

An Interdisciplinary Approach
Robert Sedgewick • Kevin Wayne

## 17. Introduction to Theoretical CS

Section 7.2
http://introcs.cs.princeton.edu

Introduction to theoretical computer science

## Fundamental questions

-What can a computer do?

- What can a computer do with limited resources?

General approach

- Don't talk about specific machines or problems.
- Consider minimal abstract machines.
- Consider general classes of problems.


Surprising outcome. Sweeping and relevant statements about all computers.

## Why study theory?

In theory...

- Deeper understanding of computation.
- Foundation of all modern computers.
- Pure science.
- Philosophical implications.

In practice...

- Web search: theory of pattern matching.
- Sequential circuits: theory of finite state automata.
- Compilers: theory of context free grammars.
- Cryptography: theory of computational complexity.
- Data compression: theory of information.
- ...

"In theory there is no difference between theory and practice. In practice there is. "


## Programming

in Java



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## 17. Introduction to Theoreticaal CS

- Regular expressions
- DFAs
- Applications
- Limitations


## Pattern matching

Pattern matching problem. Is a given string a member of a given set of strings?
Example 1 (from genomics)
A nucleic acid is represented by one of the letters $\mathrm{a}, \mathrm{c}, \mathrm{t}$, or g .
A genome is a string of nucleic acids.
A Fragile X Syndrome pattern is a genome having an occurrence of gcg, followed by any number of cgg or agg triplets, followed by ctg.

Note. The number of triplets correlates with Fragile X Syndrome, a common cause of mental retardation.
Q. Does this genome contain a such a pattern?
gcggcgtgtgtgcgagagagtgggtttaaagctggcgcggaggcggctggcgcggaggctg
A. Yes.
gcg start mark


## Pattern matching

Example 2 (from computational biochemistry)
An amino acid is represented by one of the characters CAVLIMCRKHDENQSTYFWP.

A protein is a string of amino acids.

A $\mathrm{C}_{2} \mathrm{H}_{2}$-type zinc finger domain signature is

- C followed by 2, 3, or 4 amino acids, followed by
- C followed by 3 amino acids, followed by
- L, I, V, M, F, Y, W, C, or X followed by 8 amino acids, followed by
- H followed by 3, 4, or 5 amino acids, followed by
- H.
Q. Is this protein in the $\mathrm{C}_{2} \mathrm{H}_{2}$-type zinc finger domain?



## Pattern matching

Example 3 (from commercial computing)

An e-mail address is

- A sequence of letters, followed by
- the character "@", followed by
- the character "." , followed by a sequence of letters, followed by
- [any number of occurences of the previous pattern]
- "edu" or "com" (others omitted for brevity).
Q. Which of the following are e-mail addresses?
A.
rs@cs.princeton.edu $\checkmark$
not an e-mail address $\quad x$
wayne@cs.princeton.edu $\checkmark$
eve@airport $x$
Ooops, need to fix description $\longrightarrow$ rsl23@princeton.edu $\boldsymbol{x}$

Challenge. Develop a precise description of the set of strings that are legal e-mail addresses.

## Regular expressions

A regular expression (RE) is a notation for specifying sets of strings.

## An RE is

- A sequence of letters or "."
- The union of two REs
- The closure of an RE
- May be delimited by ().

| operation | example RE | matches <br> (IN the set) | does not match <br> (NOT in the set) |
| :---: | :---: | :---: | :---: |
| concatenation | aabaab | aabaab | every other string |

## More examples of regular expressions

The notation is surprisingly expressive.
\(\left.\begin{array}{c|cc|c}regular expression \& matches \& does not match <br>
contains the trigraph spb \& raspberry <br>

crispbread\end{array}\right]\)| subspace |
| :---: |
| subspecies |

## Generalized regular expressions

Additional operations futher extend the utility of REs.

| operation | example RE | matches | does not match |
| :---: | :---: | :---: | :---: |
| one or more | $\mathrm{a}(\mathrm{bc})+\mathrm{de}$ | abcde <br> abcbcde | ade <br> bcde |
| character class | $[\mathrm{A}-\mathrm{Za}-\mathrm{z}][\mathrm{a}-\mathrm{z}] *$ | lowercase <br> Capitalized | came1Case <br> $4 \mathrm{i} 17 \mathrm{lega7}$ |
| exactly k | $[0-9]\{5\}-[0-9]\{4\}$ | $08540-1321$ <br> $19072-5541$ | 111111111 <br> $166-54-1111$ |
| negation | $[\wedge$ aeiou]\{6\} | rhythm | decade |
| white space | $\backslash s$ | any whitespace char <br> (space, tab, newline...) | every other character |

Note. These operations are all shorthand.
They are very useful but not essential.

