decision boundary?
 ✓ not really, but it can be inferred

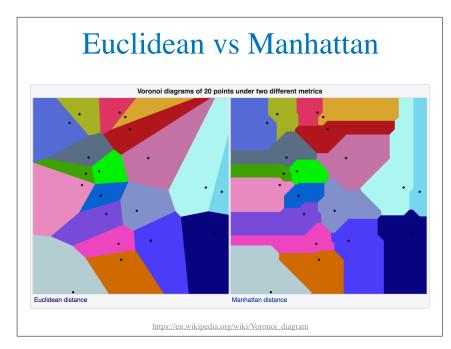


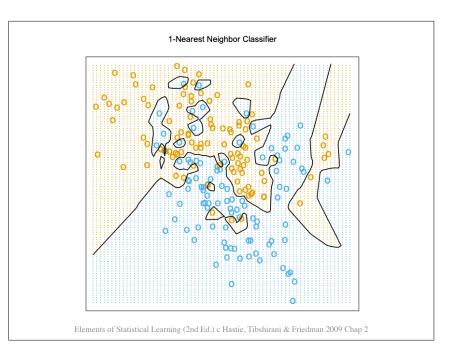
is the diagram sensitive to k?

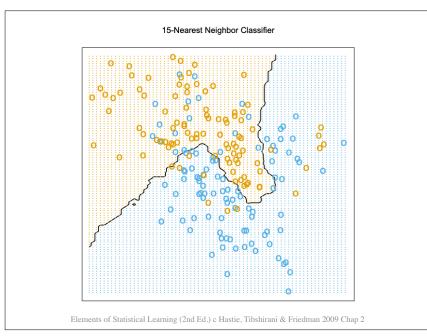
is the diagram sensitive to the distance function?

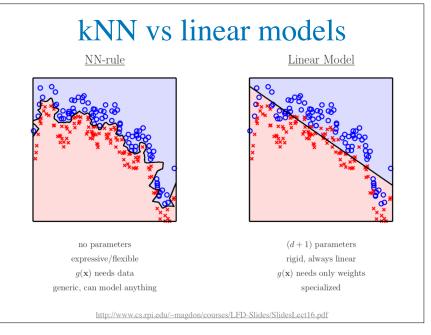
Nearest neighbor Voronoi tesselation

http://www.cs.rpi.edu/~magdon/courses/LFD-Slides/SlidesLect16.pdf





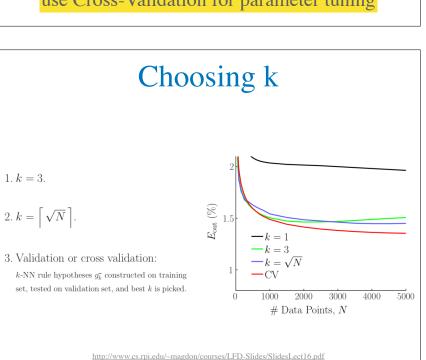


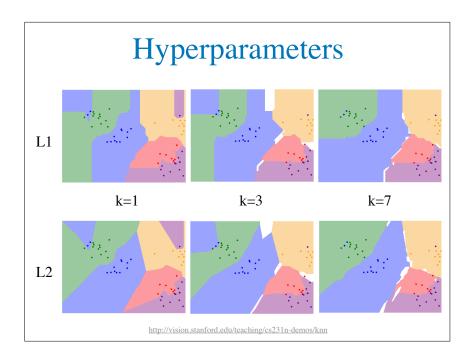


## Hyperparameters

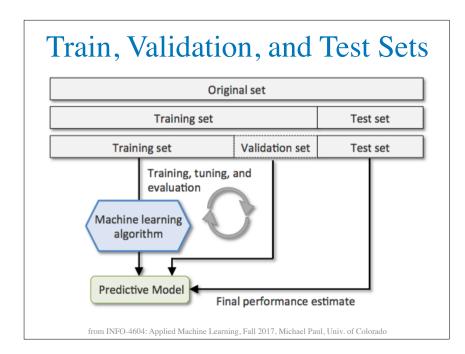
- ▶ The number of neighbors **k** 
  - ✓ too small, sensitive to noise
  - ✓ <u>too large</u>, neighborhood includes points from other classes
- **▶ Distance** function
- ▶ How to find a value that may generalize better?

