Overfitting and Cross Validation

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Most contents (text or images) of course slides are from the following textbook
Provost, Foster, and Tom Fawcett. Data Science for Business: What you need to
know about data mining and data-analytic thinking. "O'Reilly Media, Inc.", 2013

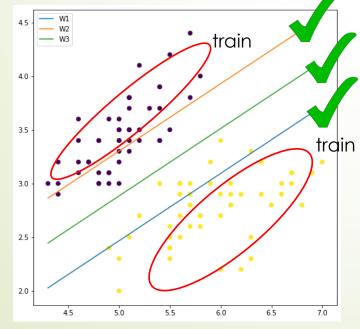
Outline

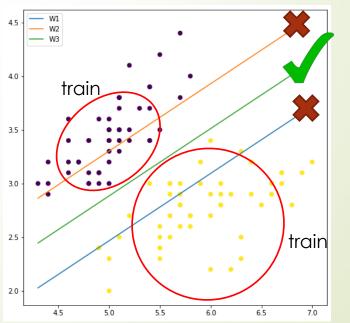
- Cross Validation
- Overfitting Avoidance
- Similarity and Distance

Drawbacks of Holdout Evaluation

- While a holdout set will indeed give us an estimate of generalization performance, it is just a single estimate.
 - A single estimate of model accuracy might have just been a single particularly lucky (or unlucky) choice of training and test data
 - ✓ Should we have any confidence in our estimation?







Repeated Holdout Evaluation with Randomness

Random splits do NOT guarantee that all splits will be different, although this is still very likely for sizeable datasets and many runs.

train_test_split():

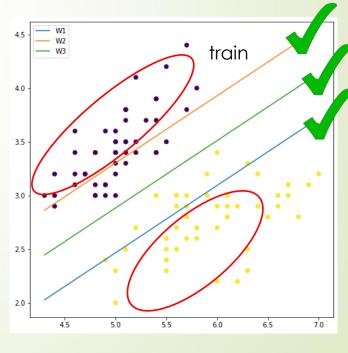
shuffle: boolean, optional (default=True)

Whether or not to shuffle the data before splitting. If shuffle=False then stratify must be None.





怎么老是你?

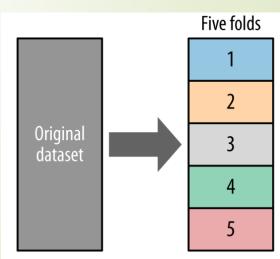


1st Run

n-th Run

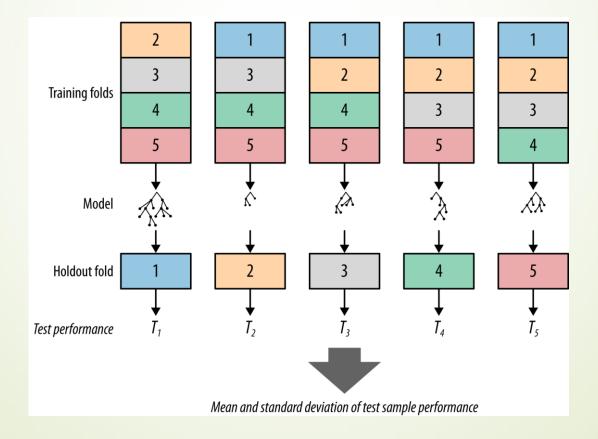
Cross-validation

- Cross-validation (CV) computes the estimation of out_performance over the whole training data by performing multiple splits (training and holdout subgroups)
 - ✓ Conduct multiple times of <u>independent</u> holdout test.
 - ✓ Not only a simple estimate of the generalization performance, but also some statistics on the estimated performance (mean, variance)
 - ✓ Different implementations: **k-fold**, leave-p/1-out, Monte Carlo cross-validation
 - k-fold cross-validation split whole training data into k partitions called *folds*
 - \checkmark Iterate training and testing k times
 - ✓ In each iteration, a different fold is chosen as the holdout(test)data; and the other k 1 folds are combined to form the training data



