



Data Structures and Algorithms in Rust



Problem solving with algorithms and data structures using Rust

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· Rust China Community ·



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structures using Rust*

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Preface

The emergence of the transistor sparked a revolution in integrated circuits and chips, leading to the development of the central processing unit, large-capacity storage, and convenient communication facilities. Unix^[1] was born out of the failure of Multics^[2], which subsequently gave rise to the Linux kernel^[3] and its various distributions^[4]. By combining open source with network technology, the rapid development of IT was made possible. Technological progress provides a platform and tools for practical ideas, while social progress creates new demands and passions, further promoting technological progress. Although the upper layer of the computer world has given rise to the Internet, cloud computing, blockchain, AI, and the Internet of Things, the basic principles underlying the bottom layer remain unchanged. The fundament of computing is the combination of hardware and its abstract data types and algorithms. Whether it's a regular computer, a supercomputer, or a quantum computer^[5], their functions are built on some abstraction of data structures and algorithms.

This book focuses on the design, implementation, and use of abstract data types as they play an important role in computer science. By learning to design abstract data types, one can better implement programming and deepen their understanding of them. The algorithm implementations in this book are not the most optimal or general-purpose engineering implementations, as these tend to be verbose and fail to focus on key principles, which can be harmful to learning. Instead, the code in this book takes different simplification measures for different cases, some using generics and others using specific types. These implementation measures are intended to simplify the code and ensure that each segment of code can be compiled and executed separately to obtain a result.

Prerequisite Knowledge

Understanding abstract data types does not require specific forms, but implementing code requires consideration of specific forms, which in turn requires a certain level of proficiency in Rust. Although readers' familiarity with Rust and coding style may vary, the basic requirements remain the same. To read this book, readers should ideally have the following abilities and interests:

- Ability to implement complete programs using Rust, including the use of Cargo, rustc, test, etc.
- Ability to use basic data types and structures, including structs, enums, loops, matches, etc.
- Ability to use Rust generics, lifetimes, ownership system, pointers, unsafe code, macros, etc.
- Ability to use built-in libraries (crates) and external libraries, and to set up Cargo.toml.

If readers lack these abilities, they can refer to the section on Rust learning materials in Chapter 1 and find some recommended books and resources to learn Rust first, before returning to this book. The `code` for this book is available on Github, organized according to chapter and name. Readers are welcome to download and use it, and to point out any errors they find.

This book and all code were written in the Ubuntu 20.04 environment, with Rust version 1.58. The code environment has two types: code with and without line numbers. The former is used to show code, while the latter displays results or other content. All code except for simple or explanatory code, which does not give results, provides outputs. Shorter outputs are commented within the current code box, while more complex outputs are placed in a code box without line numbers.

What This Book Covers

To provide readers with a clear understanding of the structure of the book and to help advanced users select chapters, we have summarized the contents of the entire book below. The book is divided into ten chapters. Chapter 1 organizes Rust learning materials and reviews basic knowledge. Chapters 2 and 3 introduce the concepts of computer science and algorithm analysis, forming the foundation of the book. Chapters 4 to 7 cover the design and implementation of simple data structures and algorithms, while Chapters 8 and 9 cover more complex tree and graph data structures, which are widely used. These two chapters build upon the higher-level topics covered in the previous chapters. The final chapter contains several practical projects that use the data structures and algorithms to solve real problems.

Readers are encouraged to start with the first three chapters and then progress through the middle chapters and practical project in order. However, readers can also choose to learn certain chapters first and then learn other content.

Chapter 1: Rust Basics. This chapter covers the basics of Rust, including installation of the Rust toolchain, organization of Rust learning resources, a review of Rust fundamentals, and a project. Readers who are already familiar with these topics can choose to skip this chapter. The final project summarizes the learned basic knowledge and demonstrates how Rust modules and code are organized.

Chapter 2: Computer Science. This chapter introduces the definition and concepts of computer science, which can help readers analyze real problems. This includes establishing abstract data type models for data structures, designing and implementing algorithms, and verifying algorithm performance. Abstract data types describe the logic of data operations and hide implementation details, which better abstracts the essence of the problem.

Chapter 3: Algorithm Analysis. Algorithm analysis is a method for understanding program execution time and space performance, and can help determine algorithm performance efficiency. The standard method of algorithm analysis is the Big-O notation analysis.

Chapter 4: Basic Data Structures. This chapter covers basic data structures in Rust. These data structures include arrays, slices, Vec, and derivative stacks, queues, and more which are based on the linear memory model. The chapter focuses on using Vec to implement these basic data structures.

Chapter 5: Recursion. Recursion is an algorithmic technique that is another form of iteration. It must satisfy the three laws of recursion, and tail recursion is an optimization of recursion. Dynamic programming is an efficient algorithmic technique, which can be implemented using recursion or iteration.

Chapter 6: Search. This chapter covers search algorithms that are used to find elements in a data set or to determine if an element exists. Sequential and non-sequential search algorithms are explained based on whether the data set is ordered or not.

