

# Introduction to Machine Learning

## Fundamentals of Machine Learning

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## Course context

- This lecture is part of the course **Fundamentals of Machine Learning**.
- Goal of this first lecture:
  - Build intuition for what machine learning (ML) is.
  - Motivate why ML is useful across many domains.
  - Introduce the basic ML paradigm and terminology.
  - Discuss feature representation and distance metrics.
- We deliberately avoid complex algorithms and heavy mathematics.

## Plan of the lectures

- 1 Introduction to Machine Learning + Practical Work (4h)
- 2 Supervised Learning: Classification, Regression + PW (4h)
- 3 Supervised Learning: More Methods, Model Evaluation + PW (4h)
- 4 Unsupervised Learning: Clustering + PW (4h)
- 5 Machine Learning Project with LLM (3h)
- 6 Competition session (3h)

# What you will learn in this course

- How to translate real-world problems into ML tasks.
- The main families of ML methods and when to use them.
- How to represent data as **feature vectors**.
- How to train, validate and evaluate ML models.
- How to reason about generalization, overfitting and underfitting.
- Hands-on ML practice in Python / common ML libraries.

# Where is Machine Learning?

- **Digital services**

- Recommendation systems (movies, music, products).
- Search ranking and query suggestion.
- Online advertisement targeting.

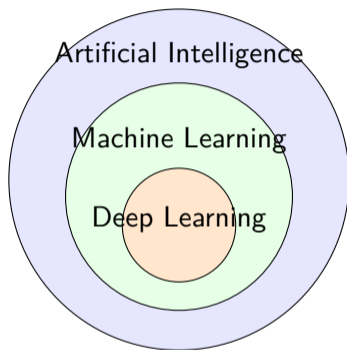
- **Perception**

- Image and face recognition.
- Speech recognition and voice assistants.
- Photo filters and style transfer.

- **Decision and control**

- Self-driving cars and robotics.
- Fraud detection and credit scoring.
- Medical diagnosis support systems.

# AI, Machine Learning and Deep Learning



- **AI**: broad field of making machines perform tasks requiring human intelligence.
- **ML**: subset of AI that learns patterns from data instead of using hand-crafted rules.
- **DL**: subset of ML based on deep neural networks, powerful for unstructured data.

# What is Machine Learning? — definitions

- Informal idea: build programs that **improve their performance with experience**.
- Common textbook-style definition: ML studies computer algorithms that **learn from data** and **generalize** to unseen data, instead of following hard-coded instructions.
- In practice:
  - We observe example pairs of input and output.
  - We construct a model that captures regularities in the data.
  - We use the model to make predictions or decisions for new inputs.

# Machine Learning vs. Traditional Programming

## Traditional programming

- Developer writes explicit rules.
- Program + input data  $\Rightarrow$  output.
- Works well when rules are clear and stable.

## Machine learning

- Developer provides data and a learning algorithm.
- Algorithm learns rules (model) from data.
- Useful when rules are complex or unknown.

*One program (the learning algorithm) can be reused to solve many problems by changing only the data.*

## A motivating example: game retention (1/3)

- Imagine you launched a mobile game.
- Some players:
  - Play for a few hours and never return.
  - Others come back every day and even pay for in-game items.
- Business question:

How can we learn the behaviour of players and design strategies to keep them playing?

## A motivating example: game retention (2/3)

- Step 1: **Collect data**
  - For each player, record features:
    - Total play time.
    - Levels completed.
    - Number of friends invited.
    - In-game purchases, etc.
  - Record outcome:
    - 1 if the player is still active after 7 days.
    - 0 otherwise.
- Each player becomes one data point in our dataset.

## A motivating example: game retention (3/3)

- Step 2: **Train an ML model**
  - Input: player features.
  - Output: probability that the player will remain active.
- Step 3: **Use the model**
  - Identify at-risk players (low predicted probability).
  - Offer them a special item or bonus (e.g., powerful sword).
  - Measure whether the intervention improves retention.
- This is a typical **supervised classification** task.

