



# **MACHINE LEARNING PARADIGMS IN BIOINFORMATICS**

**Presented By**

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A background network diagram consisting of numerous small grey dots (nodes) connected by thin grey lines (edges). The nodes are distributed across the slide, with some clusters and many isolated connections, creating a web-like structure. The overall color scheme is light grey and white.

# Introduction

In this short presentation, we would like to talk about and explore the various ways popular machine learning algorithms can help us extract useful and actionable data from seemingly ordinary datasets with its implementation. We would also explore some popular machine learning algorithms and how they work.

# Outline

- **Introduction**
- **What is Machine Learning?**
- **Types of Machine Learning Algorithms**
- **Implementation of a Working Model**
- **Planned Future Work**
- **Conclusion**
- **Acknowledgement**
- **References**

# What is Machine Learning?

Some popular definitions are as follows

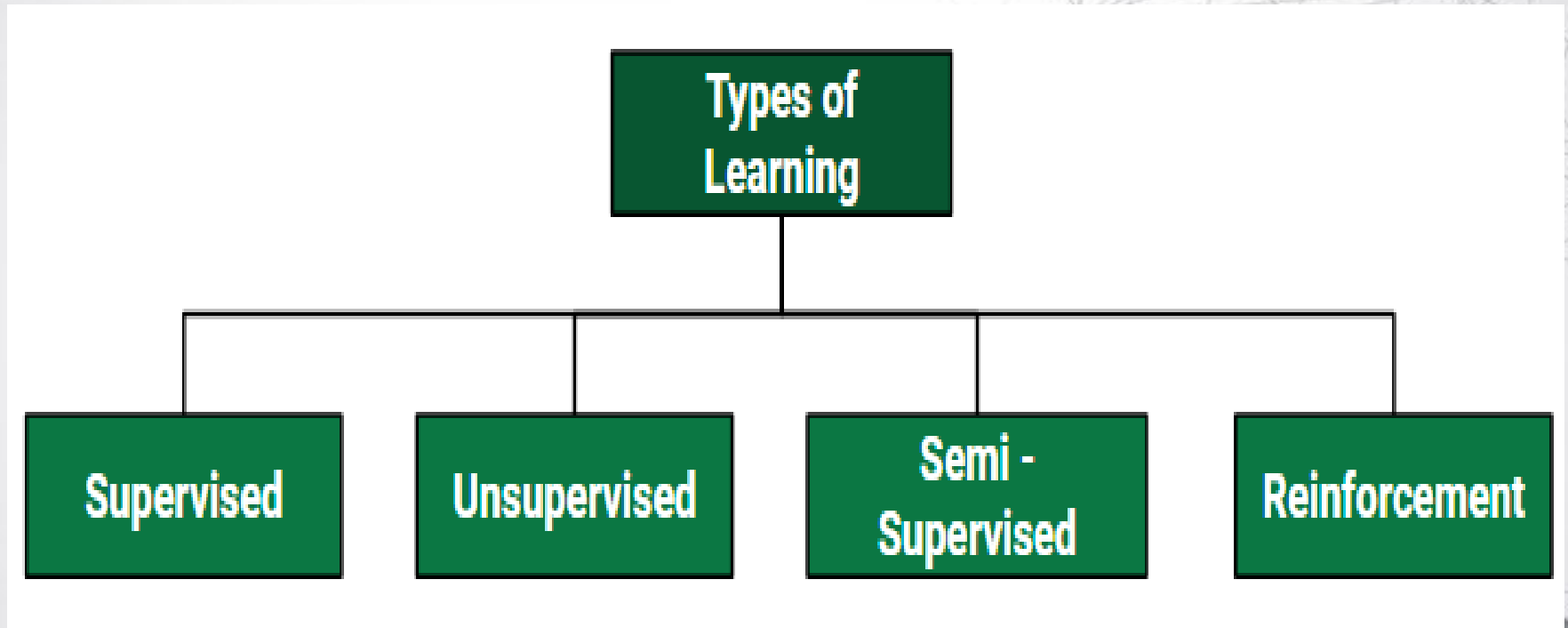
- **Arthur Samuel(1959)**: Field of study that gives computers the ability to learn without being explicitly programmed.
- **Tom Mitchell(1998)**: A computer program is said to learn from experience(**E**) with respect to some task (**T**) and some performance measure (**P**) , if its performance on **T** , as measured by **P** , improves with experience **E**.

The background of the slide features a complex, abstract network of interconnected nodes and lines, resembling a neural network or a data graph. The nodes are represented by small circles of varying sizes and shades of gray, while the lines are thin, light gray connections between them. The overall aesthetic is clean and modern, with a light gray background.