Chapter 7: Sorting. In the previous chapter, the sequential search algorithm required ordered data, but data is generally unordered. Thus, sorting algorithms are needed. This chapter covers ten common sorting algorithms, including bubble sort, quick sort, insertion sort, shell sort, merge sort, selection sort, heap sort, bucket sort, counting sort, and radix sort. Tim sort algorithm, a combination of some common sorting algorithms which is very efficient, has become the default sorting algorithm in many programming languages.

Chapter 8: Trees. This chapter explains how non-linear data structures can be constructed on linear systems. Trees are non-linear data structures that point to subtrees through pointers or references. Binary trees, binary heaps, binary search trees and balanced binary trees are explained. B-trees, B+ trees, and red-black trees are more complex trees.

Chapter 9: Graphs. Unlike trees, graphs have no parent-child node relationship and have connections that can go in any direction. This chapter explains how graphs, a non-linear data structure, are used to handle problems with a large number of nodes and connections, such as network traffic, and path searching. Graph storage forms, including adjacency lists and adjacency matrices, are also explained.

Chapter 10: Practical Projects. This chapter applies the knowledge we learned in the previous chapters to solve practical problems. Distance algorithms, trie, filters, cache eviction algorithms, consistent hashing algorithms, and blockchain are covered. These practical projects help readers deepen their understanding of data structures and improve their Rust coding skills.

Acknowledgments

Rust is an excellent language that provides efficient, secure, and convenient engineering management tools, and it is gradually being integrated into the Linux kernel, making it a potential replacement for some C/C++ work in the future. Although several Rust books are available in the market, the author noticed that there are no algorithm books about Rust, which led to many obstacles in the learning process. Therefore, the author decided to create a simple and convenient Rust book that can help newcomers learn algorithms and data structures. After extensive research^[6], thinking, organizing, and combining the author's learning experience, this book was completed. Despite the steep learning curve of Rust, with the right direction and good resources, one can learn the language well. The author hopes that this book will make a small contribution to the Rust learning community.

The primary aim of writing this book is to learn and promote Rust and give back to the entire open-source community, which has enabled the author to learn and grow. The author expresses gratitude to PingCap for developing TiDB, the operating open-source community, and online courses. The author also thanks Zhang Handong, Mike Tang and other members of the Rust Chinese community for organizing Rust conferences and maintaining the community, Ling Hu Yi Chong for sharing Rust learning videos on Bilibili Danmu website, Zhang Handong for promoting Rust language through his book, "The Tao of Rust Programming," and RustMagazine Chinese monthly. The author also acknowledges the Rust Foundation^[7], established by Mozilla, AWS, Facebook, Google, Microsoft, and Huawei, for creating a platform that motivates the author to write this book, given the shortage of learning resources.

Finally, the author thanks the University of Electronic Science and Technology of China(UESTC) for providing resources and environment and the care and help of the author's mentor and fellow students in the KC404 teaching and research section where the author learned various technologies and cultures, grew up, and found the direction for his life journey.

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Chapter 1

Rust Basic

1.1 Objectives

- Install Rust and learn its toolchain
- Explore learning resources for Rust in various areas
- Review the fundamentals of the Rust programming language

1.2 Install Rust and its Toolchain

Ubuntu 22.04 and later versions come with pre-installed Rust, so you can start using it right away. If you're using other Unix-like systems, use the command below to install Rust. For Windows, refer to the [official website](#). As the code in this book is written in Linux, we recommend readers use Linux or Mac OS to avoid any inconsistencies and errors caused by different environments.

```
$ curl --proto '=https' --tlsv1.2 \
  -sSf https://sh.rustup.rs | sh
```

After installing Rust on Linux, you need to set environment variables to enable the system to locate Rust tools like the `rustc` compiler. Add the following three lines to the end of `~/.bashrc`.

```
# Rust env-path variables
export RUSTPATH=$HOME/.cargo/bin
export PATH=$PATH:$RUSTPATH
```

Then, save and execute `source ~/.bashrc`. If you find this process cumbersome, you can download the `install_rust.sh` script from the first chapter of the source code of this book and run it to complete the Rust toolchain installation. The instructions above will install the `rustup` tool, which manages and installs the Rust toolchain, including the `rustc` compiler, `cargo` project management tool, `rustup` toolchain management tool, `rustdoc` documentation tool, `rustfmt` formatting tool, and `rust-gdb` debugging tool.