RE: $(a|b| c|d| e)(a|b| c|d| e) *$
shorthand: (a-e)+

## Example of describing a pattern with a generalized RE

A $\mathrm{C}_{2} \mathrm{H}_{2}$-type zinc finger domain signature is

- C followed by 2, 3, or 4 amino acids, followed by
- C followed by 3 amino acids, followed by
- L, I, V, M, F, Y, W, C, or X followed by 8 amino acids, followed by
- H followed by 3, 4, or 5 amino acids, followed by
- H.
Q. Give a generalized RE for all such signatures.
A. C. $\{2,4\}$ C...[LIVMFYWCX]. $\{8\} \mathrm{H} .\{3,5\} \mathrm{H}$
"Wildcard" matches any of the letters
CAVLIMCRKHDENQSTYFWP



## Example of a real-world RE application: PROSITE

## 1ras <br> Database of protein domains, families and functional sites

PROSITE consists of documentation entries describing protein domains, families and functional sites as well as associated patterns and profiles to identify them [More... / References / Commercial users]
PROSITE is complemented by ProRule, a collection of rules based on profiles and patterns, which increases the discriminatory power of profiles and patterns by providing additional information about functionally and/or structurally critical amino acids [More...].
Forthcoming changes: information can be found here.

## Release 20.97, of 08-Nov-2013 (1673 documentation entries, 1308 patterns, 1056 profiles and 1062 ProRule)

Browse

- by documentation entry
by ProRule description
by taxonomic scope
- by number of positive hits


## Another example of describing a pattern with a generalized RE

## An e-mail address is

- A sequence of letters, followed by
- the character "@", followed by
- the character "." , followed by a sequence of letters, followed by
- [any number of occurences of the previous pattern]
- "edu" or "com" (others omitted for brevity).
Q. Give a generalized RE for e-mail addresses.
A. $[a-z]+@([a-z]+\backslash)+.(e d u \mid c o m)$

Exercise. Extend to handle rs123@princeton.edu, more suffixes such as .org, and any other extensions you can think of.

Next. Determining whether a given string matches a given RE.

## TEQ 1 on REs

Q. Which of the following strings match the RE $a * b b(a b \mid b a) * ?$
is in the set
it describes

1. abb
2. aaba
3. abba
4. bbbaab
5. cbb
6. bbababbab

## TEQ 1 on REs

Q. Which of the following strings match the RE $a * b b(a b \mid b a) * ?$
is in the set
it describes


## TEQ 2 on REs

Q. Give an RE for genes

- Characters are a, c, t or g.
- Starts with atg (a start codon).
- Length is a multiple of 3.
- Ends with tag, taa, or ttg (a stop codon).



## TEQ 2 on REs

Q. Give an RE for genes

- Characters are a, c, t or g.
- Starts with atg (a start codon).
- Length is a multiple of 3.
- Ends with tag, taa, or ttg (a stop codon).

A. $\operatorname{atg}((a|c| t \mid g)(a|c| t \mid g)(a|c| t \mid g)) *(t a g|t a a| t t g)$


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## Deterministic finite state automata (DFA)

A DFA is an abstract machine that solves a pattern matching problem.

- A string is specified on an input tape (no limit on its length).
- The DFA reads each character on input tape once, moving left to right.
- The DFA lights "YES" if it recognizes the string, "NO" otherwise.

Each DFA defines a set of strings (all the strings that it recognizes).


## Deterministic finite state automata details and example

A DFA is an abstract machine with a finite number states, each labeled Y or N , and transitions between states, each labelled with a symbol. One state is the start state.

- Begin in the start state (denoted by an arrow from nowhere).
- Read an input symbol and move to the indicated state.
- Repeat until the last input symbol has been read.
- Turn on the "YES" or "NO" light according to the label on the current state.



## Deterministic finite state automata details and example

A DFA is an abstract machine with a finite number states, each labeled Y or N and transitions between states, each labelled with a symbol. One state is the start state.

- Begin in the start state.
- Read an input symbol and move to the indicated state.
- Repeat until the last input symbol has been read.
- Turn on the "YES" or "NO" light according to the label on the current state.



