K-Nearest Neighbors, Decision Trees and Random Forests

Machine Learning 1 — Lecture 10 28th May 2024

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Algorithms for Supervised Learning

- So far, we discussed parametric models for supervised learning
 - linear models
 - linear models with non-linear features
 - neural networks
- Each of these models was specified via a parameter vector θ and we specified a training loss $\mathcal{L}_{train}(\theta)$ which we aimed to minimize w.r.t. θ
- There are, however, many other approaches to supervised learning

Algorithms for Supervised Learning

- "No free lunch theorem:" for any machine learning algorithm, there is some task where the algorithm does not perform best when provided with finitely many data
- Thus, getting best performance requires trying different learning algorithms
- Today, we are going to discuss
 - K-nearest neighbours (KNN)
 - Decision trees
 - Random forests

K-Nearest Neighbours

- Assume a training set $\mathcal{D} = \{(\mathbf{x}^{(1)}, y^{(1)}), \dots, (\mathbf{x}^{(N)}, y^{(N)})\}$
- No training required: just dump the dataset in your memory
- Assume a test sample x^* for which we don't know target y^*
- Let i be the index of the **nearest neighbour** of x^* , i.e.

$$i = \arg\min_{i} \|\boldsymbol{x}^* - \boldsymbol{x}^{(i)}\|$$

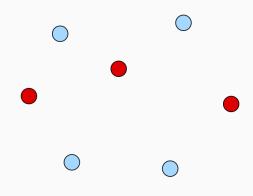
for some norm $\|\cdot\|$

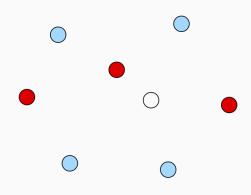
 The nearest neighbour predictor returns the target of the nearest neighbour

$$f_{\mathsf{NN}}(\mathbf{x}^*) = y^{(i)}$$

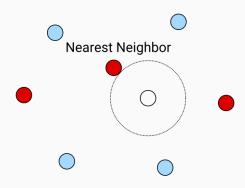
Note that this works for both classification and regression



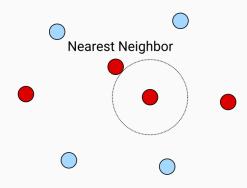












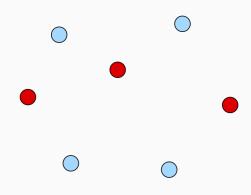
K-Nearest Neighbors (KNN)

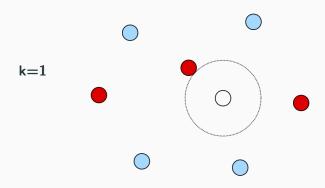
- Why should we restrict to only one neighbour?
- Instead, we can combine the targets of the K nearest neighbours
- Sort training points according to distance to x*:

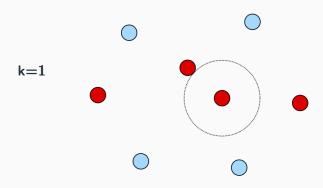
$$\|\mathbf{x}^{(i_1)} - \mathbf{x}^*\| \le \|\mathbf{x}^{(i_2)} - \mathbf{x}^*\| \le \dots \le \|\mathbf{x}^{(i_K)} - \mathbf{x}^*\| \le \dots \le \|\mathbf{x}^{(i_N)} - \mathbf{x}^*\|$$

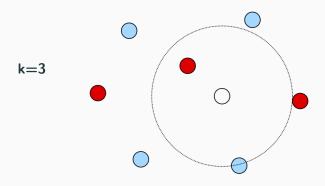
- Classification: Label test sample according to majority vote
 of K nearest neighbours. Break ties randomly.
- **Regression:** Average targets of *K* nearest neighbours

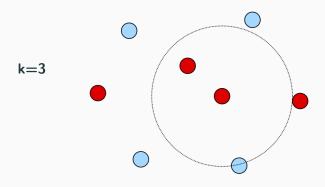
$$\hat{y} = \frac{1}{K} \sum_{k=1}^{K} y^{(i_k)}$$

