

Unit 4

Graphical Approaches & Grammatical Inference in Syntactic Pattern Recognition

Graphical Approaches:

Definition: Graphical approaches in syntactic pattern recognition involve using graph-based structures to represent and analyze patterns within data. These approaches leverage the power of graphs to model relationships between different components of a pattern, allowing for a more flexible and comprehensive representation of complex structures. The main idea is to use nodes and edges in a graph to encapsulate the components and their relationships in the pattern.

Graphs are particularly useful in syntactic pattern recognition because they can represent hierarchical and relational information effectively. The structure of the graph can be used to define a grammar that describes the possible patterns within a given domain.

Example: Consider the problem of recognizing a simple handwritten character, such as the letter "A."

- **Graph Representation:**
 - **Nodes:** Each stroke or line segment of the character "A" can be represented as a node in a graph.
 - **Edges:** The connections between these strokes (e.g., the angles or intersection points) can be represented as edges in the graph.

In this graphical model, the relationships between the different strokes of the letter "A" (like the horizontal stroke connecting the two slanting lines) are captured. The graph can

then be used to match and recognize other instances of the letter "A" by checking if the graph structure matches the defined pattern (grammar) of an "A."

- **Grammar in Graphical Approach:** The grammar in this context could be a set of rules that describe how the nodes (strokes) should be connected by edges (angles, intersections) to form a valid letter "A." For instance, the grammar might specify that two slanted lines must be connected by a horizontal line in the middle to form the letter.

Use Cases:

- **Computer Vision:** Recognizing shapes, objects, or letters by modeling them as graphs.
- **Biometric Recognition:** Graphs can be used to model and recognize complex patterns such as fingerprints or facial features.
- **Structural Pattern Recognition:** Graphical approaches can model and recognize patterns in molecular structures in chemistry and biology.

Graphical approaches in syntactic pattern recognition provide a powerful tool for modeling and recognizing complex patterns by representing them as graphs and applying grammatical inference to match and analyze the structures.

Graph Based Structural Representation:

Definition: Graph-based structural representation is a method used in syntactic pattern recognition where patterns are represented as graphs. This approach is particularly effective in capturing the structural and relational information inherent in complex patterns. In a graph-based representation, the components of a pattern are depicted as nodes, and the relationships or interactions between these components are represented as edges connecting the nodes.

Key Concepts:

- **Nodes:** Represent the basic elements or components of the pattern (e.g., parts of an object, characters in a string, etc.).
- **Edges:** Represent the relationships or connections between the components (e.g., spatial relationships, connectivity, dependencies, etc.).

- **Graph Grammar:** A set of rules that define how nodes and edges can be combined to form valid patterns.

This method is particularly useful when the patterns have a hierarchical or relational structure that is difficult to capture with traditional feature-based approaches.

Example: Consider a scenario where you want to recognize different types of **chemical molecules**.

- **Nodes:** Each atom in the molecule can be represented as a node in the graph.
- **Edges:** The bonds between atoms can be represented as edges connecting the nodes.

In this graph-based structural representation, the molecule's structure is captured by the graph, where the types of atoms and their bonding relationships are explicitly represented.

- **Graph Grammar:** The graph grammar could define the valid types of bonds (e.g., single, double, triple bonds) and how different atoms can be connected according to the rules of chemistry. This grammar helps in recognizing and differentiating between different molecules by analyzing the graph structure.

Applications:

- **Molecular Structure Analysis:** Graph-based representations are used to analyze and recognize molecular structures in chemistry.
- **Scene Understanding in Computer Vision:** Objects and their spatial relationships in an image can be represented as a graph, enabling more robust scene understanding.
- **Handwriting Recognition:** Individual strokes in handwritten characters can be represented as nodes, with edges representing the connections between strokes.

Advantages:

- **Flexibility:** Graph-based representation can model a wide variety of patterns with complex internal structures.

- **Expressiveness:** Graphs can capture both the components of a pattern and the relationships between them, providing a richer representation than linear or feature-based models.