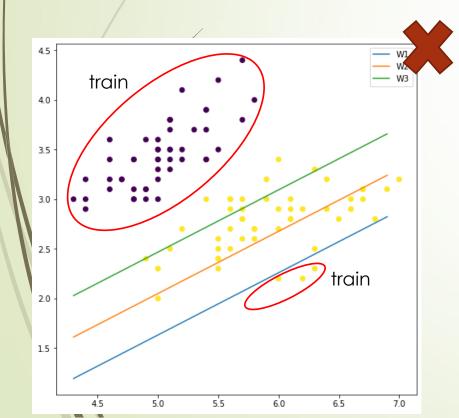
K-fold Cross-validation

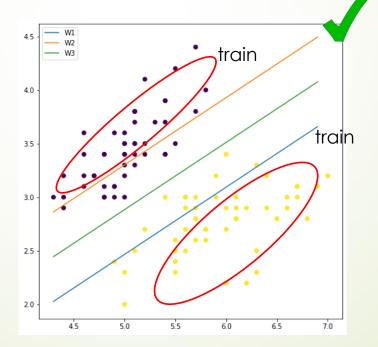
- Every example will be used once for testing but k-1 times for training
- \blacksquare Compute average and standard deviation from the evaluation results of k folds

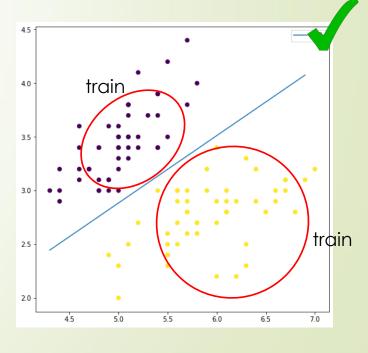


Stratified CV

- Different proportion of labels in different train-test splits
 - Stratify =True parameter makes a split preserving the percentage of samples for each class.



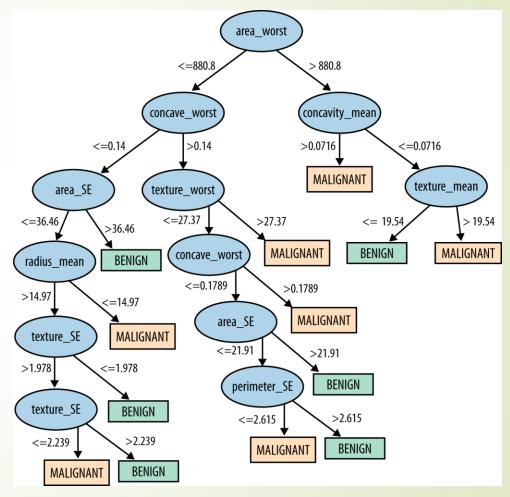




Revisit "Logistic Regression vs. Decision Tree"

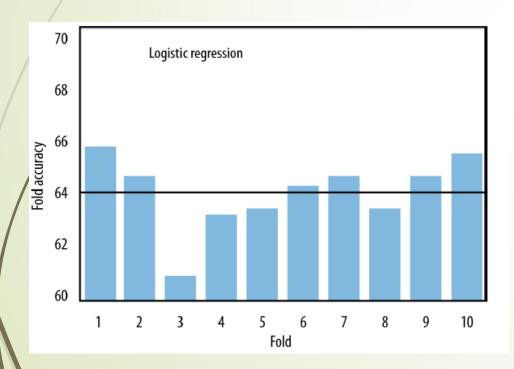
Attribute	Weight (learned parameter)
SM00THNESS_worst	22.3
CONCAVE_mean	19.47
CONCAVE_worst	11.68
SYMMETRY_worst	4.99
CONCAVITY_worst	2.86
CONCAVITY_mean	2.34
RADIUS_worst	0.25
TEXTURE_worst	0.13
AREA_SE	0.06
TEXTURE_mean	0.03
TEXTURE_SE	-0.29
COMPACTNESS_mean	-7.1
COMPACTNESS_SE	-27.87
w ₀ (intercept)	-17.7

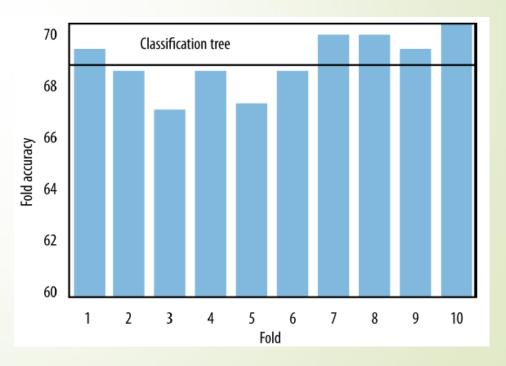
Linear equation learned by Logistic regression acc. = 98.9% (In_performance)