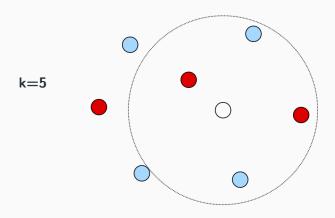


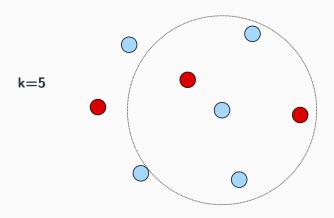












- Assume a classification setting, i.e. y is discrete
- Assume we would know the true data-generating distribution p_{true}(x, y) (Recall Lecture 8)
- Then we could compute $p_{true}(y \mid \mathbf{x}) = \frac{p_{true}(\mathbf{x}, y)}{p_{true}(\mathbf{x})} = \frac{p_{true}(\mathbf{x}, y)}{\sum_{y} p_{true}(\mathbf{x}, y)}$
- Then, a natural classification rule would be:

$$\hat{y} = \arg\max_{y} \, p_{true}(y \,|\, \boldsymbol{x})$$

- This rule is known as the Bayes optimal classifier
- It suffers the least **true classification error**, i.e.

$$\mathbb{E}_{\mathbf{x}, y \sim p_{true}} \left[\mathbb{1}(\hat{y} \neq y) \right]$$

is minimal among all possible classifiers.

Consistency of KNN

KNN is simple, but theoretically sound!

In particular, if

- number of training points $N \to \infty$
- number of neighbours $K \to \infty$
- ullet N grows quicker than K, i.e. $rac{K}{N}
 ightarrow 0$

then KNN converges towards the Bayes optimal classifier.

A classifier with this property is called **consistent**.

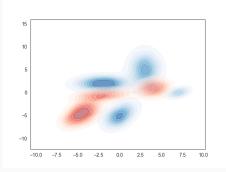
Furthermore, for regression, KNN converges to the **optimal** regression function $\mathbb{E}_{p_{true}(y \mid x)}[y]$.

Bayes Consistency of KNN

Example

We can demonstrate Bayes consistency of KNN on a toy dataset.

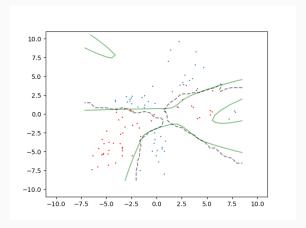
Assume a binary classification problem and assume the following true data distribution $p_{true}(x, y)$ (red and blue for the two classes):



Thus, we can actually compute the Bayes optimal classifier!

Bayes Consistency of KNN cont'd

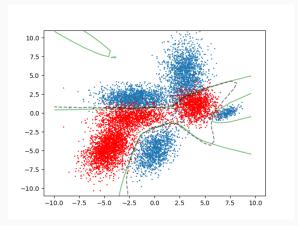




Green line: Bayes optimal classifier (94.67% test accuracy) Dashed: KNN $N_{train}=100$ and K=9 (90.26% test accuracy)

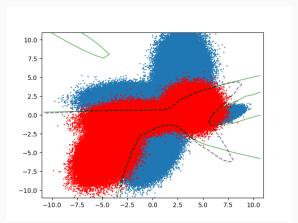
(accuracies computed on a 1 million test samples.)

Bayes Consistency of KNN cont'd



Green line: Bayes optimal classifier (94.67% test accuracy) Dashed: KNN $N_{train}=10k$ and K=99 (94.56% test accuracy) (accuracies computed on a 1 million test samples.)

Bayes Consistency of KNN cont'd



Green line: Bayes optimal classifier (94.67% test accuracy) Dashed: KNN $N_{train}=1M$ and K=999 (94.65% test accuracy) (accuracies computed on a 1 million test samples.)

- simple but consistent predictor
- training is very easy: just store the data
- testing is $\mathcal{O}(N)$
- k-d trees organize the data in hierarchical way, then testing can be done on O(log N)