For simple projects, you can use the `rustc` compiler for compilation. However, for larger projects, you need the `cargo` tool for management, which internally calls `rustc` for compilation. `Cargo` is an excellent tool that handles project building, testing, compilation, and publishing, making it highly efficient. Programmers who have worked with C/C++ projects will appreciate the `cargo` tool's greatness in handling various libraries and dependencies. Most of the time, we use the `rustc` compiler instead of the `cargo` tool, as the projects in this book are not too complex. `Rustup` manages the installation, upgrade, and uninstallation of Rust tools. Note that the Rust language includes stable and nightly versions, both of which can coexist. You can install Nightly using the following command.

```
$ rustup default nightly
```

After installation, you can use rustup to check the current version in use.

```
$ rustup toolchain list
stable-x86_64-unknown-linux-gnu
nightly-x86_64-unknown-linux-gnu (default)
```

To switch between stable and nightly versions, please use the following command.

```
$ rustup default stable # nightly
```

1.3 Learning Resources

Rust is a systems programming language and has a certain level of difficulty. It is impossible to learn Rust without good resources. Below is a collection of learning materials in various fields of Rust compiled by the author. It may not be complete, but most fields are covered.

1.3.1 Books and Documents

Basic: *The Rust Programming Language*, *Easy to Understand Rust*, *Rustlings*, *Rust by Example*, *Rust Primer*, *Rust Cookbook*, *Rust in Action*, *Rust Course*

Advanced: *Cargo Documentation*, *The Tao of Rust Programming*, *Learn Rust With Entirely Too Many Linked Lists*, *Rust Design Patterns*

High-level: *rustc book*, *The Little Book of Rust Macros*, *The Rustonomicon*, *Asynchronous Programming in Rust*

1.3.2 Related Topics

Wasm: <https://wasmer.io>, <https://wasmtime.dev>, <https://wasmedge.org>

HTTP/3: <https://github.com/cloudflare/quiche>

Algorithms: <https://github.com/TheAlgorithms/Rust>

Games: <https://github.com/bevyengine/bevy>

Tools: <https://github.com/rustdesk/rustdesk>, <https://github.com/uutils/coreutils>

Blockchain: <https://github.com/w3f/polkadot>

Databases: <https://github.com/tikv>, <https://github.com/tensorbase/tensorbase>

Compilers: https://github.com/rust-lang/rustc_codegen_gcc

Operating Systems: <https://github.com/Rust-for-Linux>, <https://github.com/rcore-os>

Web Front-end: <https://github.com/yewstack/yew>, <https://github.com/denoland/deno>

Web Back-end: <https://actix.rs/>, <https://github.com/tokio-rs/axum>, <https://github.com/poem-web/poem>

1.3.3 Community and Blogs

Rust Official: <https://www.rust-lang.org>

Rust Source Code: <https://github.com/rust-lang/rust>

Rust Documentation: <https://doc.rust-lang.org/stable>

Rust Reference: <https://doc.rust-lang.org/reference>

Awesome Rust: <https://github.com/rust-unofficial/awesome-rust>

Rust lib/crate: <https://crates.io>, <https://lib.rs>

Rust Sources: <https://www.yuque.com/zhoujiping/programming/rust-materials>

Rust Cheat Sheet: <https://cheats.rs>

1.4 Review

Similar to C/C++, Rust is a systems programming language. This means that concepts learned in those languages can be applied to understand Rust better. However, Rust introduces unique concepts like mutability, ownership, borrowing, and lifetimes, which can be both advantages and challenges.

1.4.1 The History of Rust

Rust is a highly efficient and reliable general-purpose compiled language that utilizes LLVM as its backend. It offers a balance between development efficiency and execution efficiency, making it a popular language among developers. The language has consistently been ranked as the most loved language by developers on Stack Overflow for many years. In 2021, Google, Amazon, Huawei, Microsoft, and Mozilla established a foundation for Rust, and Google even funded the rewriting of the Internet infrastructure Apache httpd and OpenSSL in Rust. Rust also has its own mascot, a red crab named [Ferris](#).

Rust was originally a personal project of Mozilla employee Graydon Hoare, and it received support from the Mozilla Research Institute starting in 2009. Rust was released to the public in 2010. Mozilla supported Rust because the Firefox Gecko engine was behind the times due to various vulnerabilities, historical baggage, and performance bottlenecks. Rust replaced the OCaml-written compiler implementation for self-hosting in 2010-2011 and released version 1.0 in 2015. Rust has a strong and active [community](#), and a stable and testing version are released every six weeks, with a major edition released every three years. Rust employs modern engineering management tool Cargo and has over 100,000 packages published on [crates.io](#).

In addition to development efficiency, Rust also exhibits excellent performance. In 2017, a team of six Portuguese researchers conducted a survey ^[8] on the performance of various programming languages, and Rust was found to be highly performant. They wrote solutions to ten problems based on 27 languages using the same algorithm, and then ran these solutions, recording the electricity consumption, speed, and memory usage of each language.