Deducted decision tree(J48) acc. = 99.1% (In_performance)

Logistic Regression vs. Decision Tree

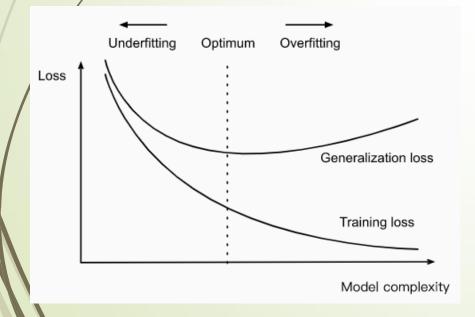




Churn Prediction

Interpreting Fitting Graph

- Focusing on the performance on the holdout data among cross validation
 - Validation curve in sklearn: training and test scores for varying parameter values
- In-performance may help deterring the direction of model complexity
 - More complex models will offer more higher flexibility to pick up complex but effective patterns (good in performance)



$$f(\mathbf{x}) = w_0' + w_1' * x_1 + w_2' * x_2 + w_3' * x_3 + w_4' * x_4$$

$$\begin{bmatrix} w_0' = w_0 \\ w_1' = w_1 \\ w_2' = w_2 \\ w_3' = 0 \\ w_4' = 0 \end{bmatrix}$$

$$f(\mathbf{x}) = w_0 + w_1 * x_1 + w_2 * x_2$$

Outline

- Cross Validation
- Overfitting Avoidance
- Similarity and Distance
- Quiz

Generals Method for Avoiding Overfitting

- 1-Control the model complexity (assuming that each model has a parameter C to control complexity)
 - ✓ <u>Complexity parameter C</u>: decision tree number of nodes; mathematical functionweight of regularization (larger weight of regularization → lower model complexity)
 - ✓/ Occam's razor: choose the simplest model/hypothesis that can fit the data well
 - How to set parameter C?
- 2-Feature Selection use fewer attributes
 - ✓ What is the subset of features should be retained?
- 3-Collect more data
 - ✓ Using the learning curve

Avoiding Overfitting in Tree Induction

- Pre-pruning: stop growing the tree before it gets too complex
 - ✓ Stop criteria: the information gain/nodes in leaf is lower than a threshold.
 - ✓ Decide the threshold by experience, or using hypothesis test
 - ✓ Computational efficient, but easy to be under-fitted
- **Post-pruning:** takes a fully-grown decision tree and discards unreliable parts
 - Estimate whether replacing a set of leaves or a branch with a leaf would reduce accuracy? If not, do the replacement
 - Continue process iteratively, until any removal or replacement would reduce accuracy

Complexity Control in Mathematical Functions

For models described as a mathematical function, its loss on the whole training data can also be represented as a mathematical function *TrainLoss(w)*. Then we try to find the best model parameter w that generates least loss on training data

$$\min_{\mathbf{w}} TrainlLoss(\mathbf{w})$$

- Occam's razor: choose the simplest model/hypothesis that can fit the data well
- Regularization: Add the penalty of model complexity to the objectives of optimizations
 - ✓ Weight C determines how much importance the optimization procedure should place on the penalty, compared to TrainLoss
 - ✓ Trade-off between model simplicity and its in_performance

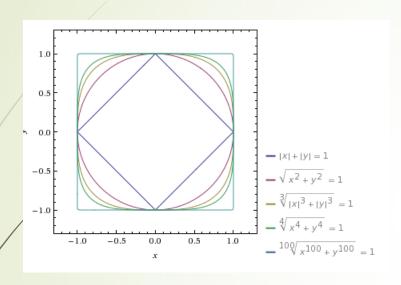
$$\min_{\mathbf{w}} TrainlLoss(\mathbf{w}) + C \cdot penalty(\mathbf{w})$$