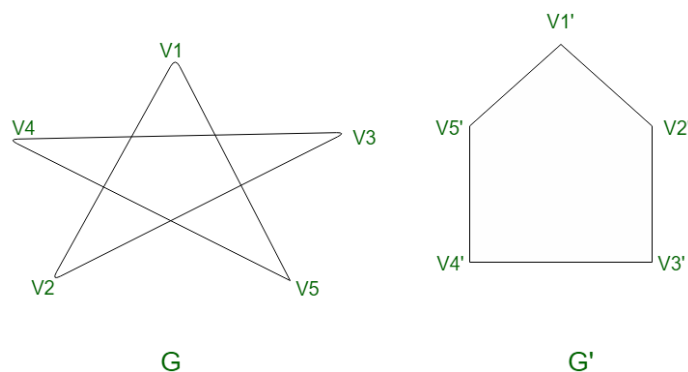
Disadvantages:

- **Complexity:** Graph-based methods can be computationally intensive, especially for large and complex graphs.
- **Matching and Parsing Challenges:** Comparing and parsing graphs to recognize patterns can be more complex than matching simpler feature vectors.

Graph Isomorphism:

In Graph Theory, What is Graph Isomorphism?

A graph isomorphism is a bijection between the vertex sets of two graphs that preserves the adjacency relationship. In other words, two graphs G and H are isomorphic if there is a one-to-one correspondence between their vertices such that two vertices are adjacent in G if and only if their corresponding vertices are adjacent in H .



In Pattern Recognition:

In computer vision and pattern recognition, graph isomorphism are used to match patterns and shapes.

Definition: Graph isomorphism in syntactic pattern recognition refers to the problem of determining whether two graphs are isomorphic, meaning they are structurally identical in terms

of their connectivity, even if their node labels or the specific ordering of nodes and edges are different. In other words, two graphs are isomorphic if there is a one-to-one correspondence between their nodes and edges such that the connectivity (relationships between nodes) is preserved.

Formal Definition: Two graphs $G_1 = (V_1, E_1)$ and $G_2 = (V_2, E_2)$ are said to be isomorphic if there exists a bijection $f : V_1 \rightarrow V_2$ such that for every pair of nodes $u, v \in V_1$, there is an edge between u and v in G_1 if and only if there is an edge between $f(u)$ and $f(v)$ in G_2 .

Example:

Scenario: Software Component Comparison

In software engineering, consider a scenario where you need to verify if two different implementations of a software module are functionally equivalent. These modules can be represented as control flow graphs (CFGs), where:

- **Nodes** represent basic blocks of code (e.g., sequences of instructions or statements without any branches).
- **Edges** represent the control flow between these blocks (e.g., conditional branches, loops, function calls).

Example Graphs:

- **Graph 1 (Module A):**
 - Nodes: N_1, N_2, N_3, N_4
 - Edges: $(N_1 \rightarrow N_2), (N_2 \rightarrow N_3), (N_3 \rightarrow N_4), (N_4 \rightarrow N_2)$
- **Graph 2 (Module B):**
 - Nodes: M_1, M_2, M_3, M_4
 - Edges: $(M_1 \rightarrow M_2), (M_2 \rightarrow M_4), (M_4 \rightarrow M_3), (M_3 \rightarrow M_2)$

Observations:

- The nodes in both graphs represent basic blocks of code, though the nodes have different labels in each graph.
- The edges in both graphs represent the control flow between these blocks, which might appear differently at first glance.

Graph Isomorphism:

To determine if Graph 1 and Graph 2 are isomorphic, you would check if there's a mapping between the nodes of one graph to the nodes of the other such that the edges correspond.

- **Mapping:**
 - Map N_1 in Graph 1 to M_1 in Graph 2
 - Map N_2 in Graph 1 to M_2 in Graph 2
 - Map N_3 in Graph 1 to M_3 in Graph 2
 - Map N_4 in Graph 1 to M_4 in Graph 2

With this mapping, every edge in Graph 1 corresponds exactly to an edge in Graph 2, preserving the control flow structure between the basic blocks of code.

Conclusion:

Since there is a one-to-one correspondence between the nodes and edges of Graph 1 and Graph 2, the graphs are **isomorphic**. This implies that the two software modules, despite having different internal structures (such as different variable names, order of operations, etc.), are functionally equivalent in terms of their control flow.

Application:

In software engineering, graph isomorphism can be used to:

- **Code Refactoring:** Ensure that a refactored version of a program is functionally equivalent to the original by comparing their control flow graphs.

- **Software Plagiarism Detection:** Detect cases where the structure of a program has been copied and slightly modified by comparing CFGs of different programs.
- **Optimization Validation:** Verify that optimized code is functionally equivalent to the original by comparing their CFGs.