Decision Trees

Decision Trees

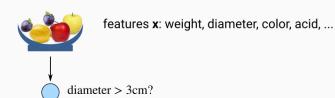


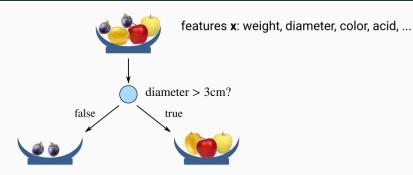


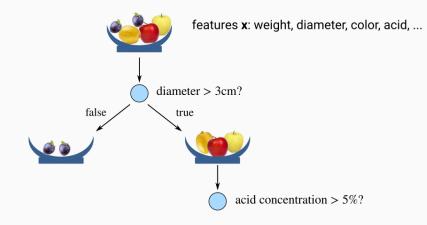


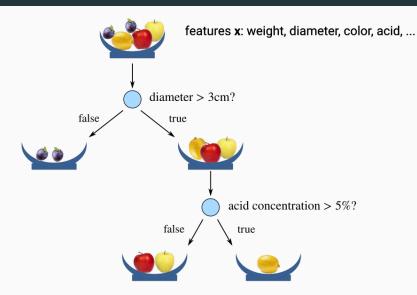
features $\mathbf{x}\!:$ weight, diameter, color, acid, \dots











Given a set of features $X_1, X_2, ..., X_D$ (discrete or continuous) and a class variable Y, a Decision Tree is a **directed binary tree** with two types of nodes:

Decision Nodes (internal nodes)
 Decision nodes are associated with a feature X_i and a boolean function (decision, test)

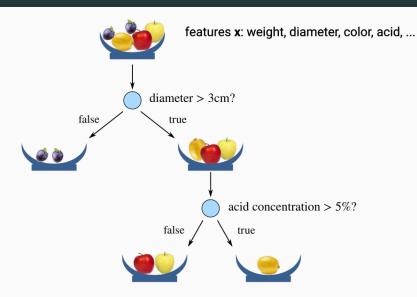
$$f: X_i \mapsto \{\mathit{true}, \mathit{false}\}$$

Decision nodes have 2 children labeled true and false, which are selected by the output of f.

- Prediction Nodes (leaves)
 Prediction nodes are associated with either
 - a fixed label y
 - a distribution over labels p(Y)

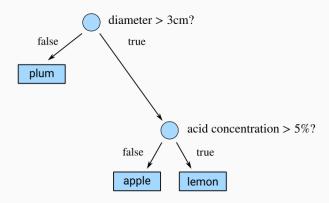
Classification using a Decision Tree

- Parse decisions from root to leaf
- Since the decision tree is a tree, there is a unique path from root to each leaf
- Leaf is selected, if all tests on this path are true
- Selected leaf is guaranteed to be unique, since decision nodes select exactly one child
- Use the predictor in the leaf for classification:
 - a fixed label y
 - a distribution over labels p(Y)
- Easily generalized to non-binary decision nodes
- Decision trees can also be used for regression (not discussed)



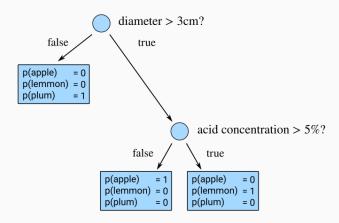


features x: weight, diameter, color, acid, ...



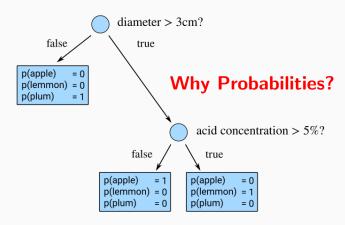


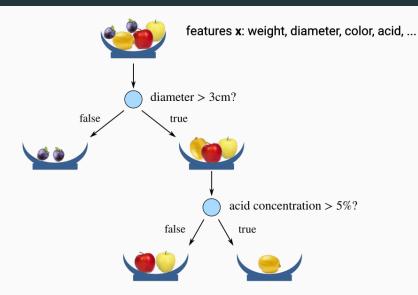
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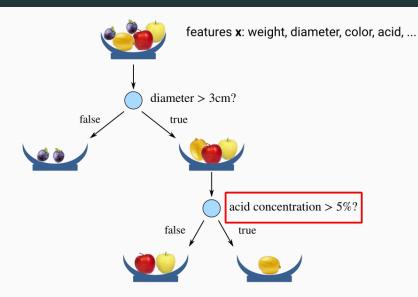


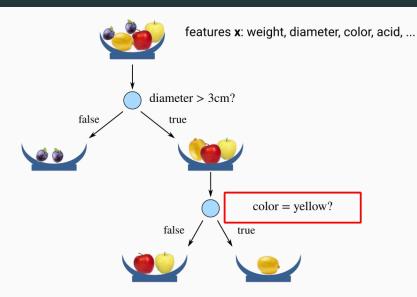


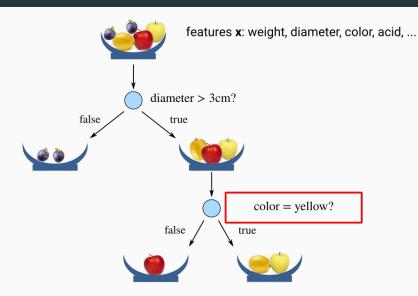
features x: weight, diameter, color, acid, ...