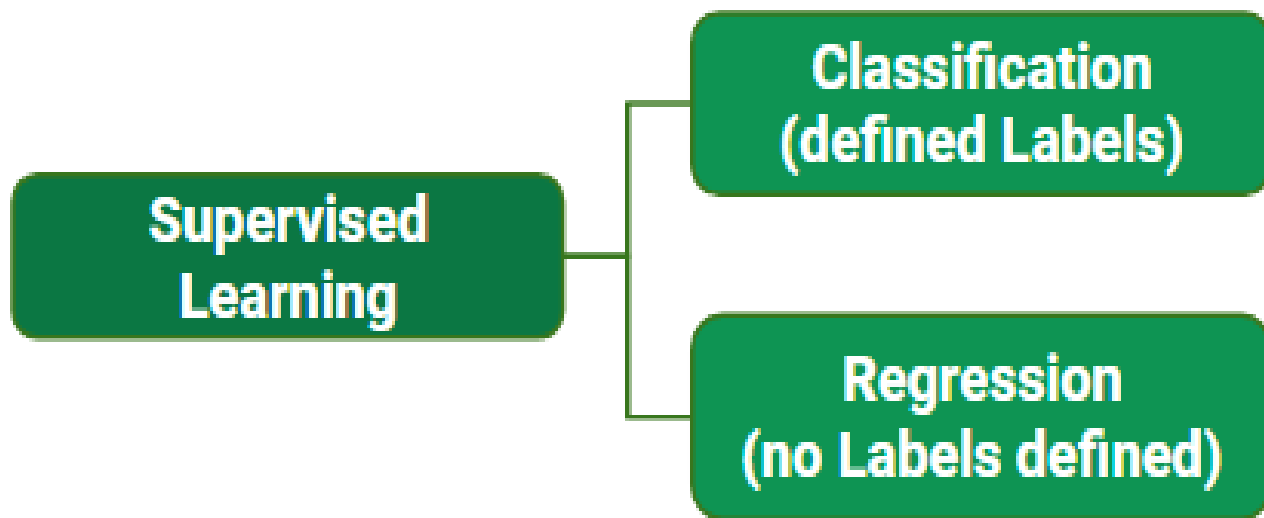
# Types of Machine Learning Algorithms



# Machine Learning Algorithms



# Supervised Learning



## Supervised Learning

- Supervised learning is when the model is getting trained on a labelled dataset.
- **Labelled** dataset is one which have both input and output parameters. In this type of learning both training and validation datasets are labelled
- Example:  
Linear Regression  
Logistic Regression



The background of the slide is a light gray with a complex network of thin, dark gray lines connecting various sized dark gray dots, creating a web-like or molecular structure.

## Types of Supervised Learning

### 1. Classification Algorithms

These are the kinds of algorithms with produces defined labels or discrete values as output.

Ex. - Logistic Regression, Neural Networks

## Types of Supervised Learning

**Logistic Regression** is a Machine Learning classification algorithm that is used to predict the probability of a categorical dependent variable.

In logistic regression, the dependent output variable is a binary variable that contains data coded as 1 (yes, success, etc.) or 0 (no, failure, etc.).

# Types of Supervised Learning

## 1. Regression Algorithms

Here, we fit a curve / line to the data points, in such a manner that the differences between the distances of data points from the curve or line is minimized. We obtain a continuous output variable.

Ex. - Linear Regression, Support Vector Regression



## Types of Supervised Learning

**Linear Regression** is a supervised machine learning algorithm where the predicted output is continuous and has a constant slope. It's used to predict values within a continuous range, (e.g. sales, price) rather than trying to classify them into categories (e.g. cat, dog).

## Unsupervised Learning

It is the training of machine using information that is neither classified nor labeled and allowing the algorithm to act on that information without guidance. Here the task of machine is to group unsorted information according to similarities, patterns and differences without any prior training of data.

Ex. K-Means Clustering, Principle Component Analysis



## Types of Unsupervised Learning

**K-Means Clustering** is an iterative clustering algorithm which helps you to find the highest value for every iteration. Initially, the desired number of clusters are selected. In this clustering method, you need to cluster the data points into  $k$  groups.

The output of the algorithm is a group of "labels." It assigns data point to one of the  $k$  groups. In  $k$ -means clustering, each group is defined by creating a centroid for each group, they capture the points closest to them and adds them to the cluster.