# How are things learned? Memorization

- **Memorization**
  - Accumulation of individual facts:
    - “Player A stayed 10 days.”
    - “Player B left after 2 days.”
  - Limited by:
    - Time to observe all possible situations.
    - Memory to store all examples.
- Pure memorization does **not** help us make good predictions for new players.

# How are things learned? Generalization

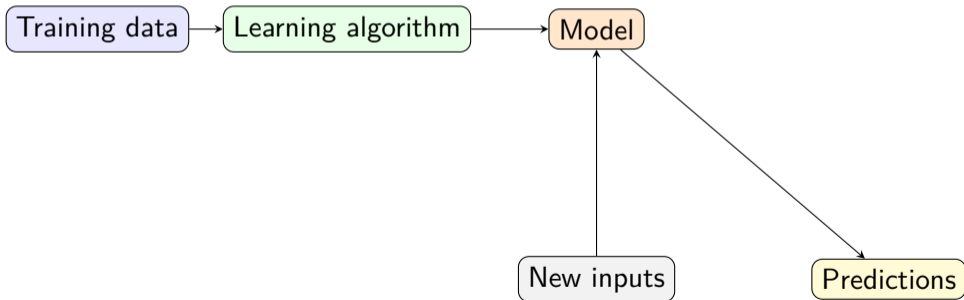
- **Generalization**
  - Deduce new facts from old facts.
  - Look for regularities, patterns and structure in the data.
  - Essentially a **predictive activity**.
- Assumption: **the past is informative about the future**.
- Goal of ML: build programs that generalize well from training data to unseen data.

# Declarative vs. Imperative knowledge

- **Declarative knowledge**
  - Facts: “this email is spam”, “this image contains a cat”.
  - Easy to store, hard to use directly for new cases.
- **Imperative knowledge**
  - Procedures: how to decide if an email is spam.
  - ML aims to infer such procedures from many labelled examples.
- We are interested in programs that infer **useful procedures** from data.

# Basic paradigm of Machine Learning

- 1 Observe a set of examples: **training data**.
- 2 Infer an internal model of the process that generated the data.
- 3 Use this model to make predictions on **test data** (unseen examples).



# Two main variations on the paradigm

## Supervised learning

- Training data: feature/label pairs  $(x_i, y_i)$ .
- Goal: learn a function that predicts  $y$  for a new  $x$ .
- Tasks: classification, regression, sequence labelling.

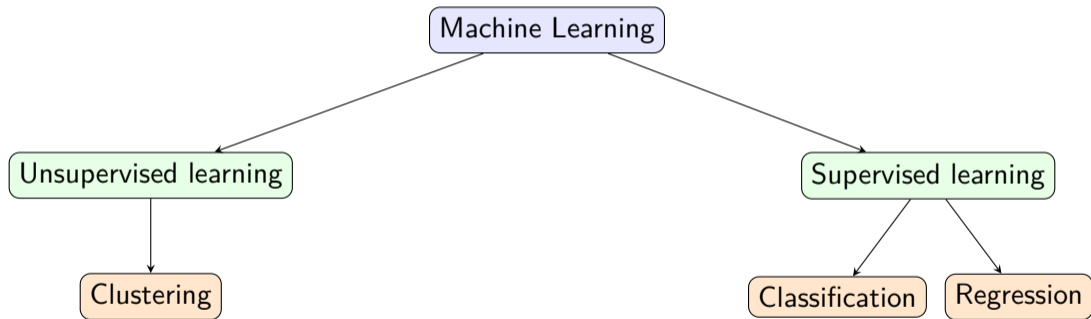
## Unsupervised learning

- Training data: feature vectors  $x_i$  without labels.
- Goal: discover structure, e.g., clusters, latent factors.
- Tasks: clustering, dimensionality reduction, density estimation.

## Other learning paradigms (high-level view)

- **Semi-supervised learning**
  - Use both labelled and unlabelled data to improve performance.
- **Reinforcement learning**
  - Agent interacts with an environment.
  - Receives rewards and learns a policy to maximize long-term reward.
- **Online learning**
  - Data arrives sequentially; the model is updated continuously.
- In this lecture we focus on basic supervised and unsupervised intuition.