A Structured Strategy to Compare Attribute Graphs:

Definition: Attribute graphs are an extension of basic graphs where nodes and edges have additional information (attributes) associated with them. Comparing attribute graphs involves not only checking the structural isomorphism (i.e., the connectivity of nodes and edges) but also ensuring that the corresponding attributes match.

Strategy to Compare Attribute Graphs:

1. Structural Comparison (Graph Isomorphism):

- First, perform a structural comparison to check if the two graphs are isomorphic, meaning their nodes can be mapped one-to-one while preserving the connectivity of edges.

2. Attribute Matching:

- For each pair of corresponding nodes and edges in the isomorphism mapping, compare their attributes (e.g., labels, weights, colors).
- Ensure that the attributes match according to the specific rules or criteria defined for the application.

3. Similarity Metrics (Optional) Matching:

For each pair of corresponding :

- If the graphs are not exactly isomorphic but share a similar structure, a similarity metric can be used to quantify the degree of similarity, taking into account both structure and attributes.

Example:

Consider two networks representing social connections where:

- Nodes represent people.
- Edges represent friendships.
- Node Attributes: Each node has an attribute like "age."
- Edge Attributes: Each edge has an attribute like "frequency of interaction."

Graph 1:

- Nodes: A (Age 30), B (Age 25), C (Age 28)
- Edges: A-B (Frequency: High), B-C (Frequency: Medium)

Graph 2:

- Nodes: D (Age 30), E (Age 25), F (Age 28)
- Edges: D-E (Frequency: High), E-F (Frequency: Medium)

Comparison Process:

1. Structural Isomorphism:

Map A to D, B to E, and C to F.

The edges match in connectivity, so the graphs are structurally isomorphic.

2. Attribute Matching:

- Compare the node attributes: A's age matches D's, B's age matches E's, and C's age matches F's.
- Compare the edge attributes: The frequency of interaction between A-B matches D-E, and B-C matches E-F.
- Since both the structural and attribute comparisons match, the two social networks are considered equivalent in this context.

Application: This strategy can be applied in areas like social network analysis, where not only the structure but also the characteristics of the connections are important for recognizing patterns and equivalences.

(Edge A-B (Frequency: High):
<ul style="list-style-type: none">• This means that the interaction between entities A and B happens frequently. For instance, in a social network, A and B might communicate with each other daily.
Edge B-C (Frequency: Medium):
<ul style="list-style-type: none">• This means that the interaction between entities B and C happens at a moderate rate. In the same social network example, B and C might communicate weekly.)

Other Attributed Graph Distance or Similarity measures:

In pattern recognition, comparing attributed graphs (graphs with additional information or attributes on nodes and edges) often requires specialized distance or similarity measures. These measures help quantify how similar or different two attributed graphs are, considering both their structure and their attributes. Here are some commonly used distance or similarity measures for attributed graphs:

1. Graph Edit Distance (GED):

- **Description:** GED measures the minimum number of edit operations (e.g., insertion, deletion, substitution of nodes or edges) required to transform one graph into another. The edits can also take into account the attributes of nodes and edges.
- **Usage:** GED is widely used in various applications such as error-tolerant graph matching, where small differences between graphs are expected.
- **Example:** In a network of roads (represented as a graph), GED can help compare different road layouts by considering both the connections between intersections and the types of roads (attributes).

2. Subgraph Isomorphism:

- **Description:** This method checks whether one graph is a subgraph of another, meaning one graph can be mapped into a part of the other graph while preserving the structure and attributes.

- **Usage:** Sub graph isomorphism is useful in applications where you need to find patterns or motifs within a larger graph, such as in bioinformatics to identify specific molecular structures within a larger network.
- **Example:** Identifying a common sub network in different social networks where certain attributes like "job title" or "communication frequency" match.

3. Hamming Distance (for labeled graphs):

- **Description:** Hamming distance counts the number of different node or edge labels between two graphs. It's a simple measure used when the graphs are of the same size and structure but have different labels.
- **Usage:** This is applicable in cases where only the labels or attributes differ between graphs, such as in comparing different configurations of the same network.
- **Example:** Comparing two graphs representing organizational structures where nodes represent employees and labels represent their roles.