Total					
	Energy		Time		Mb
(c) C	1.00	(c) C	1.00	(c) Pascal	1.00
(c) Rust	1.03	(c) Rust	1.04	(c) Go	1.05
(c) C++	1.34	(c) C++	1.56	(c) C	1.17
(c) Ada	1.70	(c) Ada	1.85	(c) Fortran	1.24
(v) Java	1.98	(v) Java	1.89	(c) C++	1.34
(c) Pascal	2.14	(c) Chapel	2.14	(c) Ada	1.47
(c) Chapel	2.18	(c) Go	2.83	(c) Rust	1.54
(v) Lisp	2.27	(c) Pascal	3.02	(v) Lisp	1.92
(c) Ocaml	2.40	(c) Ocaml	3.09	(c) Haskell	2.45
(c) Fortran	2.52	(v) C#	3.14	(i) PHP	2.57
(c) Swift	2.79	(v) Lisp	3.40	(c) Swift	2.71
(c) Haskell	3.10	(c) Haskell	3.55	(i) Python	2.80
(v) C#	3.14	(c) Swift	4.20	(c) Ocaml	2.82
(c) Go	3.23	(c) Fortran	4.20	(v) C#	2.85
(v) Dart	3.83	(v) F#	6.30	(i) Hack	3.34
(v) F#	4.13	(i) JavaScript	6.52	(v) Racket	3.52
(i) JavaScript	4.45	(i) Dart	6.67	(i) Ruby	3.97
(v) Racket	7.91	(v) Racket	11.27	(c) Chapel	4.00
(i) TypeScript	21.50	(i) Hack	26.99	(v) F#	4.25
(i) Hack	24.02	(i) PHP	27.64	(i) JavaScript	4.59
(i) PHP	29.30	(v) Erlang	36.71	(i) TypeScript	4.69
(v) Erlang	42.23	(i) Jruby	43.44	(v) Java	6.01
(i) Lua	45.98	(i) TypeScript	46.20	(i) Perl	6.62
(i) Jruby	46.54	(i) Ruby	59.34	(i) Lua	6.72
(i) Ruby	69.91	(i) Perl	65.79	(v) Erlang	7.20
(i) Python	75.88	(i) Python	71.90	(i) Dart	8.64
(i) Perl	79.58	(i) Lua	82.91	(i) Jruby	19.84

The figure above provides readers with a clear understanding of the resource usage of various programming languages. Rust's energy consumption, time consumption, and memory consumption indicators are all very favorable. While this comparison may not be completely accurate, the overall trend is clear: Rust is energy-saving and efficient. Considering the critical point of climate change we currently face, the use of energy-saving and efficient languages like Rust to develop software is in line with the trend of Carbon Peaking and Carbon Neutrality Goals. I believe that it can be an important tool for enterprise transformation and look forward to the industry and society reaching a consensus.

Rust has a wide range of applications that extends to command-line tools, DevOps tools, audio and video processing, game engines, search engines, blockchain, the Internet of Things, browsers, cloud-native, network servers, databases, and operating systems. Many universities and enterprises at home and abroad are using Rust extensively. For example, Tsinghua University uses Rust in its new students' rCore and zCore operating systems, ByteDance's Feishu, remote desktop software RustDesk, PingCap's TiDB database, js/ts runtime Deno, and Google's Fuchsia operating system.

1.4.2 Keywords, Comments, Naming conventions

Rust uses 39 keywords (which may increase in the future), and some of these keywords, such as `Self` and `self`, can be challenging to learn.

<code>Self</code>	<code>enum</code>	<code>match</code>	<code>super</code>
<code>as</code>	<code>extern</code>	<code>mod</code>	<code>trait</code>
<code>async</code>	<code>false</code>	<code>move</code>	<code>true</code>
<code>await</code>	<code>fn</code>	<code>mut</code>	<code>type</code>
<code>break</code>	<code>for</code>	<code>pub</code>	<code>union</code>
<code>const</code>	<code>if</code>	<code>ref</code>	<code>unsafe</code>
<code>continue</code>	<code>impl</code>	<code>return</code>	<code>use</code>
<code>crate</code>	<code>in</code>	<code>self</code>	<code>where</code>
<code>dyn</code>	<code>let</code>	<code>static</code>	<code>while</code>
<code>else</code>	<code>loop</code>	<code>struct</code>	

Keywords are crucial to programming languages and reflect the designers' thought process behind program design. Different keywords indicate different trade-offs and considerations for various tasks. For instance, Rust's `match` is powerful, while C's similar `switch` functionality is less so, resulting in different coding styles and ways of thinking.

To facilitate code understanding and explain complex logic, appropriate explanations must be provided within the program. Comments serve as a useful place for these explanations. Rust supports two types of comments: regular comments and documentation comments. Regular comments have three comment formats.

```

1 // comment style 1
2
3 /* comment style 2 */
4
5 /*
6  * comment style 3
7  * multi-line-comments
8  * multi-line-comments
9  * multi-line-comments
10 */

```

Documentation and testing are highly emphasized in Rust, which is why it features a special documentation comment feature. This feature enables the writing of documentation and test code within comments, allowing code testing through cargo test and document generation through cargo doc.

```

1  /// This symbol placed at the top of a module file to control
2  /// the generation of library documentation and to describe
3  /// the functionality of the entire module.
4  /// This symbol is placed above the object being described to
5  /// control the generation of library documentation and to
6  /// describe functions or structures.