## Simulating the operation of a DFA

```
public class DFA
{
    private int state; symbol table to map
    private int start;
    privale int start;
    private String[] action;
    private ST<Character, Integer>[] next;
    public DFA(In in)
    { /* Fill in data structures */ }
    public String simulate(String input)
    {
        state = start;
        for (int i = 0; i < input.length(); i++)
                state = next[state].get(input.charAt(i));
            return action[state];
    }
    public static void main(String[] args)
    {
        DFA dfa = new DFA(new In(args[0]));
        while (!StdIn.isEmpty())
        {
            input = StdIn.readString();
            StdOut.println(dfa.simulate(input));
        }
    }
}
```


\% more b3.txt
\# states $\longrightarrow 3$
alphabet $\longrightarrow \mathrm{ab}$
start state $\longrightarrow 0$
Yes 01
No 12
No 20
\% java DFA b3.txt bababa
Yes
bb
No
abbabbababbbabaaa
Yes
abbabbababbba
No

## TEQ 1 on DFAs

Q. Which of the following strings does this DFA accept?


1. Bitstrings that end in 1
2. Bitstrings with an equal number of occurrences of 01 and 10
3. Bitstrings with more 1 s than 0 s
4. Bitstrings with an equal number of occurrences of 0 and 1
5. Bitstrings with at least one 1

## TEQ 1 on DFAs

Q. Which of the following strings does this DFA accept?

A.

1. Bitstrings that end in 1

X 0110
2. Bitstrings with an equal number of occurrences of 01 and 10

X 011010
3. Bitstrings with more 1 s than 0 s

X 0001
4. Bitstrings with an equal number of occurrences of 0 and 1

X 0001
5. Bitstrings with at least one 1
$\checkmark$

## TEQ 2 on DFAs

Q. Which of the following strings does this DFA accept?


1. Bitstrings with at least one 1
2. Bitstrings with an equal number of occurrences of 01 and 10
3. Bitstrings with more 1 s than 0 s
4. Bitstrings with an equal number of occurrences of 0 and 1
5. Bitstrings that end in 1

## TEQ 2 on DFAs

Q. Which of the following strings does this DFA accept?

A.

1. Bitstrings with at least one 1

X 0100
2. Bitstrings with an equal number of occurrences of 01 and 10
$x \quad 0110$
3. Bitstrings with more 1 s than 0 s

X 010000001
4. Bitstrings with an equal number of occurrences of 0 and 1

X 01111111
5. Bitstrings that end in 1
$\checkmark$

## Kleene's theorem

$S \equiv$ the set of $a b$ strings where the number of occurrences of $b$ is a multiple of 3
Two ways to define a set of strings

- Regular expressions (REs).
- Deterministic finite automata (DFAs).

Remarkable fact. DFAs and REs are equivalent.

## Equivalence theorem (Kleene)

Given any RE, there exists a DFA that accepts the same set of strings. Given any DFA, there exists an RE that matches the same set of strings.

Consequence: A way to solve the RE pattern matching problem

- Build the DFA corresponding to the given RE.
- Simulate the operation of the DFA.


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## GREP: a solution to the RE pattern matching problem

An algorithm for the RE pattern matching problem?

- Build the DFA corresponding to the given RE.
- Simulate the operation of the DFA.

Practical difficulty: The DFA might have exponentially many states.

A more efficient algorithm: use Nondeterministic Finite Automata (NFA)

- Build the NFA corresponding to the given RE.
- Simulate the operation of the NFA.
"GREP" (Generalized Regular Expression Pattern matcher).
- Developed by Ken Thompson, who designed and implemented Unix.
- Indispensible programming tool for decades.
- Found in most development environments, including Java.


Interested in
details? Take a
course in
algorithms.

## REs in Java

Java's String class implements GREP.

```
public class String
    boolean matches(String re) does this string match the given RE?
```

    String re = "C. \(\{2,4\} \mathrm{C} .\). [LIVMFYWC]. \(\{8\} \mathrm{H} .\{3,5\} \mathrm{H} "\);
    String zincFinger = "CAASCGGPYACGGAAGYHAGAH";
boolean test $=$ zincFinger.matches(re);
true!

CAASCGGPYACGGWAGYHAGWH


## Java RE client example: Validation

```
public class Validate
{
    public static void main(String[] args)
    {
        String re = args[0];
        while (!StdIn.isEmpty())
        {
            String input = StdIn.readString();
            StdOut.println(input.matches(re));
        }
    }
}
```


## Applications

- Scientific research.
- Compilers and interpreters.
- Internet commerce.
- ...

Does a given string match a given RE?

- Take RE from command line.
- Take strings from Stdln.



## Beyond matching

Java's String class contains other useful RE-related methods.