4. Graph Spectral Distance:

- **Description:** Spectral methods use the eigenvalues of graph-related matrices (like the adjacency matrix) to compute a distance between graphs. The difference in eigenvalues or eigenvectors provides a measure of similarity or difference between the graph structures.
- **Usage:** Spectral methods are often used in graph clustering and in cases where the global structure of the graph is important.
- **Example:** Comparing the overall structure of two large social networks by analyzing the connectivity patterns rather than individual nodes or edges.

Learning Via Grammatical Inference:

Definition: Grammatical inference, also known as grammar induction, is the process of learning grammars and languages from a set of observations, typically a finite set of strings or patterns. In the context of syntactic pattern recognition, this involves learning the underlying grammatical rules that generate a set of observed patterns, which can then be used to recognize or generate new patterns.

Key Concepts:

1. **Grammar:** A formal set of rules that describe how strings or patterns are generated in a language. Grammars are often represented using formal languages like context-free grammars (CFGs) or regular grammars.

What is Context-Free Grammar?

Context Free Grammar is formal grammar, the syntax or structure of a formal language can be described using context-free grammar (CFG), a type of formal grammar. The grammar has four tuples: (V,T,P,S).

V - It is the collection of variables or non-terminal symbols.

T - It is a set of terminals.

P - It is the production rules that consist of both terminals and non-terminals.

S - It is the starting symbol.

A grammar is said to be the Context-free grammar if every production is in the form of :

$G \rightarrow (VUT)^*$, where $G \in V$

- And the left-hand side of the G, here in the example, can only be a Variable, it cannot be a terminal.
- But on the right-hand side here it can be a Variable or Terminal or both combination of Variable and Terminal.

The above equation states that every production which contains any combination of the 'V' variable or 'T' terminal is said to be a context-free grammar.

For example, the grammar $A = \{ S, a, b \}$ having productions:

- Here S is the starting symbol.

- $\{a, b\}$ are the terminals generally represented by small characters.
- S is the variable.

$S \rightarrow aS$

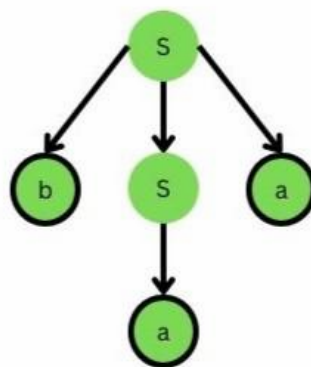
$S \rightarrow bSa$

but

$a \rightarrow bSa$, or

$a \rightarrow ba$ is not a CFG as on the left-hand side there is a variable which does not follow the CFGs rule.

Lets consider the string “**aba**” and try to derive the given grammar from the productions given. we start with symbol S , apply production rule $S \rightarrow bSa$ and then $S \rightarrow aS$ ($S \rightarrow a$) to get the string “**aba**”.



Context-free grammars are frequently used, especially in the areas of formal language theory, compiler development, and natural language processing. It is also used for explaining the syntax of programming languages and other formal languages.

Limitations of Context-Free Grammar:

Apart from all the uses and importance of Context-Free Grammar in the Compiler design and the Computer science field, there are some limitations that are addressed, that is CFGs are less expressive, and neither English nor programming language can be expressed using Context-Free Grammar. Context-Free Grammar can be ambiguous means we can generate multiple parse trees of the same input. For some grammar, Context-Free Grammar can be less efficient because of

the exponential time complexity. And the less precise error reporting as CFGs error reporting system is not that precise that can give more detailed error messages and information.

2. **Learning:** The process of inferring these rules from a given set of examples (positive or negative) and using them to model the data or predict new data.

Learning, in the context of data science, artificial intelligence, and machine learning, refers to the process of inferring rules or patterns from a given set of data (examples) and using those patterns to make predictions, classify new data, or model the underlying structure of the data. Here's a detailed breakdown:

1. Types of Learning:

- **Supervised Learning:**

- In supervised learning, the system is trained on a labeled dataset, meaning that each input comes with the correct output (label). The task of the system is to learn the mapping from inputs to outputs by inferring rules from the data.
- Examples include classification (e.g., determining whether an email is spam or not) and regression (e.g., predicting house prices).
- **Process:** The model receives a dataset of input-output pairs, learns patterns from these examples, and uses them to predict outputs for new, unseen inputs.

- **Unsupervised Learning:**

- In unsupervised learning, the system is provided with input data without explicit labels. The system's task is to infer patterns, groupings, or structures from the data itself.
- Examples include clustering (e.g., customer segmentation) and dimensionality reduction (e.g., principal component analysis).
- **Process:** The model analyzes the structure or distribution of the input data and attempts to discover underlying patterns or relationships.