# Machine learning methods



## Typical ML workflow

- 1 Formulate the problem and define the goal.
- 2 Collect and clean data.
- 3 Represent objects with feature vectors.
- 4 Choose a model family and objective function.
- 5 Train the model using an optimization algorithm.
- 6 Validate and tune hyperparameters.
- 7 Evaluate on test data.
- 8 Deploy and monitor the model in production.

## What all ML methods require

- A suitable **training dataset** and evaluation method.
- A **representation** of the features.
- A **distance** or similarity measure between feature vectors (for many methods).
- An **objective function** (loss) and possibly constraints.
- An **optimization method** to learn model parameters.

# What is a feature?

- A **feature** is a measurable property of an object.
- Examples for an email:
  - Number of exclamation marks.
  - Presence of certain keywords.
  - Sender domain, time of day.
- We group features into a **feature vector**:

$$x = (x_1, x_2, \dots, x_d) \in \mathbb{R}^d.$$

- Good features make patterns easier to learn and generalize.

# Feature engineering

- The art of constructing informative features from raw data.
- Goals:
  - Facilitate generalization.
  - Avoid overfitting by reducing noise and irrelevant dimensions.
  - Encode prior knowledge about the problem.
- Strategies:
  - Transform variables (log, normalization, binning).
  - Create new features (ratios, interactions, counts).
  - Select a subset of useful features.

## Reptile classification example — setting

- Suppose we want to classify animals into:
  - **Reptile** vs. **Not reptile**.
- We start with candidate features:
  - Egg-laying (yes/no).
  - Has scales (yes/no).
  - Poisonous (yes/no).
  - Cold-blooded (yes/no).
  - Number of legs (integer).
- Our labels come from a biology expert.

# Reptile classification — initial model

- First attempt:

## Rule 1

An animal is a reptile if it:

- lays eggs,
  - has scales,
  - is poisonous,
  - is cold-blooded,
  - has no legs.
- 
- Obviously this is too strict: many reptiles will not satisfy all conditions.
  - We need to refine our feature set and rules.

## Reptile classification — refining the rule (1/3)

- Consider cobra, rattlesnake and boa constrictor.
- We observe:
  - All have scales, are cold-blooded and have no legs.
  - Egg-laying and poisonous may vary.
- Updated model:

### Rule 2

Reptile if:

- has scales,
- is cold-blooded,
- has no legs.
- This rule correctly classifies these snakes as reptiles.

## Reptile classification — refining the rule (2/3)

- Now we add chicken, which:
  - lays eggs,
  - is warm-blooded,
  - has two legs,
  - has no scales.
- Rule 2 correctly identifies chicken as **not** a reptile.
- Our model is improving but remains incomplete.

## Reptile classification — refining the rule (3/3)

- Add alligator and dart frog.
- Revised idea:

### Rule 3

Reptile if:

- has scales,
- is cold-blooded,
- has 0 or 4 legs.
- Problem:
  - Some non-reptiles (e.g., certain fish) may also satisfy these features.
  - Difficult to find a simple exact rule with our limited features.

## Imperfect but useful models

- With current features, we may obtain a rule that:
  - rarely misses true reptiles (**few false negatives**),
  - but may mistakenly label some animals as reptiles (**false positives**).
- In many applications this trade-off is acceptable.
- Feature engineering often aims at:
  - reducing false positives,
  - without increasing false negatives too much.

## Feature engineering lessons from reptiles

- Features must be **informative** for the task.
- Too few features  $\Rightarrow$  cannot separate classes.
- Too many irrelevant features  $\Rightarrow$  noise and overfitting.
- Sometimes we must change representation:
  - continuous vs. binary,
  - absolute counts vs. ratios,
  - domain-specific transformations.

## From animals to feature vectors

- Encode animals as binary / integer feature vectors.
- Example:

rattlesnake = (1, 1, 1, 1, 0)

boa constrictor = (0, 1, 0, 1, 0)

dart frog = (1, 0, 1, 0, 4)

- We now need a way to measure how similar two animals are.