- RE search and replace
- RE delimited parsing
public class String

| $\ldots$ |  |
| :--- | :--- |
| String | replaceA11 (String re, String to) |
| String[] | split(String re) |
|  | $\ldots$ |
|  | split the string around matches of the given $R E$ |

Tricky notation (typical in string processing): \signals "special character" so "<br>" means "\" Examples using the RE "<br>s+" (matches one or more whitespace characters). and "<br>s" means "\s"

Replace each sequence of at least one whitespace character with a single space.

```
String s = StdIn.readA11();
    s = s.replaceA11("\\s+", " ");
```

Create an array of the words in StdIn (basis for StdIn. readA11Strings() method)

```
String s = StdIn.readA11();
    String[] words = s.split("\\s+'');
```


## Way beyond matching

Java's Pattern and Matcher classes give fine control over the GREP implementation.

[A sophisticated interface designed for pros, but very useful for everyone.]

## Java pattern matcher client example: Harvester

    }
    }

Harvest information from input stream

- Take RE from command line.
- Take input from file or web page.
- Print all substrings matching RE.

```
```

import java.util.regex.Pattern;

```
```

import java.util.regex.Pattern;
import java.util.regex.Matcher;
import java.util.regex.Matcher;
public class Harvester
public class Harvester
{
{
public static void main(String[] args)
public static void main(String[] args)
{
{
String re = args[0];
String re = args[0];
In in = new In(args[1]);
In in = new In(args[1]);
String input = in.readAl1();
String input = in.readAl1();
Pattern pattern = Pattern.compile(re);
Pattern pattern = Pattern.compile(re);
Matcher matcher = pattern.matcher(input);
Matcher matcher = pattern.matcher(input);
while (matcher.find())
while (matcher.find())
StdOut.println(matcher.group());

```
            StdOut.println(matcher.group());
```

```
% java Harvester "gcg(cgg|agg)*ctg" chromosomeX.txt
```

% java Harvester "gcg(cgg|agg)*ctg" chromosomeX.txt

```
% java Harvester "gcg(cgg|agg)*ctg" chromosomeX.txt
gcgcggcggcggcggcggctg
gcgcggcggcggcggcggctg
gcgcggcggcggcggcggctg
gcgctg
gcgctg
gcgctg
gcgctg
gcgctg
gcgctg
gcgcggcggcggaggcggaggcggctg
gcgcggcggcggaggcggaggcggctg
gcgcggcggcggaggcggaggcggctg
% java Harvester "[a-z]+@([a-z]+\.)+(edu|com)" http://www.cs.princeton.edu/people/faculty
% java Harvester "[a-z]+@([a-z]+\.)+(edu|com)" http://www.cs.princeton.edu/people/faculty
% java Harvester "[a-z]+@([a-z]+\.)+(edu|com)" http://www.cs.princeton.edu/people/faculty
...
...
...
rs@cs.princeton.edu
rs@cs.princeton.edu
rs@cs.princeton.edu
wayne@cs.princeton.edu
```

wayne@cs.princeton.edu

```
wayne@cs.princeton.edu
```

```
harvest email addresses from web for spam campaign.
```

```
harvest email addresses from web for spam campaign.
```

```
harvest email addresses from web for spam campaign.
```


## Java pattern matcher real-world example: Parsing a data file

A typical situation

- An institution publishes data on the web to be shared by all.
- The data is published in human-readable form.
- You want to strip out everything but the raw data in order to process it.

Example: National Center for Biotechnology Information genome data


## Java pattern matcher real-world example: Parsing a data file

Key challenge: Develop an appropriate RE.


Extract data after spaces followed by a line number.

Ignore everything else

```
LOCUS AC146846 128142 bp DNA linear HTG 13-NOV-2003
DEFINITION Ornithorhynchus anatinus clone CLM1-393H9,
ACCESSION AC146846
VERSION AC146846.2 GI:38304214
KEYWORDS HTG; HTGS_PHASE2; HTGS_DRAFT.
SOURCE Ornithorhynchus anatinus (platypus)
ORIGIN first (only) group
lst }\longrightarrow1\mathrm{ tgtatttcat ttgaccgtgc tgttttttcc cggtttttca gtacggtgtt agggagccac / in 2nd match
    61 gtgattctgt ttgttttatg ctgccgaata gctgctcgat gaatctctgc atagacagct // a comment
    121 gccgcaggga gaaatgacca gtttgtgatg acaaaatgta ggaaagctgt ttcttcataa
1 2 8 1 0 1 ~ g g a a a t g c g a ~ c c c c c a c g c t ~ a a t g t a c a g c ~ t t c t t t a g a t ~ t g ~
//
```