```

Rust's documentation uses Markdown syntax, so using the # symbol is essential.

```

1  /// Math mod    <-- doc comment, describe mode
2  ///
3  /// # Add <-- doc comment, discribe function, test case
4  /// This function sum the inputs
5  ///
6  /// # Example    <-- test code, use case
7  /// use math::add;
8  /// assert_eq!(3, add(1, 2));
9  fn add(x: i32, y: i32) -> i32 {
10     // sum        <-- regular comment
11     x + y
12 }

```

Naming conventions have been a topic of interest in programming, and Rust has its recommended practices. Rust suggests using UpperCamelCase for class-level content and snake_case for value-level content.

Item	Convention
Crate	snake_case
Type	UpperCamelCase
Trait	UpperCamelCase
Enum	UpperCamelCase
Function	snake_case
Method	snake_case
Constructor	new / with_more_details
Converter	from_other_type
Macros	snake_case!
Local variable	snake_case
Static variable	SCREAMING_SNAKE_CASE
Constants	SCREAMING_SNAKE_CASE
Type Param	The alphabeta of UpperCamelCase, like T, U, K
Lifetime	lowercase, such as 'a, 'src, 'dest

In UpperCamelCase, initialisms and abbreviations of compound words count as one word. For example, `Useize` should be used instead of `USize`. In snake_case or SCREAMING_SNAKE_CASE, words should not be composed of single letters unless it is the last word. Therefore, `btree_map` should be used instead of `b_tree_map`, and `PI_2` instead of `PI2`. The following code exemplifies Rust's naming conventions, which this book follows, and readers are encouraged to follow as well.

```

1  // Enum
2  enum Result<T, E> {
3      Ok(T),
4      Err(E),
5  }

```

```

6
7 // Trait
8 pub trait From<T> {
9     fn from<T> -> Self;
10 }
11
12 // Struct
13 struct Rectangle {
14     height: i32,
15     width: i32,
16 }
17 impl Rectangle {
18     // constructor
19     fn new(height: i32, width: i32) -> Self {
20         Self { height, width }
21     }
22
23     // function
24     fn calc_area(&self) -> i32 {
25         self.height * self.width
26     }
27 }
28
29 // static and constant variables
30 static NAME: &str = "kew";
31 const AGE: i32 = 25;
32
33 // Macro definition
34 macro_rules! add {
35     ($a:expr, $b:expr) => {
36         {
37             $a + $b
38         }
39     }
40 }
41
42 // variable and macro call
43 let sum_of_nums = add!(1, 2);

```

As Rust's popularity grows and its usage expands, a unified coding standard becomes necessary. Currently, Professor Zhang Handong leads a good [Rust Coding Standard](#), which readers can refer to.

1.4.3 Constants, Variables, Data Type

Variables and constants are fundamental concepts in programming, and Rust is no exception. However, Rust introduces mutability and immutability, which can be confusing, especially for beginners transitioning from other programming languages. In Rust, there are three types of values: constants, variables, and static variables. Novices may face difficulties when dealing with mutability and immutability, and they may have to struggle with the compiler to get their code to compile. By understanding the differences between constants, variables, and static variables, Rust programmers can write more concise and efficient code.

Constants are defined using `const` and are bound to a name that cannot be changed or reassigned.

```

1 // define constant(like #define used in C/C++)
2 const AGE: i32 = 1984;
3 // AGE = 1995; error, mutation not allowed
4
5 const NUM: f64 = 233.0;
6 // const NUM: f64 = 211.0; error, defined

```

Variables are defined using `let` and can have a value that is mutable or immutable depending on whether `mut` is used during its definition.

```

1 let x: f64 = 3.14; // let defines x which can be reassigned
2                  // but not mutated.
3 // x = 6.28; error, x is unmutable
4 let x: f64 = 2.71 // reassign x
5
6 let mut y = 985; // let mut defines y which can be reassigned
7                // and mutated.
8 y = 996; // y is mutable
9 let y = 2019; // reassign y

```

Finally, static variables are defined using `static` and can be mutable or immutable depending on whether `mut` is used during its definition.

```

1 static NAME: &str = "shieber" // static variable which can be
2                               // used as a constant
3 // NAME = "kew"; error,NAME is unmutable
4
5 static mut NUM: i32 = 100; // static mutable variable
6 unsafe {
7     NUM += 1; // NUM is mutable
8     println!("Num:{}",NUM);
9 }

```

The `"mut"` keyword is a constraint in Rust programming, restricting variables from being changed unless `"mut"` is added in front of it. Otherwise, the variable is immutable. This differs from other programming languages. Although static variables and constants have similarities, they're different in reality. Constants are replaced inline as many times as they're used, while static variables are referenced, with only one global instance. Static variables defined with `"static mut"` are wrapped in `"unsafe"` to indicate that they're not safe. It's recommended to use constants and variables instead, forget about static variables, and avoid coding errors.