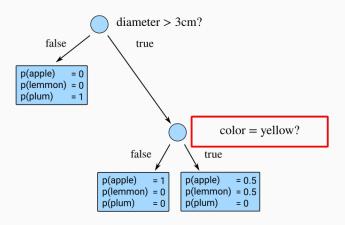






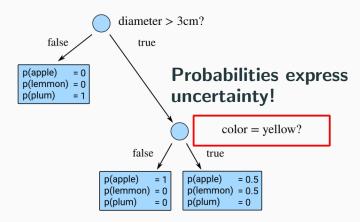


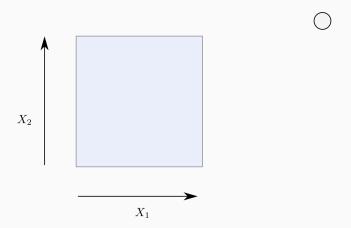
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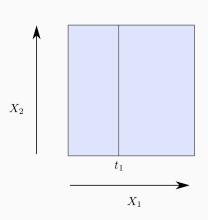


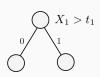


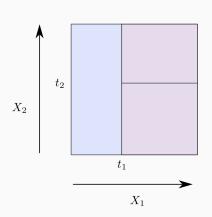
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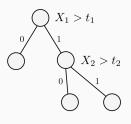


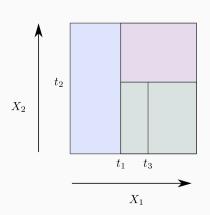


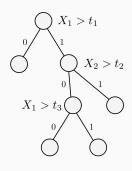


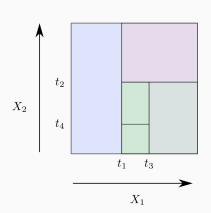


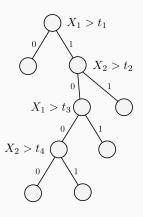


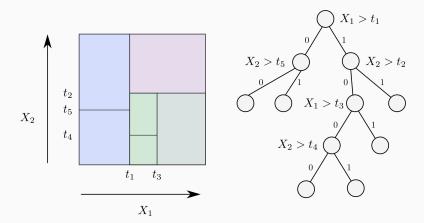












Learning Algorithms for Decision Trees

- CART (Classification and Regression Tree) [Breiman'84]
- ID3 [Quinlan'86]
- C4.5 [Quinlan'93]
- ...

CART

```
1 input: Data (x_1, y_1), (x_2, y_2), \dots, (x_N, y_N)
 2 result: Decision Tree
3 Initialize single leaf L, assign all data points \mathbf{x}^{(i)} to L, call split(L)
 4 split(L):
 5 if stopping criterion is true for L then
        Learn p(Y) from class proportions of samples assigned to L
        return
8 end
 9 for considered variables/decisions X_d, f do
      compute cost(X_d, f)
11 end
12 Let X_{min}, f_{min} be the decision with minimal cost
13 Apply f_{min}(X_{min}) to L, yielding new leaves L_0 and L_1
14 Assign \mathbf{x}^{(i)} with f(\mathbf{x}^{(i)}) = \text{false} to L_0, call \mathbf{split}(L_0)
15 Assign \mathbf{x}^{(i)} with f(\mathbf{x}^{(i)}) = true to L_1, call \mathbf{split}(L_1)
```

CART

```
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```

Possible Decisions