- **Semi-supervised Learning:**

- This is a hybrid between supervised and unsupervised learning, where the model is trained on a small amount of labeled data and a larger amount of unlabeled data.
- **Process:** The labeled data helps the model understand some patterns, while the unlabeled data allows it to generalize better.
- **Reinforcement Learning:**
 - In reinforcement learning, the system learns by interacting with an environment. It makes decisions (actions) based on the current state of the environment and receives feedback (rewards or punishments) based on the outcome of those decisions.
 - **Process:** The model continuously refines its strategy (policy) to maximize the cumulative reward over time.

2. Steps in the Learning Process:

- **Data Collection:**
 - The process begins by gathering examples (data points) that represent the task to be learned. These examples can be positive (e.g., an image of a cat) or negative (e.g., an image of something that is not a cat).
- **Data Preprocessing:**
 - Raw data often requires cleaning and transformation. This step may involve removing outliers, handling missing data, normalizing features, or converting categorical data into numerical formats.
- **Feature Selection/Engineering:**
 - Features are the individual measurable properties or characteristics of the data. Feature selection involves choosing the most relevant features, while feature engineering involves creating new features from the raw data that might be more informative to the learning algorithm.

- **Model Selection:**

- Once the data is ready, the next step is to choose a learning algorithm (model). Different algorithms are suitable for different types of data and tasks (e.g., decision trees, neural networks, support vector machines).

- **Training:**

- The model is trained on the data. In this phase, the model infers the rules by adjusting its internal parameters to minimize the error (the difference between the predicted outputs and the actual outputs).
- Techniques like gradient descent, back propagation (in neural networks), or optimization algorithms help adjust the model's parameters to fit the data.

- **Evaluation:**

- The performance of the model is assessed by testing it on a separate set of data that it hasn't seen before. Common evaluation metrics include accuracy, precision, recall, F1-score (for classification), and mean squared error (for regression).

- **Tuning:**

- If the model's performance isn't satisfactory, hyper parameters (settings that control the learning process) are tuned to improve results. This may involve adjusting learning rates, the complexity of the model, or even trying different algorithms.

- **Prediction:**

- Once the model is trained and evaluated, it can be deployed to predict new, unseen data.

3. Key Concepts in Learning:

- **Generalization:**

- A good model doesn't just memorize the training data but generalizes well to new, unseen data. Over fitting (where a model is too complex and fits noise in the

training data) can harm generalization, while under fitting (where a model is too simple) can lead to poor performance.

- **Bias-Variance Tradeoff:**

- **Bias** refers to errors introduced by oversimplifying the model (under fitting), while **variance** refers to errors introduced by overcomplicating the model (over fitting). There's often a tradeoff between these two, and the goal is to find a balance that allows the model to generalize well.

- **Regularization:**

- Techniques like L1 or L2 regularization penalize overly complex models to prevent over fitting. Regularization adds a term to the loss function to discourage large or complex weights/parameters in the model.

- **Cross-validation:**

- A technique used to assess the model's performance more reliably. The dataset is split into several parts (folds), and the model is trained and tested on different combinations of these parts, ensuring the model's performance is consistent across different subsets of the data.

4. Inference:

- After learning the patterns or rules from the training data, the model can infer results for new, unseen data. Inference is essentially the process of applying the learned model to new data to make predictions or classify instances.

5. Example:

- Suppose we are developing a spam detection model.
 - **Data Collection:** We collect a dataset of emails, some of which are labeled as spam and some as non-spam.
 - **Preprocessing:** The emails are cleaned by removing unnecessary information (like HTML tags), and important features like word frequencies are extracted.
 - **Model Selection:** A decision tree or a logistic regression model might be chosen for this task.
 - **Training:** The model is trained on a portion of the labeled emails to learn which features (e.g., certain keywords) distinguish spam from non-spam.

- **Evaluation:** The model's accuracy is assessed by testing it on a separate set of emails.
- **Prediction:** Once satisfied with the model's performance, it can be deployed to classify new emails as spam or non-spam.

3. Inference: Involves generalizing from specific examples to broader rules that can be applied to unseen data.

Types of Grammatical Inference:

1. Positive Example Learning:

- The learner is given a set of strings that belong to the language (positive examples) and must infer a grammar that generates these strings.