Data types are the building blocks of a language, and Rust's data types are similar to C, while some are similar to Go. There are two types of basic data types in Rust: scalar and compound. Scalar types represent a single value, such as integers, floating-point numbers, Boolean types, and character types. Compound types combine multiple values into one, such as tuples and arrays, which are Rust's two native compound types.

Integers in Rust are numbers without a decimal part, classified into 12 types based on whether they're signed or unsigned and their length, denoted by `"i"` for signed types and `"u"` for unsigned types. A 64-bit machine can handle a 128-bit number by storing it in segments and processing it using multiple registers. `"isize"` and `"usize"` are integer types that match the machine architecture. So on a 64-bit machine, `"isize"` and `"usize"` represent `"i64"` and `"u64"`, respectively. On a 32-bit machine, they represent `"i32"` and `"u32"`.

Size	Signed	Unsigned
8	i8	u8

16	i16	u16
32	i32	u32
64	i64	u64
128	i128	u128
arch	isize	usize

Floating-point numbers in Rust are numbers with a decimal part, having two types: "f32" and "f64," with the default type being "f64," and both types being signed.

Size	Signed
32	f32
64	f64 (default)

The boolean type in Rust is represented by "bool" and has only two values: true and false, consistent with other programming languages.

The character type "char" in Rust is the most primitive type, similar to the C language. Characters are declared using single quotes, while string literals are declared using double quotes. The variables "c" and "c_str" are completely different types, with characters being four-byte Unicode scalar values, and strings represented as arrays.

```
1 // unicode scalar value
2 let c = 's';
3
4 // dynamic arrays
5 let c_str = "s";
```

Tuples are a type of composite value that combines multiple values of other types. Once declared, the length of a tuple cannot be increased or decreased. Tuples use parentheses to enclose the values, separated by commas. To retrieve values from a tuple, pattern matching and dot notation can be used, with indices starting at 0.

```
1 let tup1: (i8, f32, i64) = (-1, 2.33, 8000_0000);
2 // pattern match
3 let (x, y, z) = tup1;
4
5 let tup2 = (0, 100, 2.4);
6 let zero = tup2.0; // use symbol . to get value
7 let one_hundred = tup2.1;
```

In Rust, the let keyword is not just used to define variables, but can also destructure values through pattern matching. For example, x, y, and z can each obtain the values of -1, 2.33, and 80000000, respectively, through pattern matching. Defining variables is also a form of pattern matching, as let is essentially a pattern matching operation. The unit type in Rust is represented by an empty tuple (), and when there is only one value, it is written as (). The unit value is a special value of this type, and is implicitly returned when an expression does not return any value.

Unlike tuples, arrays in Rust must have elements of the same type, and the length cannot be changed once declared. If a mutable collection is needed, Vec can be used instead. It allows for dynamic resizing and is the preferred option in most situations.

```
1 // define arrays
2 let genders = ["Female", "Male", "Bigender"];
3 let gender_f = genders[0]; // indice element
4
5 // [type; num] define array
```

```

6 let digits[i32; 5] = [0, 1, 2, 3, 4];
7 let zeros = [0; 10]; // define an array which holds ten '0'

```

Rust allows for explicit type conversion between data types using the `as` keyword, but does not provide implicit type conversion between primitive types. Integer conversions in Rust follow the conventions of the C language, and all integer conversions are well-defined. For instance, converting `i8` to `i32` is a reasonable conversion that can be done explicitly.

```

1 // type_transfer.rs
2 #![allow(overflowing_literals)] // Ignore overflow warnings
3                                 // for type conversion.
4 fn main() {
5     let decimal = 61.3214_f32;
6     // let integer: u8 = decimal; // Error, f32 cannot be
7                                 // converted to u8.
8     let integer = decimal as u8; // Correct, use as
9     let character = integer as char;
10    println!("1000 as a u16: {}", 1000 as u16);
11    println!("1000 as a u8: {}", 1000 as u8);
12 }

```

For some complex types, Rust also provides two traits, `From` and `Into`, for conversion.

```

1 pub trait From<T> {
2     fn from<T> -> Self;
3 }
4 pub trait Into<T> {
5     fn into<T> -> T;
6 }

```

With these two traits, you can provide conversion functionality for various types.

```

1 // integer_to_complex.rs
2 #[derive(Debug)]
3 struct Complex {
4     real: i32, // real quantity
5     imag: i32 // imaginary quantity
6 }
7
8 // Implement a conversion from i32 to a complex number, where
9 // the i32 is converted to the real part and the imaginary
10 // part is set to 0.
11 impl From<i32> for Complex {
12     fn from(real: i32) -> Self {
13         Self { real, imag: 0 }
14     }
15 }
16
17 fn main() {
18     let c1: Complex = Complex::from(2_i32);
19     // implemented Into by default
20     let c2: Complex = 2_i32.into();
21     println!("c1: {:?}, c2: {:?}", c1, c2);
22 }