```
input: Data (x_1, y_1), (x_2, y_2), \dots, (x_N, y_N)
   Result: Decision Tree
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Assign \mathbf{x}^{(i)} with f(\mathbf{x}^{(i)}) = true to L_1, call \mathbf{split}(L_1)
```

Which Decisions to Consider?

Continuous Feature X_d

- thresholds $X_d > t$
- sort data values for X_d
- consider thresholds halfway between consecutive values
- only finitely many decisions to consider
- e.g., for values $X_d \in [0.1,\ 1.3,\ 3.1415]$, we consider thresholds $\frac{0.1+1.3}{2}=0.7$ and $\frac{1.3+3.1415}{2}=2.22$

Discrete Feature X_d

- one-vs-all $\{1,2,3,4\} \rightarrow \{1,2,4\},\{3\}$
- complete split (not a binary tree)
 {1,2,3,4} → {1}, {2}, {3}, {4}

Cost

```
input: Data (x_1, y_1), (x_2, y_2), \dots, (x_N, y_N)
   Result: Decision Tree
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```



- "Pure leaves are good"
- Let N be the number of data points at leaf L
- For a particular candidate decision let N_0 , N_1 be the number of data points at new leaves L_0 and L_1 , respectively
- $cost = \frac{N_0}{N} impurity(L_0) + \frac{N_1}{N} impurity(L_1)$
- Note that $rac{N_0}{N} + rac{N_1}{N} = 1$ (weighted average)

Impurity Measures

Let p(Y) be the empirical frequencies of class labels, e.g.



$$p(plum) = \frac{5}{8} = 0.625$$

 $p(lemon) = \frac{3}{8} = 0.375$

• Gini Impurity

$$G=1-\sum_{y}p(y)^{2}$$

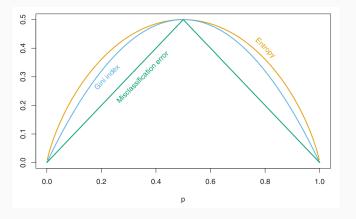
Entropy

$$E = -\sum_{y} p(y) \log(p(y))$$

Self-Classification Error

$$C = 1 - \max_{y} p(y)$$

Impurity measures for binary classification, as a function of p(y=0) (and p(y=1) due to symmetry). Entropy has been scaled to pass through (0.5, 0.5).



Cost

- Let N be the number of data points at leaf L
- Let N₀, N₁ be the number of data points at new leaves L₀ and L₁, respectively.
- $cost = \frac{N_0}{N} impurity(L_0) + \frac{N_1}{N} impurity(L_1)$
- Where impurity might be
 - Gini Impurity