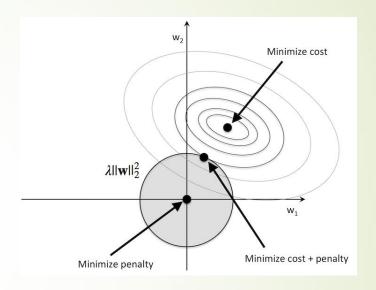
Popular Complexity Penalty

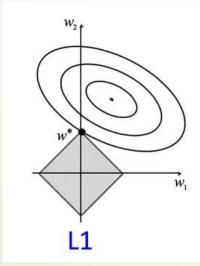
We try to find optimal \mathbf{w} by optimizing the following regularized objective $\min_{\mathbf{w}} TrainlLoss(\mathbf{w}) + C \cdot penalty(\mathbf{w})$

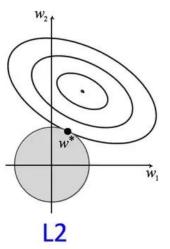
- ▶ L2-norm penalty: $penalty(\mathbf{w}) = \|\mathbf{w}\|_2 = \sqrt{w_1^2 + w_2^2 + \dots + w_n^2}$
 - ✓ The sum of the squares of the weights (avoid large negative/positve weights)
 - ✓ L2-norm + standard least-squares linear regression = ridge regression
- ▶ L1-norm penalty: $penalty(\mathbf{w}) = \|\mathbf{w}\|_1 = |w_1| + |w_2| + ... + |w_n|$
 - ✓ The sum of the absolute values of the weights
 - ✓ L1-norm + standard least-squares linear regression = LASSO regression
 - ✓ Zero out entries of weights (drop corresponding feature)

Intuition of L1/L2-norm Penalty*









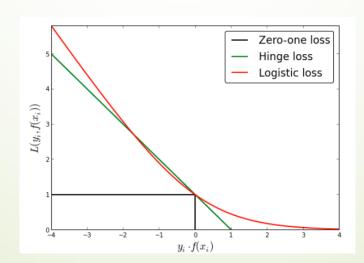
SVM as Regularized Linear Model

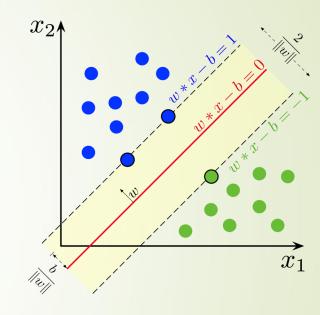
SVM tries to optimize

$$\min_{\mathbf{w}} HingeLoss(\mathbf{w}) + C \cdot \|\mathbf{w}\|_2$$

Linear SVM learning is almost equivalent to L2-regularized Logistic regression as follows

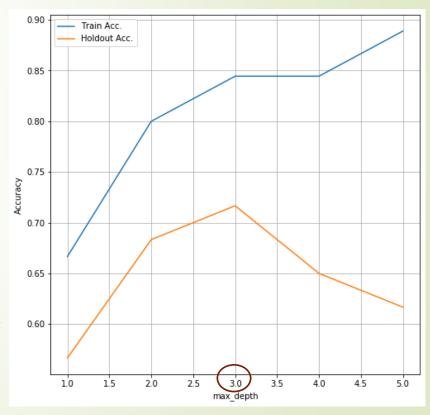
$$\min_{\mathbf{w}} LogisticLoss(\mathbf{w}) + C \cdot \|\mathbf{w}\|_{2}$$





Parameter Selection with Cross-validation

- Treat models with different complexity parameter C or with different subset of features as different models
 - Examples of C: trade coefficient of SVM, max_depth for decision tree ...
 - Use cross-validation to estimate their generalization performance respectively and select the best model accordingly.
 - The corresponding parameters of the best model are the optimal parameter C_{best}
- Train the model with parameter C_{best} on the whole training data to get the final trained model (during CV we only trained on k-1 folds of training data)



GridSearchCV for Multiple Parameters

SVM with kernel = 'poly'

C: float, optional (default=1.0)

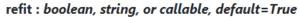
Regularization parameter. The strength of the regularization is inversely proportional to C. Must be strictly positive. The penalty is a squared I2 penalty.

kernel: string, optional (default='rbf')

Specifies the kernel type to be used in the algorithm. It must be one of 'linear', 'poly', 'rbf', 'sigmoid', 'precomputed' or a callable. If none is given, 'rbf' will be used. If a callable is given it is used to pre-compute the kernel matrix from data matrices; that matrix should be an array of shape (n samples).