match

## Java pattern matcher real-world example: Parsing a data file

```
```

import java.util.regex.Pattern;

```
```

import java.util.regex.Pattern;
import java.util.regex.Matcher;
import java.util.regex.Matcher;
public class ParseNCBI
public class ParseNCBI
{
{
public static void main(String[] args)
public static void main(String[] args)
{
{
String re = "[ ]*[0-9]+([actg ]*).*";
String re = "[ ]*[0-9]+([actg ]*).*";
Pattern pattern = Pattern.compile(re);
Pattern pattern = Pattern.compile(re);
In in = new In(args[0]);
In in = new In(args[0]);
while (in.hasNext Line())
while (in.hasNext Line())
{
{
String line = in.readLine();
String line = in.readLine();
Matcher matcher = pattern.matcher(line);
Matcher matcher = pattern.matcher(line);
if (matcher.find())
if (matcher.find())
StdOut.print(matcher.group(1).replaceAl1(" ", ""));
StdOut.print(matcher.group(1).replaceAl1(" ", ""));
}
}
StdOut.println();
StdOut.println();
}
}
}

```
```

}

```
```

\% java ParseNCBI platypus.txt tgtatttcatttgaccgtgctgttttttcccgg tttttcagtacggtgttagggagccacgtgatt ctgtttgttttatgctgccgaatagctgctcga tgaatctctgcatagacagctgccgcagggaga aatgaccagtttgtgatgacaaaatgtaggaaa gctgtttcttcataa...

## Applications of REs

Pattern matching and beyond.

- Compile a Java program.
- Scan for virus signatures.
- Crawl and index the Web.
- Process natural language.
- Access information in digital libraries.
- Search-and-replace in a word processors.
- Process NCBI and other scientific data files.
- Filter text (spam, NetNanny, ads, Carnivore, malware).
- Validate data-entry fields (dates, email, URL, credit card).
- Search for markers in human genome using PROSITE patterns.
- Automatically create Java documentation from Javadoc comments.

GREP and related facilities are built in to Java, Unix shell, PERL, Python ...

OH NO! THE KILLER MUST HAVE POLIOWED HER ON VACATION!


BUT TO FIND THEM WE'D HAVE TO SEARCH THROUGH 200 MB OF EMAILS LOOKING FOR SOMETHING FORMATIED LIKE AN ADDRESS!


http://xkcd.com/208/

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## Summary

## Programmers

- Regular expressions are a powerful pattern matching tool.
- Equivalent DFA/NFA paradigm facilitates implementation.
- Combination greatly facilitates real-world string data processing.



## Theoreticians

- REs provide compact descriptions of sets of strings.
- DFAs are abstract machines with equivalent descriptive power.
- Are there languages and machines with more descriptive power?

You

- CS core principles provide useful tools that you can exploit now.
- REs and DFAs provide an introduction to theoretical CS.


## Basic questions

Q. Are there sets of strings that cannot be described by any RE?
A. Yes.

- Bitstrings with equal number of 0s and 1 s .
- Strings that represent legal REs.
- Decimal strings that represent prime numbers.
- DNA strings that are Watson-Crick complemented palindromes.
- ...
Q. Are there sets of strings that cannot be described by any DFA?
A. Yes.
- Bit strings with equal number of 0 s and 1 s .
- Strings that represent legal REs.

- Decimal strings that represent prime numbers.
- DNA strings that are Watson-Crick complemented palindromes.
- ...


## A limit on the power of REs and DFAs

Proposition. There exists a set of strings that cannot be described by any RE or DFA.

Proof sketch. No DFA can recignize the set of bitstrings with equal number of 0 s and 1 s .

- Assume that you have such a DFA, with $N$ states.
- It recognizes the string with $N+10$ s followed by $N+11 \mathrm{~s}$.
- Some state is revisited when recognizing that string.
- Delete the substring between visits.
- DFA recognizes that string, too.
- It does not have equal number of 0 s and 1 s .
- Proof by contradiction: the assumption that such a DFA exists must be false.

Ex. $N=10$

```
0
0
    0
    0 5
```


## Another basic question

Q. Are there abstract machines that are more powerful than DFAs?
A. Yes. A 1 -stack DFA can recognize

- Bitstrings with equal number of 0s and 1 s .
- Strings that represent legal REs.

Proof. [details omitted]


## Yet another basic question

Q. Are there abstract machines that are more powerful than a 1 -stack DFA?
A. Yes. A 2 -stack DFA can recognize

- Decimal strings that represent prime numbers.
- Strings that represent legal Java programs.
- ...
[stay tuned for next lecture]



## One last basic question

Q. Are there machines that are more powerful than a 2-stack DFA?
A. No! Not even a roomful of supercomputers (!!!)
[stay tuned for next lecture]


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