```

1.4.4 Statement, Expression, Computers and Flow Control

In Rust, there are two basic syntax categories: Statements and Expressions. Statements refer to operations to be executed, while expressions are simply used to evaluate values. Rust statements include declaration statements and expression statements. Declaration statements are used to declare variables, static variables, constants, structs, functions, external packages, and external modules. Statements that end with a semicolon are expression statements.

```

1 // Declaration statement for function
2 fn sum_of_nums(nums: &[i32]) -> i32{
3     nums.iter().sum:<i32>()
4 }
5
6 let x = 5;      // The entire line is a statement, x = 5 is an
7                // expression that calculates the value of x
8 x + 1;         // The entire line is an expression
9 let y = x + 1; // The entire line is a statement, y = x + 1
10              // is an expression
11 println!("{y}");
12
13 let z = [1,2,3];
14 println!("sum is {:?}", sum_of_nums(&z));

```

Operators are symbols used to specify the type of calculation performed by an expression. Rust supports common types of operators including arithmetic, relational, logical, assignment, and reference operators. Here are some Rust operators and their functions:

Operator	Example	Explanation
+	expr + expr	addition
+	trait + trait, 'a + trait	compound type constraint
-	expr - expr	subtraction
-	- expr	negation
*	expr * expr	multiply
*	*expr	dereference
*	*const type, *mut type	raw pointer
/	expr / expr	division
%	expr % expr	remainder
=	var = expr, ident = type	assignment/equality
+=	var += expr	addition and assignment
-=	var -= expr	subtraction and assignment
*=	var *= expr	multiply and assignment
/=	var /= expr	division and assignment
%=	var %= expr	remainder and assignment
==	expr == expr	equality comparison
!=	var != expr	inequality comparison
>	expr > expr	greater-than comparison
<	expr < expr	less-than comparison
>=	expr >= expr	>= comparison
<=	expr <= expr	<= comparison
&&	expr && expr	logical AND
	expr expr	logical OR
!	!expr	bitwise NOT or logical NOT

<code>&</code>	<code>expr & expr</code>	bitwise AND
<code>&</code>	<code>&expr, &mut expr</code>	borrow
<code>&</code>	<code>&type, &mut type,</code>	pointer/reference
<code> </code>	<code>pat pat</code>	pattern matching
<code> </code>	<code>expr expr</code>	bitwise OR
<code>^</code>	<code>expr ^ expr</code>	bitwise XOR
<code><<</code>	<code>expr << expr</code>	left shift
<code>>></code>	<code>expr >> expr</code>	right shift
<code>&=</code>	<code>var &= expr</code>	bitwise AND and assignment
<code> =</code>	<code>var = expr</code>	bitwise OR and assignment
<code>^=</code>	<code>var ^= expr</code>	bitwise XOR and assignment
<code><<=</code>	<code>var <<= expr</code>	left shift and assignment
<code>>>=</code>	<code>var >>= expr</code>	right shift and assignment
<code>.</code>	<code>expr.ident</code>	member access
<code>..</code>	<code>.., expr..,</code>	
	<code>..expr, expr..expr</code>	right-exclusive range
<code>..</code>	<code>..expr</code>	struct update syntax
<code>..=</code>	<code>..=expr, expr..=expr</code>	right-inclusive range
<code>:</code>	<code>pat: type, ident: type</code>	type constraint
<code>:</code>	<code>ident: expr</code>	struct field initialization
<code>:</code>	<code>'a: loop {...}</code>	loop label
<code>;</code>	<code>[type; len]</code>	fixed-size array type
<code>=></code>	<code>pat => expr</code>	match arm
<code>@</code>	<code>ident @ pat</code>	pattern binding
<code>?</code>	<code>expr?</code>	error propagation
<code>-></code>	<code>fn(...) -> type,</code>	
	<code> ... -> type</code>	function or closure return type

The ability to decide whether to execute a piece of code based on a condition or to repeatedly run a piece of code depending on whether a condition is satisfied is a feature found in most programming languages. In Rust, `if` expressions and loops are the most common structures used to control execution flow.

```

1 let a = 3;
2
3 if a > 5 {
4     println!("Greater than 5");
5 } else if a > 3 {
6     println!("Greater than 3");
7 } else {
8     println!("less or equal to 3");
9 }
```

`if` and `let` can also be combined to control code execution. One way to do this is with a `let if` statement, which returns the result to the `let` section if the condition is met. Note that there is a semicolon at the end because `let c = ..;` is a statement.

```

1 let a = 3; let b = 2;
2 let c = if a > b {
3     true
```

```

4         } else {
5             false
6         };

```

Another way is with an if let statement, which executes code by pattern matching the value on the right.

```

1 let some_value = Some(100);
2 if let Some(value) = some_value {
3     println!("value: {value}");
4 } else {
5     println!("no value");
6 }