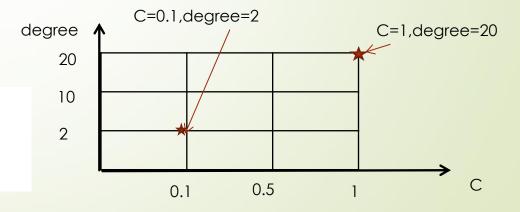
degree: int, optional (default=3)

Degree of the polynomial kernel function ('poly'). Ignored by all other kernels.



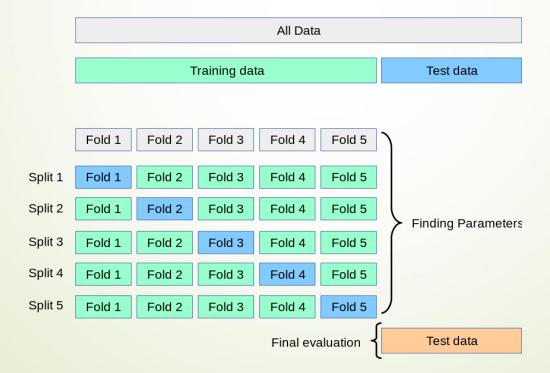
Refit an estimator using the best found parameters on the whole dataset

For multiple metric evaluation, this needs to be a string denoting the scorer that would be used to find the best parameters for refitting the estimator at the end.



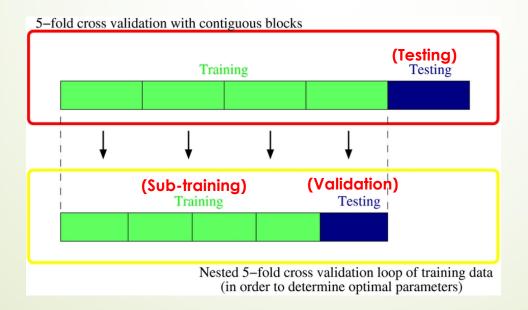
Estimated Generalization Performance

- Report the estimated generalization performance by simple cross-validation is overly optimistic
 - We choose the best model by estimated generalization performance, and then report the generalization performance directly



Nested Cross Validation*

- Nested cross validation
 - ✓ (Outer) Repeatedly split whole training data into training/testing data. (Fit selected model to training data, and report performance on testing data)
 - (Inner) Do cross-validation on the training data by further repeatedly splitting the training into sub-training/validation(select the best model/parameter)



2-Feature Selection

- Sequential forward selection (SFS) can help us to select features based on nested cross-validation
 - ✓ Pick the best individual feature by looking at all models built with just one feature
 - ✓ Then, test all models that add a second feature to first chosen feature
 - ✓ Proceed similarly with three, four, ... features
- SFS is computationally expensive by evaluating all combinations of features
- Sequential backward elimination (Using all features initially, but delete feature one by one based on nested cross-validation)
- L1-norm for feature selection (drop the features associated with zero weights)

