

# **Formal Language Simulator – User Manual**

## **Overview**

The Formal Language Simulator is a console-based educational tool designed to demonstrate how different classes of formal languages are recognized using different computational models. The program integrates finite automata, approximate pattern matching, and pushdown automata into a single system.

It is intended for students studying formal languages, automata theory, compilers, and bioinformatics. No prior in-depth knowledge of automata theory is required to use the program.

The simulator supports:

- Regular language recognition using NFA and DFA
- NFA to DFA conversion (subset construction)
- Approximate pattern matching for biological sequences (DNA)
- Context-free language recognition using a Pushdown Automaton (PDA)

## **How the Program Works (Simplified)**

The program is divided into four main functional components, each corresponding to a different class of formal language or recognition technique.

### **1. Pattern Input (Regular Language)**

You first enter a pattern consisting of literal characters. Each character is treated as a symbol in a regular language.

**Important Note:** The program supports literal concatenation only. Special regex operators such as \* or | are treated as normal characters.

#### **Example Pattern**

-ab

This pattern represents the regular language containing exactly the string ab.

From this pattern, the program:

- Constructs an NFA (Non-deterministic Finite Automaton)
- Converts the NFA into an equivalent DFA (Deterministic Finite Automaton)

### **2. Exact Match Using NFA and DFA**

After building the automata, the program asks for a test string.

Both the NFA and DFA simulate the input string:

- The NFA explores all possible transitions simultaneously
- The DFA follows a single deterministic transition path

If the simulation ends in a final state, the string is accepted.

### **Sample Input**

Pattern: ab

Test string: ab

Output

NFA ACCEPT

DFA ACCEPT

### **3. Approximate Pattern Matching (DNA Sequence Analysis)**

The simulator supports approximate matching using an edit-distance-based dynamic programming algorithm.

This feature models real-world DNA sequence analysis, where small differences (mutations, insertions, deletions) are common.

The program checks whether the pattern appears as a substring in the given DNA sequence within a user-defined maximum number of errors.

### **Sample Input**

Pattern: ACG

DNA sequence: TACGGA

Maximum errors: 1

Output

Approximate match found

This means the pattern ACG appears in the DNA sequence with at most one difference.

### **4. Transition Visualization (NFA and DFA)**

For transparency and learning purposes, the program prints the transition tables for both automata.

Example NFA Output

0 --a--> 1

1 --b--> 2

Start: 0

Final: 2

Example DFA Output

0 --a--> 1

1 --b--> 2

Start: 0

Final: 2

These outputs help illustrate how strings are processed step by step by each automaton.

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## 5. Context-Free Language Recognition (Pushdown Automaton)

To demonstrate the limitations of regular languages, the program includes a Pushdown Automaton (PDA).

The PDA recognizes the classic context-free language:

$$L = \{ a^n b^n \mid n \geq 1 \}$$

This language cannot be recognized by finite automata alone and requires a stack, which is provided by the PDA.

PDA Behavior

- Push one symbol onto the stack for each a
- Pop one symbol from the stack for each b
- Accept if the stack is empty and the input is fully consumed

### Sample Input

aabb

Output

PDA ACCEPT ( $a^n b^n$ )

Rejected Example

aabbb

Output:

PDA REJECT

Step-by-Step Usage

1. Compile the program

```
g++ -std=c++17 searchsystem.cpp -static-libgcc -static-libstdc++ -O2 -o  
formal_sim.exe
```

2. Run the program
3. Enter a pattern (literal concatenation only)
4. Enter a string for exact matching
5. Enter a DNA sequence and maximum allowed errors
6. Enter a string for PDA testing ( $a^n b^n$  form)
7. Review the results and transition tables

### Sample Complete Run

Enter regex: ab

Enter string for exact match: ab

NFA ACCEPT

DFA ACCEPT

Enter DNA sequence: xabx

Enter max errors: 1

Approximate match found

Enter string for PDA test: aaabbb

PDA ACCEPT ( $a^n b^n$ )

### Tips for Users

- Use uppercase A, C, G, T for DNA input
- Avoid spaces in inputs (the program reads tokens)
- Remember that regex operators like \* are treated as literal characters
- Use the PDA section to understand why some languages are not regular

### Summary

This simulator demonstrates: - Regular language recognition using NFA and DFA - Performance-oriented conversion from NFA to DFA - Approximate matching for biological sequences - Context-free language recognition using Pushdown Automata

It clearly illustrates the boundaries between regular and context-free languages using practical, executable examples.