$$G=1-\sum_{y}p(y)^{2}$$

Entropy

$$E = -\sum_{y} p(y) \log(p(y))$$

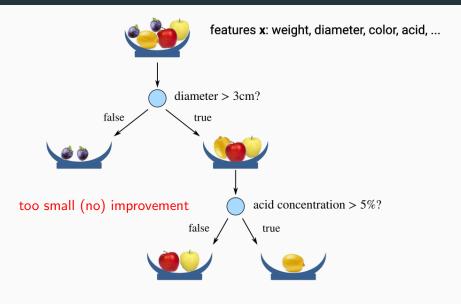
Self-Classification Error

$$C = 1 - \max_{y} p(y)$$

Stopping Criterion

```
input: Data (x_1, y_1), (x_2, y_2), \dots, (x_N, y_N)
   Result: Decision Tree
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```

Stopping Criterion



too small (no) improvement too few data points

Stopping Criterion and Hyperparameters

- Threshold determining when improvement of cost is too small
- Another threshold for determining when there are too few data points
- Furthermore, one introduce a maximal tree depth
- These thresholds are hyper-parameters of CART

Random Forests

- Decision trees: well interpretable models, but don't perform too well in practice
- Random Forests: ensemble of decision trees
- Output of random forest is computed by aggregating the outputs of the individual decision trees
 - Majority vote for classification
 - Averaging for regression
- A theoretical underpinning for ensembles is the bias-variance trade-off

Bias-Variance Trade-Off

- ullet Let ${\mathcal H}$ be our considered **hypothesis class**
- Recall from Lecture 8 that the ideal goal of supervised learning is to find an optimal predictor f* minimizing the true loss:

$$f^* = \arg\min_{f \in \mathcal{H}} \mathcal{L}_{true}(f) = \mathbb{E}_{p_{true}}[\ell(f(\mathbf{x}), y)]$$

- However, we only have a finite training set of N samples drawn from p_{true}
- At best, we can find a minimizer f^e of the **empirical loss**:

$$f^e = \arg\min_{f \in \mathcal{H}} \mathcal{L}_{empirical}(f) = \frac{1}{N} \sum_{i=1}^{N} \ell(f(\mathbf{x}^{(i)}, y^{(i)}))$$

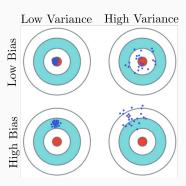
The Bias-Variance Trade-Off

- Figure that we repeatedly draw training sets of size N from p_{true}:
 - the training set is random
 - thus, the learned model femp is random
 - thus, $\mathcal{L}_{true}(f^{emp})$ is also random
 - thus, the excess loss $\delta = \mathcal{L}_{true}(f^{emp}) \mathcal{L}_{true}(f^*)$ is also random
- The $\mathbb{E}[\delta]$ is denoted as **bias**, measuring how far $\mathcal{L}_{true}(f^{emp})$ is from $\mathcal{L}_{true}(f^*)$ on average
- The variance $var[\delta] = \mathbb{E}\left[(\delta \mathbb{E}[\delta])^2\right]$ measures how much the true loss of f^{emp} varies

The Bias-Variance Trade-Off cont'd

Bias and variance depend on the model complexity:

- simple models tend to have high bias and low variance
 ⇒ underfitting
- expressive, flexible models tend to have low bias and high variance ⇒ overfitting



Why Ensembles?

- Individual models (like decision trees) easily overfit, i.e. they have low bias but high variance
- **Idea:** instead of training one model, train *K* models and aggregate them
- $\bullet~$ We can expect that $\mbox{\it var}(\mbox{ensemble}) \approx \frac{\mbox{\it var}(\mbox{single model})}{\mbox{\it K}}$
- What is the problem with this idea?

- We have only one dataset—use bootstrapping to generate synthetic copies of the training data
- In statistics, bootstrapping is a resampling method to produces uncertainty estimates
- Given N samples, generate K new datasets of size N by sampling with replacement
- These new datasets are called bootstraps
- For large N, each bootstrap contains only $1-\frac{1}{e}\approx 63.21\%$ of the original samples

Original Data	x ₁	x ₂	x ₃	x ₄	x ₅	x 6	x ₇	x 8	x 9	x ₁₀
Boostrap 1	x ₇	x 8	x ₁₀	x 8	x ₂	x ₅	x ₁₀	x ₁₀	x 5	X 9
Boostrap 2	x ₁	\boldsymbol{x}_1	x 9	\boldsymbol{x}_1	x ₂	x ₃	x ₂	x 7	x ₃	x ₂
<u>:</u>	:	:	:	:	:	:	:	:	:	÷
Boostrap K	x ₁	x 8	x ₅	x ₁₀	x ₅	x ₅	x 9	x 6	x ₃	x ₇

- Bagging with Decision Trees
- Bagging = Bootstrapping and Aggregating
- Generate K bootstraps, and learn a decision tree on each of them
- Additionally, randomize CART learning (to get more diverse trees): Rather than considering all variables for a new split, restrict to m randomly chosen variables, m < D
- Typical values, K = 100, 200, 500, $m \approx \frac{D}{3}$
- Run-time: K times the run-time of Decision Trees

Decision Tree vs. Random Forest

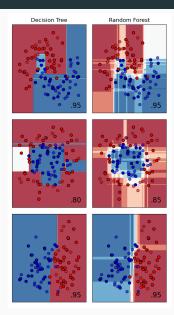
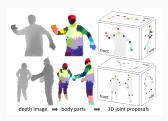


Image: https://scikit-learn.org/

Random Forests in Practice

- Kaggle reported in 2019 that decision trees and random forests are the most widely used tool in data mining and applied machine learning, after logistic regression
- Fernánandez-Delgado et al. (2014) random forests perform best among a wide range of algorithms on **tabular data**
- Prominent example: Body-part detection for skeleton tacking (Kinect 2)



- Decision trees: top-down decision diagram
- Decision nodes split input space recursively
- Leaves contain localized predictors
- CART algorithm: recursively find optimal decision (split) by minimizing impurity
- Random forest: randomized ensemble of decision trees
- Bagging: bootstrapping and aggregating