2. Negative Example Learning:

- The learner is also provided with strings that do not belong to the language (negative examples), helping refine the inferred grammar by excluding these cases.

3. Active Learning:

- The learner interacts with an oracle (such as a human expert or a more complex model) to query whether certain strings belong to the language, actively refining the grammar.

Applications in Pattern Recognition: Grammatical inference is particularly useful in areas where the patterns exhibit a structured or hierarchical nature, such as:

- **Natural Language Processing (NLP):** Learning the syntax of a language from a corpus of text, which can then be used to parse sentences, generate text, or recognize speech.
- **Bioinformatics:** Inferring patterns in DNA sequences to recognize genes or predict protein structures.

- **Document Analysis:** Learning the structure of documents (e.g., forms, reports) to automatically extract information or classify documents.
- **Robot Navigation:** Learning patterns in the environment to infer rules for robot movement and decision-making.

Example:

Imagine you are given a set of strings that represent valid sequences of commands for a robot, such as:

- `"move_forward; turn_left; move_forward"`
- `"move_forward; turn_right; move_backward"`
- `"turn_left; move_forward; turn_right; move_backward"`

The goal is to learn the underlying grammar that can generate these sequences and recognize whether new sequences are valid.

Steps:

1. **Collect Examples:** Gather a set of valid command sequences (positive examples) and optionally some invalid ones (negative examples).
2. **Infer Grammar:** Use a grammatical inference algorithm to derive a set of rules (a grammar) that explains the observed sequences. For example, a context-free grammar might include rules like:

- `S → move_command; S | turn_command; S | move_command`
- `move_command → move_forward | move_backward`
- `turn_command → turn_left | turn_right`

Learning Grammars:

Learning grammars in pattern recognition is a method used to model and recognize complex structures within data. It involves creating formal grammars (a set of rules) to generate or recognize patterns, particularly when the data has underlying structures, like sequences or hierarchical arrangements. Here's how it works and its importance in pattern recognition:

i. **Grammars:**

- a. Grammars are sets of rules or productions used to describe how strings (or patterns) are generated.
- b. In pattern recognition, grammars can describe how certain patterns are formed, allowing systems to both generate valid patterns and recognize them in data.

ii. **Types of Grammars:**

- a. **Regular Grammars:** Used for simpler pattern structures, like sequences.
- b. **Context-Free Grammars (CFGs):** Useful for hierarchical structures, such as those in natural language or certain image recognition tasks.
- c. **Context-Sensitive Grammars:** Can handle even more complex dependencies between parts of patterns.
- d. **Stochastic Grammars:** These assign probabilities to different production rules, helping with the recognition of patterns that are inherently probabilistic (like speech or handwritten text).

iii. **Learning Grammars:**

- a. **Supervised Learning:** Grammar rules are learned by training a model on a labeled dataset where the patterns and their corresponding classes

are known. This is common in fields like natural language processing (NLP).

- b. **Unsupervised Learning:** The system tries to infer grammatical rules directly from the data without explicit labels. Techniques like clustering and generative models can help in this process.
- c. **Inductive Grammar Learning:** The model induces grammar rules from examples and counterexamples of patterns. It's especially useful for tasks where clear rules can be derived from a few instances.

Problem formulation:

The **problem formulation in pattern recognition** is a crucial step, where the task of recognizing patterns from data is mathematically and conceptually framed. It helps define the goals, inputs, outputs, and methods required to solve the pattern recognition task. Proper problem formulation allows for better algorithm design, model selection, and evaluation. Here's a breakdown of how this process is typically structured:

1. Pattern Recognition Problem Definition

The goal in pattern recognition is to classify data (input patterns) into one of several predefined categories (classes). The task is to find a mapping from the input space to a set of labels or decisions. Formally, the problem can be defined as:

- Given an input space X (feature space) and a finite set of possible classes Y (label space), find a function $f: X \rightarrow Y$ that assigns the correct label to each input instance.

2. Key Components of Problem Formulation

a. Input Representation (Feature Space)

The first step is to represent each data sample (pattern) using features. These features describe the data in a way that can be processed by the recognition system. Features are usually vectors that contain numerical, categorical, or binary values derived from the raw data.

- **Examples of features:**

- For image recognition: color histograms, edges, textures.
- For speech recognition: frequency
- For text classification: word frequencies, word embeddings.

b. Class Labels (Output Space)

The output of the pattern recognition system is the class label associated with the input pattern. Classes can be discrete or continuous, depending on the type of recognition problem.