$$f(\mathbf{x}) = w_0' + w_1' * x_1 + w_2' * x_2 + w_3' * x_3 + w_4' * x_4$$

$$w_0' = w_0$$

$$w_1' = w_1$$

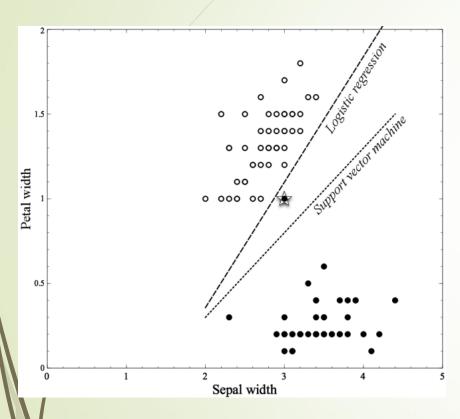
$$w_2' = w_2$$

$$w_3' = 0$$

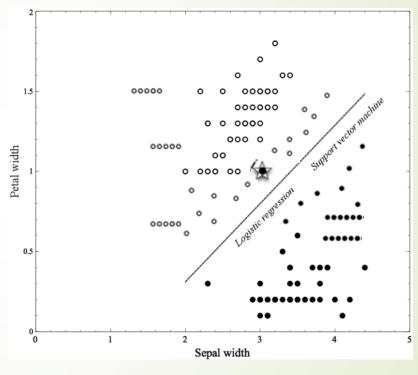
$$w_4' = 0$$

$$f(\mathbf{x}) = w_0 + w_1 * x_1 + w_2 * x_2$$

3-Collect More Data: Intuition



Training data with one outliers

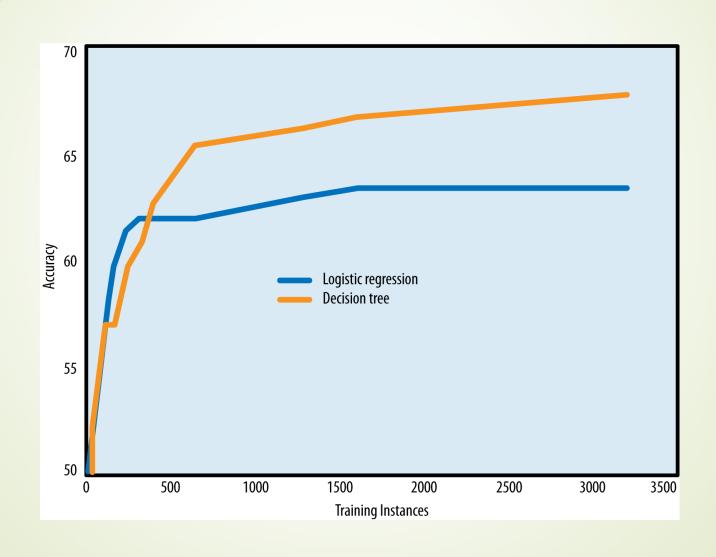


Collecting more data

3-Collect More Data: Learning Curve

- The <u>learning curve</u> is a plot of the estimated generalization performance against the amount of training data
 - ✓ Usually steep initially, but then marginal advantage of more data decreases
 - ✓ Generalization performance improves as more training data are available.
 - Fitting curve shows the performance on the training and the holdout data against model complexity (for a fixed amount of training data)
- Learning curve may give recommendations on how much to invest in training data
 - ✓ Different modeling procedures may have different performance on the same data
 - ✓ Different size of training data may result in different estimated generation performance of the same model

Example of Learning Curve



Summary of CV in Sklearn

https://scikit-learn.org/stable/modules/cross_validation.html

- CV split:
 - train_test_split : random holdout split
 - StratifiedKFold: stratified k-fold CV
 - Kfold: k-fold CV
- CV evaluation
 - cross_validate: only score
 - cross_val_score: detailed information
- CV for model selection
 - validation_curve: Single parameter
 - gridSearchCV
- CV for learning curve
 - learning_curve

Outline

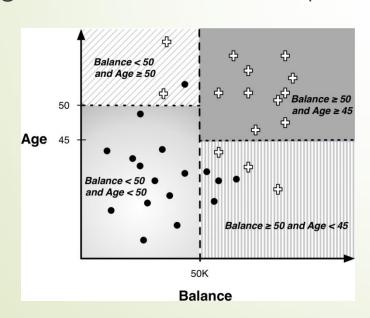
- Cross Validation
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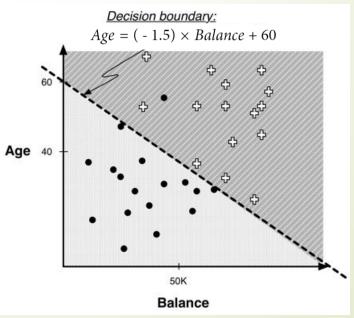
Applications of Similarity

- Similarity underlies many data science methods and solutions to business problems
- If two things (people, companies, products) are similar in some ways, they often share other characteristics as well
- Different sorts of business tasks involve reasoning from similar examples
 - Retrieve similar things: search engine
 - ✓ Classification or regression: K-Nearest-Neighbor
 - ✓ Clustering: grouping similar things tougher
 - ✓ Recommender system: Amazon -- "People who like X also like Y"
 - ✓ Reasoning from similar cases beyond business applications: lawyers and doctors

Similarity under Segmentations

- Both classification trees and linear classifiers establish boundaries between regions of differing classifications.
- They have in common the view that instances sharing a common region in space should be similar.
- What differs between classification trees and linear classifiers is how the regions/boundaries are represented and discovered.