# The Minkowski distance

## Definition

For two feature vectors  $x_1, x_2 \in \mathbb{R}^d$  and  $p \geq 1$ ,

$$\text{dist}_p(x_1, x_2) = \left( \sum_{k=1}^d |x_{1k} - x_{2k}|^p \right)^{1/p}.$$

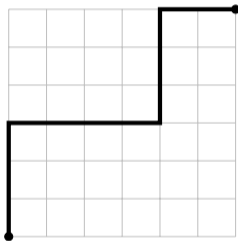
- $p = 1$ : **Manhattan distance.**
- $p = 2$ : **Euclidean distance.**
- Larger  $p$  emphasize larger coordinate differences.

## Manhattan distance: taxicab geometry

- Imagine a city with a grid of streets.
- Distance is the number of blocks you must walk:

$$d_1(x, y) = \sum_k |x_k - y_k|.$$

- Many shortest paths may exist.
- Often appropriate when different dimensions are not directly comparable.



## Euclidean distance

- Standard geometric distance in  $\mathbb{R}^d$ :

$$d_2(x, y) = \sqrt{\sum_k (x_k - y_k)^2}.$$

- Corresponds to the length of the straight line segment between two points.
- Often the default in many ML algorithms (e.g.,  $k$ -means,  $k$ -NN).
- Sensitive to scale of each dimension.

## Which distance says what?

- Consider a simple 2D grid with three points: a circle, a star and a cross.
- Question: is the circle closer to the star or to the cross?
- Using Euclidean distance:
  - cross  $\approx 2.8$
  - star  $\approx 3.0$
- Using Manhattan distance:
  - cross = 4
  - star = 3
- Different metrics can lead to different nearest neighbours.

## Scaling issues: adding the alligator

- Recall:

dart frog = (1, 0, 1, 0, 4),

boa = (0, 1, 0, 1, 0),

alligator = (1, 1, 0, 1, 4).

- Legs feature ranges from 0 to 4, others from 0 to 1.
- The “legs” dimension dominates Euclidean distance.
- Alligator may incorrectly appear closer to dart frog than to boa.

## Feature scaling and normalization

- To avoid dominated dimensions we often:
  - rescale each feature to comparable ranges,
  - or standardize to zero mean and unit variance.
- After scaling, distances better reflect meaningful similarity.
- In the reptile example, we might:
  - convert “legs” to a binary feature (has legs / no legs),
  - or divide by maximum number of legs.

## Using binary features

- Encode presence/absence rather than counts:

rattlesnake = (1, 1, 1, 1, 0)

boa = (0, 1, 0, 1, 0)

dart frog = (1, 0, 1, 0, 1)

alligator = (1, 1, 0, 1, 1)

- Now alligator is closer to the snakes than to the frog:
  - Distances reflect the intended biological similarity.
- **Feature engineering matters!**

## Summary: similarity measures

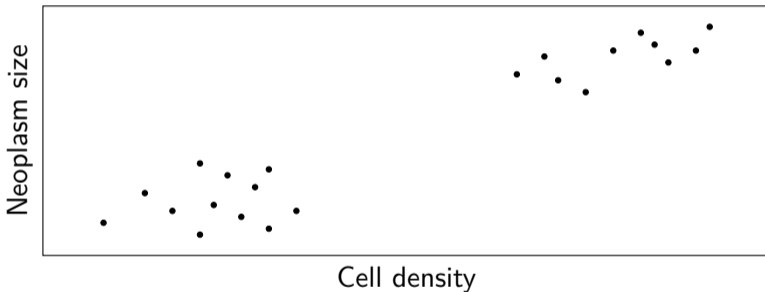
- Many ML methods rely on a notion of distance or similarity.
- Choice of metric interacts with feature representation.
- Key points:
  - Use appropriate scaling/normalization.
  - Consider Manhattan vs. Euclidean for different data types.
  - Binary / categorical features often require specialized measures.
- Poor choices here can severely hurt model performance.