- **Examples of classes:**

- Handwritten digit recognition: the digits 0 to 9.
- Object recognition in images: car, person, tree, etc.
- Disease classification: healthy, disease A, disease B.

c. Classification Function

The problem requires finding or learning a decision function that can map the input features to one of the class labels. This function is generally unknown and is learned from a dataset of labeled examples.

- **Supervised Learning:** Most pattern recognition problems are formulated as supervised learning tasks, where the classifier is trained using a set of labeled examples (training data).
- **Unsupervised Learning:** In some cases, the classes may not be predefined, and the task is to group similar patterns together (clustering).
- **Semi-supervised or Reinforcement Learning:** These approaches are used when labeled data is scarce, or decisions evolve based on feedback.

d. Error Function / Objective Function

In the pattern recognition problem, a classifier's performance is evaluated based on how well it assigns the correct class labels to new, unseen input patterns. The error or objective function helps to evaluate this performance.

- **Examples of error functions:**
 - Classification error: The proportion of misclassified instances.
 - Log-likelihood: Used in probabilistic models like Bayesian classifiers.
 - Cross-entropy loss: Common in neural networks.

3. Approaches to Solving the Problem

a. Statistical Methods

Statistical approaches assume that the patterns are generated by a probability distribution. The task is to estimate this distribution or directly model the decision boundary.

- **Bayesian Classifiers:** Based on calculating the posterior probabilities of classes given input features.

- **Maximum Likelihood Estimation (MLE):** A method for estimating parameters that maximize the likelihood of the observed data.
- **Discriminant Analysis:** Linear or quadratic discriminant functions can be used to separate classes in feature space.

b. Geometrical Methods

In geometrical approaches, the goal is to find boundaries in the feature space that separate different classes.

- **Linear classifiers:** Linear Support Vector Machines (SVM), Perceptron.
- **Non-linear classifiers:** Kernel SVM, Neural Networks.

c. Structural Methods

Structural pattern recognition deals with recognizing complex structures or relations between components of the input data, like sequences or graphs. Here, the problem is to identify these structures and classify them accordingly.

- **Grammatical approaches:** Parsing complex patterns using grammars.
- **Graph-based approaches:** Graph matching for recognition in cases like handwriting or protein structure recognition.

d. Neural Networks and Deep Learning

Deep learning models, especially Convolutional Neural Networks (CNNs) and Recurrent Neural Networks (RNNs), have become dominant in many pattern recognition tasks due to their ability to learn hierarchical feature representations directly from raw data.

- **CNNs** for image classification.
- **RNNs/LSTMs** for sequence-based data (speech, time-series).

4. Pattern Recognition System Design Steps

a. Data Collection

The first step is gathering a representative dataset for the problem. This dataset should contain sufficient examples of each pattern or class.

b. Feature Extraction

Transform the raw data into a set of features that can be used for classification. Feature extraction should capture the essential properties of the data while reducing noise and irrelevant details.

c. Classifier Design

Choose or design an appropriate classifier based on the nature of the problem and the available data. This could be a statistical model, a neural network, or a rule-based system.

d. Training

Train the classifier on the labeled dataset. This involves tuning the model's parameters to minimize error on the training set while avoiding overfitting.

e. Evaluation

The model's performance is evaluated on a test set that contains unseen examples. Common metrics include accuracy, precision, recall, F1-score, and confusion matrix analysis.

5. Challenges in Problem Formulation

a. Dimensionality of Feature Space

In some cases, the feature space can be very high-dimensional, leading to issues like the "curse of dimensionality," where the model may overfit due to the large number of features compared to the available data.

b. Imbalanced Data

Some pattern recognition tasks suffer from class imbalance, where some classes have far fewer examples than others. Special care needs to be taken to avoid biased classifiers.

c. Noise and Variability in Data

Real-world data often contains noise, missing values, or variability in how patterns appear. Handling such noise is an important aspect of problem formulation.

d. Over fitting and Generalization

Balancing model complexity is crucial. A model that is too complex may overfit the training data and fail to generalize well to unseen data.