Similarity and Distance

- Why not reason about the similarity or distance between objects directly?
- If we represent each object as a feature vector, then the closer two objects are in the space defined by the features, the more similar they are.
- For example:

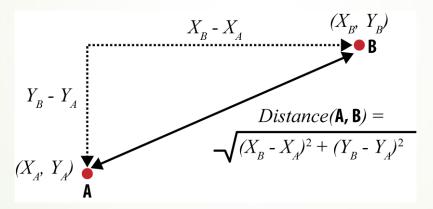
Attribute	Person A	Person B
Age	23	40
Years at current address	2	10
Residential status (1=0wner, 2=Renter, 3=0ther)	2	1

How to measure the similarity or distance between Person A and Person B?

What does it mean that two companies or two consumers are similar?

Euclidean distance

- Euclidean distance is the "ordinary" straight-line distance between two points in Euclidean space.
 - When each sample have two features, then each object is a point in a twodimensional space. (<u>Pythagorean theorem</u>)



When an object is described by n-dimensions features $(d_1, d_2, ..., d_n)$, the general equation for Euclidean distance in n dimensions is shown as

$$\sqrt{(d_{1,A} - d_{1,B})^2 + (d_{2,A} - d_{2,B})^2 + ... + (d_{n,A} - d_{n,B})^2}$$

Euclidean distance (Example)

Euclidean distance for Person A and Person B

$$d(A, B) = \sqrt{(23 - 40)^2 + (2 - 10)^2 + (2 - 1)^2}$$

$$\approx 18.8$$

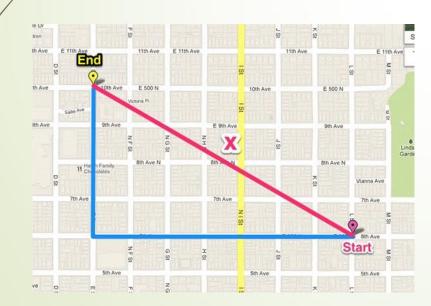
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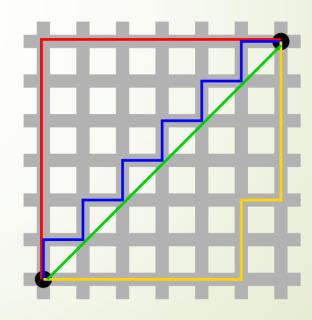
- ✓ This distance is just a number—it has no units, and no meaningful interpretation.
- Distance is only really useful for comparing the similarity of one pair of instances to that of another pair, such as d(A,B) <>= d(A,C)

Other Distance Measure (1)

- Manhattan distance: L1-norm distance
 - ✓ Application: Compressed sensing, compressed sensing ...

$$d_{\text{Manhattan}}(\mathbf{X}, \mathbf{Y}) = \| \mathbf{X} - \mathbf{Y} \|_{1} = \| x_{1} - y_{1} \| + \| x_{2} - y_{2} \| + \cdots$$

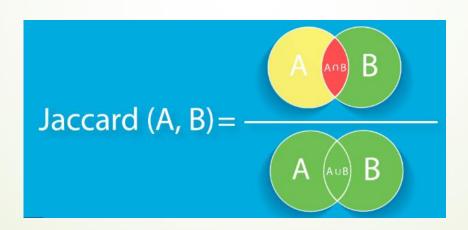




Other Distance Measure (2)

- Jaccard distance measure treats the two objects as sets of characteristics
 - ✓ The possession of a <u>common characteristic</u> between two item sets is important, but
 the <u>common absence</u> of a characteristic is not

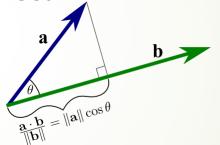
$$d_{\text{Jaccard}}(X, Y) = 1 - \frac{|X \cap Y|}{|X \cup Y|}$$