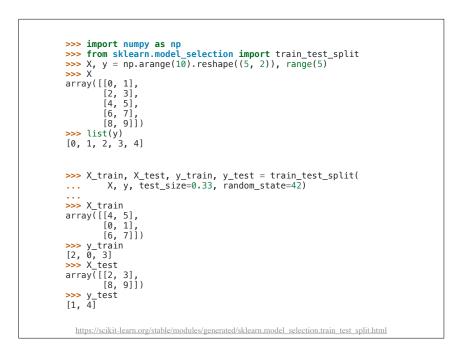
use Cross-Validation for parameter tuning











#### Parameters: \*arrays: sequence of indexables with same length / shape[0] Allowed inputs are lists, numpy arrays, scipy-sparse matrices or pandas dataframes. test\_size : float or int, default=None If float, should be between 0.0 and 1.0 and represent the proportion of the dataset to include in the test split. If int, represents the absolute number of test samples. If None, the value is set to the complement of the train size. If train\_size is also None, it will be set to 0.25. train\_size : float or int, default=None If float, should be between 0.0 and 1.0 and represent the proportion of the dataset to include in the train split. If int, represents the absolute number of train samples. If None, the value is automatically set to the complement of the test size. random\_state : int, RandomState instance or None, default=None Controls the shuffling applied to the data before applying the split. Pass an int for reproducible output across multiple function calls. See Glossary. shuffle : bool, default=True Whether or not to shuffle the data before splitting. If shuffle=False then stratify must be None. stratify : array-like, default=None If not None, data is split in a stratified fashion, using this as the class labels. Read more in the User Guide. https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.train\_test\_split.html

## Stratified split

#### Normalization

- ▶ k-NN can be sensitive to feature ranges
  - ✓ e.g., euclidean distance

$$D(\mathbf{a}, \mathbf{b}) = \sqrt{\sum_{i=1}^{d} (a_i - b_i)^2}$$

• Features can be preprocessed

✓ zero mean and unit variance

$$x_j' = \frac{x_j - \mu_j}{\sigma_j}$$



For certain datasets, the scale may be important

### Fit (train) and transform (test)

#### Normalization

- Must calculate parameters using training data
  - ✓ then transform the test data

```
>>> from sklearn import preprocessing
>>> import numpy as np
>>> X_train = np.array([[ 1., -1., 2.],
                         [ 2., 0., 0.],
[ 0., 1., -1.]])
>>> scaler = preprocessing.StandardScaler().fit(X_train)
>>> scaler
StandardScaler()
>>> scaler.mean_
array([1. ..., \overline{0}. ..., 0.33...])
>>> scaler.scale_
array([0.81..., \overline{0.81...}, 1.24...])
>>> X_scaled = scaler.transform(X_train)
>>> X scaled
array([[ 0. ..., -1.22..., 1.33...],
        [ 1.22..., 0. ..., -0.26...],
        [-1.22..., 1.22..., -1.06...]
```

#### sklearn.preprocessing: Preprocessing and Normalization

The sklearn.preprocessing module includes scaling, centering, normalization, binarization methods.