Grammatical Inference (GI) Approaches:

GI approaches used in pattern recognition:

1. **State-based approaches:** These techniques focus on inferring automata (like finite-state machines) from observed sequences of symbols. For instance, given a set of input sequences, a state-based GI approach can generate a finite automaton that models the sequences. These methods are commonly used in speech recognition and bioinformatics.
2. **Rule-based approaches:** This method tries to infer a set of production rules that define a grammar from the given data. The inferred grammar can then be used to generate new patterns or recognize existing ones. These approaches are useful in areas like natural language processing.
3. **Statistical approaches:** In this method, probabilistic models (like hidden Markov models or stochastic context-free grammars) are inferred from data. These approaches are particularly effective when the data is noisy or incomplete and are often applied in fields such as speech and handwriting recognition.
4. **Hybrid approaches:** These methods combine different techniques, such as state-based models and statistical learning, to improve performance, especially when dealing with complex pattern recognition tasks.

Procedures to Generate Constrained Grammars.

In **pattern recognition**, generating **constrained grammars** can be crucial for modeling structured patterns such as sequences in language processing, biometric data, or biological

patterns. The aim is to develop grammars that generate only valid patterns under specific constraints.

1. Define the Pattern Class and Constraints-

A. Determine the desired patterns: Choose the kind of patterns (such as shapes, sound signals, or text sequences) that you wish to be able to identify first.

B. Establish limitations: Determine the rules (such as particular lengths, repetition structures, or boundary requirements) that limit the set of acceptable patterns. This could be a restriction on shape features in image recognition or phoneme sequences in speech recognition.

C. Include a noise tolerance clause: Noise is a common component of real-world data in pattern recognition. Some pattern variability should be supported by the grammar.

2. Choose an Appropriate Grammar Formalism

Depending on the complexity of the patterns and constraints, you can choose between various grammar types:

- **Regular Grammars:** Useful for simple and repetitive patterns. For example, recognizing patterns that alternate between two symbols (like ab^*).
- **Context-Free Grammars (CFG):** Used for more structured, hierarchical patterns. For instance, recognizing valid nested parentheses or tree-like structures in syntax parsing.
- **Context-Sensitive Grammars (CSG):** Apply if the recognition depends on the context of surrounding elements (e.g., protein sequences in bioinformatics).
- **Stochastic/Probabilistic Grammars:** In real-world pattern recognition tasks, probabilistic context-free grammars (PCFG) allow capturing uncertainties (e.g., probabilistic models in speech recognition).

3. Identify Terminals and Non-Terminals

- **Terminals:** These represent the raw data elements or observed symbols (e.g., phonemes, pixels, or nucleotide bases). They are the basic units of the pattern.
- **Non-Terminals:** Non-terminals capture abstract features or larger components of the pattern (e.g., syllables in speech, geometric shapes in images).

Example:

- **Terminals:** {a, b}
- **Non-terminals:** <S> for sequences, <A> for alternating structures.

4. Define Production Rules

- **Create production rules** that reflect the structure of the pattern. Rules define how non-terminals expand into terminals or other non-terminals.
- Rules can enforce constraints like the order of symbols, repetitions, or structural properties (e.g., nested patterns, symmetric features).

Example:

- For a pattern where an object is composed of a specific number of parts:
 $\langle \text{object} \rangle ::= \langle \text{part} \rangle \langle \text{object} \rangle \mid \varepsilon$
- For recognizing a pattern of alternating signals:
 $\langle A \rangle ::= a \langle B \rangle \mid b \langle A \rangle$
 $\langle B \rangle ::= b \langle A \rangle \mid \varepsilon$

5. Incorporate Constraints into the Grammar

- **Length constraints:** Restrict the grammar to generate sequences or patterns of certain lengths. For instance, in a grammar modeling biological sequences, you might constrain the grammar to ensure a specific number of codons.
- **Structural constraints:** Apply conditions like "every opening bracket must have a closing bracket," or "each phoneme must follow specific others based on language rules."
- **Symbol frequency constraints:** Ensure that certain symbols or patterns occur with a specific frequency or order, such as the repetition of certain characters in a text pattern.

Applications of Constrained Grammars in Pattern Recognition

- **Natural Language Processing (NLP):** Parsing and generating syntactically valid sentences with specific grammar rules.
- **Speech Recognition:** Modeling phoneme patterns using stochastic context-free grammars (SCFGs) to allow for natural speech variability.
- **Image Recognition:** Describing geometric shapes or visual patterns using grammar-based techniques.
- **Bioinformatics:** Recognizing valid DNA or protein sequences using regular grammars to model nucleotide triplets or amino acids.