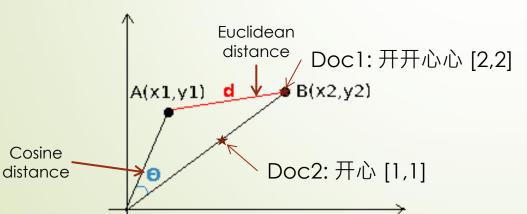
Other Distance Measure (3)

- Cosine distance: often used in text classification to measure the similarity of two documents
 - ✓ Often used in text classification to measure the similarity of two documents
 - ✓ Ignore differences in scale across instances

$$d_{cosine}(\mathbf{X}, \mathbf{Y}) = 1 - \frac{\mathbf{X} \cdot \mathbf{Y}}{\parallel \mathbf{X} \parallel_{2} \cdot \parallel \mathbf{Y} \parallel_{2}}$$



$$A = \langle 7, 3, 2 \rangle$$
 and $B = \langle 2, 3, 0 \rangle$



$$d_{\text{cosine}}(A, B) = 1 - \frac{\langle 7, 3, 2 \rangle \cdot \langle 2, 3, 0 \rangle}{\| \langle 7, 3, 2 \rangle \|_{2} \cdot \| \langle 2, 3, 0 \rangle \|_{2}}$$

$$= 1 - \frac{7 \cdot 2 + 3 \cdot 3 + 2 \cdot 0}{\sqrt{49 + 9 + 4} \cdot \sqrt{4 + 9}}$$

$$= 1 - \frac{23}{28 \cdot 4} \approx 0.19$$

Other Distance/Similarity Measure (4)

- Hamming distance (equal length)
- Edit distance
- Longest common substring
- Longest common subsequence

Example: Similar Whiskey

- How can we can find the most similar whiskey of a given type of whiskey as a data scientist
 - ✓ Construct 5 attributes that can describe the general whiskey as follows.

Color: yellow, very pale, pale, pale gold, gold, old gold, full gold, amber, etc.(14 values)Nose: aromatic, peaty, sweet, light, fresh, dry, grassy, etc.(12 values)Body: soft, medium, full, round, smooth, light, firm, oily.(8 values)Palate: full, dry, sherry, big, fruity, grassy, smoky, salty, etc.(15 values)Finish: full, dry, warm, light, smooth, clean, fruity, grassy, smoky, etc.(19 values)

- ✓ The values of attributes are NOT mutually exclusive (e.g., Aberlour's palate is described as medium, full, soft, round and smooth).
- ✓ Use a feature vector of 68 (=14+12+8+15+19) binary (0/1) attributes to reprint a type of whiskey as [0,1,1,0,...,1] (with 68 entries of 0/1)

Example: Similar Whiskey by Euclidean Distance

- Given a special type of whiskey (Bunnahabhain), we could compute the Euclidean distance between it and other whiskey, respectively
- We could rank the other whiskey by the distance in descending order as follows

Whiskey	Distance	Descriptors
Bunnahabhain	_	gold; firm,med,light; sweet,fruit,clean; fresh,sea; full
Glenglassaugh	0.643	gold; firm,light,smooth; sweet,grass; fresh,grass
Tullibardine	0.647	gold; firm,med,smooth; sweet,fruit,full,grass,clean; sweet; big,arome,sweet
Ardbeg	0.667	sherry; firm,med,full,light; sweet; dry,peat,sea;salt
Bruichladdich	0.667	pale; firm,light,smooth; dry,sweet,smoke,clean; light; full
Glenmorangie	0.667	p.gold; med,oily,light; sweet,grass,spice; sweet,spicy,grass,sea,fresh; full,long

Outline

- Cross-validation & Learning Curve
- Overfitting Avoidance
- Similarity and Distance
- Quiz

Lab Quiz

- **Deadline**: 17:59 p.m., Mar. 27, 2020
- Two questions accounting for 5% of overall score
- Upload the answer worksheet and the accomplished Python files to the Blackboard
- You may submit unlimited times but only the LAST submission will be considered
- Only the answers in answer sheet will be referred for grading
- Note: MUST attach ALL the required files in every submission/resubmission, otherwise other files will be missing.