User guide: See the Preprocessing data section for further details.

```
preprocessing .Binarizer(*[, threshold, copy]) Binarize data (set feature values to 0 or 1) according to a threshold.
preprocessing.FunctionTransformer([func, ...])
Constructs a transformer from an arbitrary callable.
preprocessing.KBinsDiscretizer([n_bins, ...]) Bin continuous data into intervals.
preprocessing.KernelCenterer()
                                                   Center an arbitrary kernel matrix K.
preprocessing.LabelBinarizer(*[, neg_label, ...]) Binarize labels in a one-vs-all fashion.
preprocessing.LabelEncoder()
                                                   Encode target labels with value between 0 and n_classes-1.
preprocessing.MultiLabelBinarizer(*[, ...])
                                                  Transform between iterable of iterables and a multilabel format.
preprocessing.MaxAbsScaler(*[,copv])
                                                   Scale each feature by its maximum absolute value.
preprocessing.MinMaxScaler([feature_range, ...]) Transform features by scaling each feature to a given range
preprocessing.Normalizer([norm, copy])
                                                   Normalize samples individually to unit norm.
preprocessing. OneHotEncoder(*[, categories, ...]) Encode categorical features as a one-hot numeric array
preprocessing.OrdinalEncoder(*[, ...])
                                                   Encode categorical features as an integer array.
preprocessing.PolynomialFeatures([degree, ...]) Generate polynomial and interaction features.
preprocessing. PowerTransformer([method, ...]) Apply a power transform featurewise to make data more Gaussian-like.
preprocessing.QuantileTransformer(*[, ...])
                                                   Transform features using quantiles information.
preprocessing.RobustScaler(*[, ...])
                                                   Scale features using statistics that are robust to outliers
preprocessing.SplineTransformer([n_knots, ...]) Generate univariate B-spline bases for features.
                                                  Standardize features by removing the mean and scaling to unit variance.
preprocessing.StandardScaler(*[, copy, ...])
```

https://scikit-learn.org/stable/modules/classes.html#module-sklearn.preprocessing

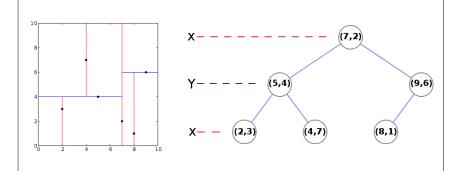
#### Irrelevant features

## k-NN Regression

- Prediction
  - ✓ instead of taking a majority vote (as in classification)
  - ✓ return the average output of the k nearest neighbors

## Computational cost

➤ Can use advanced algorithms and data structures ✓ e.g., kd-trees



## Weighted k-NN

➤ Can weight the votes according to their distance
✓ for example:

$$w = \frac{1}{d^2}$$

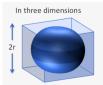
## Curse of dimensionality

• What fraction of the points lie outside the sphere?



What fraction of the square (i.e the cube) is outside the inscribed circle (i.e the sphere) in two dimensions?

$$1 - \frac{\pi r^2}{4r^2} = 1 - \frac{\pi}{4}$$



What fraction of the cube is outside the inscribed sphere in three dimensions?

$$1 - \frac{\frac{4}{3}\pi r^3}{8r^3} = 1 - \frac{\pi}{6}$$

as dimensionality increases, this fraction approaches 1!

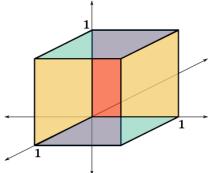
distances do not behave the same way in high dimensions

https://svivek.com/teaching/machine-learning/lectures/slides/nearest-neighbors/nearest-neighbors.pdf

# Why k-nn might work?

- ▶ Data is not always uniformly distributed over d dimensions
  - ✓ **P** may be lying on a low-dimensional subspace (low intrinsic dimensionality)
  - ✓ P may be on an underlying manifold
    - ✓ local distances (such as nearest neighbors) work better than global distances

## Curse of dimensionality



Assume **n** points are **uniformly distributed** and we are looking for the **k** nearest neighbors in **d** dimensions

Now think about the volume of the minimal enclosing box for the set of **k** nearest neighbors

$$l^d pprox rac{k}{n}$$

Solve for **l** and play with different values for **d** 

## **Summary**

- ▶ No assumptions about **P** 
  - ✓ adapts to data density
- Cost of learning is zero
  - ✓ unless a kd-tree or other data structures are used
- Need to normalize/scale the data
  - ✓ features with larger ranges dominate distances (automatically becoming more important)
  - ✓ be careful: sometimes range matters

## **Summary**

- ▶ Irrelevant or correlated attributes add noise to distance
  - ✓ may want to drop them
- Prediction is computationally expensive
  - ✓ can use **kd-trees** or **hashing techniques** like Locality Sensitive Hashing (LSH)
- Curse of dimensionality
  - ✓ data required to generalize grows exponentially with dimensionality
  - ✓ distances less meaningful in higher dimensions