The background of the slide features a complex, abstract network diagram. It consists of numerous small, dark grey circular nodes connected by thin, light grey lines. These lines form a dense web of triangles and other geometric shapes, creating a sense of interconnectedness and data flow. The overall aesthetic is clean and modern, typical of a professional presentation on technology or data science.

# **Implementation of Machine Learning for Bioinformatics**

# The Hungarian Heart Disease Dataset

	age	sex	pain	BP	...	slope	vessels	thal	diagnosis
0	29.0	1.0	2.0	120.0	...	1.894231	0.0	5.642857	0.0
1	29.0	1.0	2.0	140.0	...	1.894231	0.0	5.642857	0.0
2	30.0	0.0	1.0	170.0	...	1.894231	0.0	6.000000	0.0
3	31.0	0.0	2.0	100.0	...	1.894231	0.0	5.642857	0.0
4	32.0	0.0	2.0	105.0	...	1.894231	0.0	5.642857	0.0
..	...	...	...	...	...	...	...	...	...
288	52.0	1.0	4.0	160.0	...	1.894231	0.0	5.642857	1.0
289	54.0	0.0	3.0	130.0	...	2.000000	0.0	5.642857	1.0
290	56.0	1.0	4.0	155.0	...	2.000000	0.0	5.642857	1.0
291	58.0	0.0	2.0	180.0	...	2.000000	0.0	7.000000	1.0
292	65.0	1.0	4.0	130.0	...	2.000000	0.0	5.642857	1.0

[293 rows x 14 columns]



# Source Code

```
import pandas as pd
import numpy as np
from sklearn.impute import SimpleImputer
from sklearn.linear_model import LogisticRegression
from sklearn.model_selection import train_test_split

# info about the features used

header_row = ['age', 'sex', 'pain', 'BP', 'chol', 'fbs', 'ecg', \
              'maxhr', 'eiang', 'eist', 'slope', 'vessels', 'thal', 'diagnosis']

# Loading the dataset

dataset = pd.read_csv("processed.hungarian.data")

# Replacing the missing values with the Mean value of its column

dataset = dataset.replace("?", np.NaN)
imp = SimpleImputer(missing_values=np.NaN, strategy='mean')
imp.fit(dataset)
dataset = pd.DataFrame(imp.transform(dataset))
dataset.columns = header_row

# Extracting the features and the results

X, y = dataset.iloc[:, :-1], dataset.iloc[:, -1]

# Splitting the dataset into training and test sets to remove bias

X_train, X_test, y_train, y_test = train_test_split(X, y, \
```

## Source Code

```
X, y = dataset.iloc[:, :-1], dataset.iloc[:, -1]

# Splitting the dataset into training and test sets to remove bias

X_train, X_test, y_train, y_test = train_test_split(X, y, \
    test_size=1/3, random_state=2)

# Using Logistic Regression as the algorithm for classification

model = LogisticRegression()
model.fit(X_train, y_train)
score = model.score(X_test, y_test)

# The output suggests the % of times the model can correctly
# predict the outcome i.e. the characteristics would lead
# to a heart disease or not

print(round(score*100,2), "% Accuracy") # Output: 83.67 % Accuracy
```



# **Future Work**

1. More visually descriptive
2. Improving the accuracy
3. Cleaning the data
4. Performance measures.
5. Analyzing the original 1987 Dataset

A background network diagram consisting of numerous grey dots of varying sizes connected by thin, light grey lines, creating a complex web-like structure across the entire slide.

## Conclusion

As we can observe, there is plenty of scope in the field of Bio-Medical Informatics for us, as budding Computer Science students to explore and implement ground-breaking algorithms & machine learning pipelines to extract meaningful inferences that can directly help people. It is our conviction that we would contribute to this ever-growing field and improve our initial understandings.

The background of the slide features a complex, abstract network of thin grey lines connecting various-sized grey dots of different shades. These dots are scattered across the entire frame, creating a web-like or molecular structure that serves as a subtle backdrop for the text.

## Acknowledgement

We would like to thank our mentor Mr. Sabyasachi Mukherjee Sir, for his continuous guidance during the preparation for this topic.

We would like to thank the Head of Computer Science & Engineering department, Dr. Monish Chatterjee for providing us with the opportunity to explore such technological fields.

# References

1. Dataset was taken from the popular UCI Machine Learning Repository

<http://archive.ics.uci.edu/ml/machine-learning-databases/heart-disease/>

2. The Python in-built machine learning module "Scikit-Learn"

<https://scikit-learn.org/stable/>

3. Understanding of the Algorithms and various Data Science constructs from the popular MOOC by Andrew Ng

<https://www.coursera.org/learn/machine-learning-with-python>