## Breast cancer example — data

- Each example: breast tumor (neoplasm).
- Features:
  - Neoplasm size.
  - Cell density.
- Two underlying types:
  - **Benign.**
  - **Malignant.**
- Initially assume we do **not** know which point belongs to which type.

## Unlabelled data: scatter plot

Distribution of neoplasm size vs. cell density



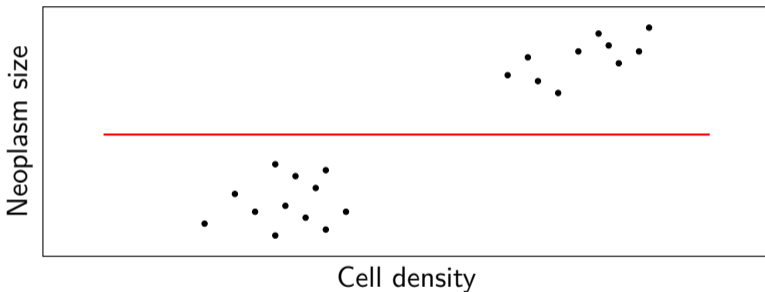
Two visually separable clusters suggest two neoplasm types.

## Task: clustering examples into groups

- **Clustering:**
  - Group examples such that points in the same cluster are similar.
  - Points in different clusters are dissimilar.
- We need:
  - A distance measure (e.g., Euclidean).
  - A number of clusters  $K$  (here, assume  $K = 2$ ).
  - An objective, e.g., minimize distance within clusters.
- Algorithms:  $k$ -means, hierarchical clustering, etc. (not covered in detail here).

## Similarity based on neoplasm size only

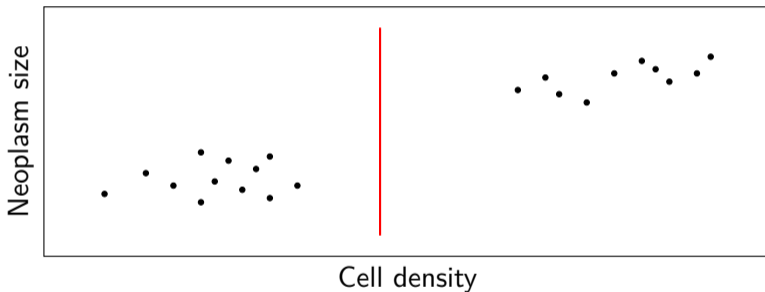
Vertical split using size only



Horizontal line splits data purely by size.

## Similarity based on cell density only

Vertical split using density only



Vertical line splits data purely by cell density.

## Using both attributes

- A better similarity notion uses both size and cell density.
- Clusters correspond to two dense regions in 2D space.
- Many clustering algorithms implicitly do this by:
  - representing each point as (density, size),
  - minimizing within-cluster distances in this 2D space.
- This illustrates again the importance of feature representation.

## If the data were labelled

- Now suppose each point is labelled:
  - blue: benign,
  - red: malignant.
- Task becomes **supervised classification**.
- Goal: find a **decision boundary** that separates the two classes.
- In 2D this is a line or curve; in higher dimensions, a surface.

## Finding classifier surfaces

- Given labelled training data, we want a function  $f(x)$  such that:
  - $f(x) = 0$  for benign,
  - $f(x) = 1$  for malignant (for example).
- The **classifier surface** is where the model switches prediction.

$$\{x : f(x) = 0.5\}$$

(for probabilistic models).

- Constraints:
  - Surface should not be too complex (avoid overfitting).
  - Some trade-off between false positives and false negatives.

## Adding new data points

- After training a classifier on past tumors, we receive a new patient.
- We measure:
  - neoplasm size,
  - cell density.
- We feed the feature vector into the model:
  - obtain a predicted probability of malignancy,
  - possibly use a threshold (e.g., 0.5) to decide the label.
- This is the standard way ML models support decision making.

# Clustering vs. classification

## Clustering (unsupervised)

- Input: unlabelled data.
- Goal: discover groups based on similarity.
- Output: cluster assignments.
- No direct notion of “correct” label.

## Classification (supervised)

- Input: labelled data.
- Goal: learn mapping from features to labels.
- Output: predicted labels or probabilities.
- Performance measured against ground truth labels.

## Sources of error

- **Overlapping classes**
  - Benign and malignant regions not perfectly separable.
- **Noisy measurements**
  - Size and density may be measured with error.
- **Limited features**
  - Important biological factors not captured.
- **Limited data**
  - Model may not see enough examples of rare cases.

# Statistical view of machine learning

- Assume input  $X$  and output  $Y$  follow an unknown joint distribution  $P(X, Y)$ .
- Training and test data are drawn i.i.d. from this distribution.
- Goal of learning:
  - Approximate either the conditional probability  $P(Y | X)$ ,
  - or a decision function  $f(X)$  that predicts  $Y$ .
- This motivates the term **statistical learning**.

# Hypothesis space and model

- We restrict ourselves to a set of candidate functions:

$$\mathcal{H} = \{f_{\theta} : \theta \in \Theta\},$$

called the **hypothesis space**.

- Examples:
  - Linear models.
  - Decision trees.
  - Neural networks of a given architecture.
- Learning  $\approx$  choosing a good  $\theta$  based on data.

## Loss, risk and empirical risk

- **Loss function**  $\ell(f(x), y)$  measures how bad a prediction is.
  - Classification: 0–1 loss, cross-entropy.
  - Regression: squared loss, absolute loss.
- **Risk** (expected loss):

$$R(f) = \mathbb{E}_{(X,Y) \sim P}[\ell(f(X), Y)].$$

- We cannot compute  $R(f)$  directly (because  $P$  is unknown).
- Instead we minimize the **empirical risk** on training data.

# Overfitting and model complexity

- Minimizing empirical risk alone may cause **overfitting**.
- Overfitting:
  - Model fits training data extremely well,
  - but performs poorly on unseen test data.
- Typically happens when:
  - Model is too complex relative to the amount of data.
  - Features contain too much noise.
- We need **regularization** and good model selection.

## Regularization idea (informal)

- Add a penalty for model complexity:

$$\text{objective} = \text{empirical risk} + \lambda \cdot \text{complexity}(f).$$

- Examples:
  - Penalize large weights in linear models.
  - Penalize deep or unpruned decision trees.
- Balances fit to training data and ability to generalize.

# Model evaluation and selection

- Split data into:
  - Training set.
  - Validation set (for model/hyperparameter selection).
  - Test set (for final evaluation).
- Use techniques such as:
  - $k$ -fold cross-validation.
  - Stratified sampling for imbalanced classes.
- Always report performance on data not used for training.

## Key takeaways (1/2)

- Machine learning builds models that **learn from data** and generalize to new cases.
- ML is a subset of AI, and deep learning is a subset of ML.
- Basic paradigm:
  - collect data,
  - choose representation and model,
  - train and evaluate.
- Feature engineering and distance metrics strongly influence results.

## Key takeaways (2/2)

- Supervised vs. unsupervised learning:
  - **Classification**: predict discrete labels.
  - **Regression**: predict continuous values.
  - **Clustering**: group similar examples without labels.
- Statistical learning view:
  - loss, risk, empirical risk,
  - overfitting and regularization.
- Always keep in mind:

The quality of data and features often matters more than the choice of algorithm.

## Next lectures

- **Supervised Learning I**
  - $k$ -nearest neighbours.
  - Linear regression, logistic regression.
- **Supervised Learning II**
  - Decision trees and ensembles.
  - Model evaluation metrics.
- **Unsupervised Learning**
  - $k$ -means and other clustering methods.
  - Dimensionality reduction.

## Suggested reading

- T. Mitchell, *Machine Learning*.
- K. Murphy, *Probabilistic Machine Learning*.
- A. Ng, *Machine Learning* (online course).
- Articles:
  - “Machine learning, explained” (MIT Sloan).
  - “What is Machine Learning?” (IBM).

Questions?

Thank you for your attention.

Questions and discussion.