```

In addition to the if let matching statement, match can also be used to control code execution.

```

1 let a = 10;
2 match a {
3     0 => println!("0 == a"),
4     1..=9 => println!("1 <= a <= 9"),
5     _ => println!("10 <= a"),
6 }

```

Rust provides various loop types, including loop, while, and for in. Using the continue and break keywords, you can jump and stop code execution on demand. The loop keyword controls repeated code execution until a condition is met.

```

1 let mut val = 10;
2 let res = loop {
3     // break loop and return value
4     if val < 0 {
5         break val;
6     }
7
8     val -= 1;
9     if 0 == val % 2 {
10        continue;
11    }
12
13    println!("val = {val}");
14 }; // a semi-colon here
15
16 // dead loop
17 loop {
18     if res > 0 { break; }
19
20     println!("{res}");
21 } // no semi-colon here

```

In contrast, a while loop computes the loop condition externally.

```

1 let num = 10;
2 while num > 0 {
3     println!("{}", num);
4     num -= 1;

```

```

5  }
6
7  let nums = [1,2,3,4,5,6];
8  let mut index = 0;
9  while index < 6 {
10     println!("val: {}", nums[index]);
11     index += 1;
12 }

```

Iterating through an array in Rust can be achieved through the `for in` loop, which is more convenient than using index and length.

```

1  let nums = [1,2,3,4,5,6];
2
3  // iterate over array
4  for num in nums {
5     println!("val: {num}");
6  }
7
8  // iterate over array in reverse order
9  for num in nums.iter().rev() {
10     println!("val: {num}");
11 }

```

This approach is not only concise, but also reduces the chance of errors, as it avoids specifying the array length explicitly. Additionally, `iter().rev()` can be used to traverse the array in reverse order.

The combination of `while` and `let` can be used to form a pattern match, eliminating the need to write a stop condition. The `let` syntax automatically determines when the condition is met before continuing `while`.

```

1  let mut v = vec![1,2,3,4,5,6];
2  while let Some(x) = v.pop() {
3     println!("{x}");
4  }

```

Overall, Rust provides many useful methods to control code flow, including `match`, `if let`, `let if`, `while let`, which align with Rust's coding specifications and are recommended.

1.4.5 Function, Program Structure

Functions are a crucial building block in programming languages, and in Rust, they are defined using the `fn` keyword followed by a snake-case function name. Function arguments are enclosed in parentheses and written as `val: type`, for example, `x: i32`. The return value is specified using `-> res` and may or may not have a value. The function implementation is wrapped in curly braces. Like C, Rust has a main function.

```

// define a function
fn func_name(parameters) -> return_types {
    code_body; // code body

    return_value // return value, no semi-colon here
}

```

Define a summation function.

```

1 // main function
2 fn main() {
3     let res = add(1, 2);
4     println!("1 + 2 = {res}");
5 }
6
7 fn add(a: i32, b: i32) -> i32 {
8     a + b
9 }

```

It is important to note that function definitions can be written before or after the main function, and the return value can be written directly on the value line without adding a semicolon, although this syntax is unique to Rust and adding a semicolon is not recommended. Functions in Rust modules are private by default and require the `pub` keyword to export them to other programs.

In Rust, a program is composed of several parts.

```

package/lib/crate/mod
variable
statement/expression
function
trait
label
comment

```

The following example shows the various elements of a program.

```

1 // rust_example.rs
2
3 // import module from standard library
4 use std::cmp::max;
5
6 // public module
7 pub mod math {
8     // public function
9     pub fn add(x: i32, y: i32) -> i32 { x + y }
10
11     // private function
12     fn is_zero(num: i32) -> bool { 0 == num }
13 }
14
15 // struct
16 #[derive(Debug)]
17 struct Circle { radius: f32, // radius }
18
19 // implement a converter to convert f32 to Circle
20 impl From<f32> for Circle {
21     fn from(radius: f32) -> Self {
22         Self { radius }
23     }
24 }
25
26 // comment: custom function

```

```

27 fn calc_func(num1:i32, num2:i32) -> i32 {
28     let x = 5;
29     let y = {
30         let x = 3;
31         x + 1 // expression
32     }; // statement
33
34     max(x, y)
35 }
36
37 // use function from math module
38 use math::add;
39
40 // main function
41 fn main() {
42     let num1 = 1; let num2 = 2;
43
44     // call function
45     println!("num1 + num2 = {}", add(num1, num2));
46     println!("res = {}", calc_func(num1, num2));
47
48     let f: f32 = 9.85;
49     let c: Circle = Circle::from(f);
50     println!("{:?}", c);
51 }

```

1.4.6 Ownership, Scope, Lifetime

Rust's ownership system is a crucial concept that deals with resource management, dangling references, and other complexities in the type system. It introduces the concepts of ownership, borrowing, and lifetime to restrict the state and scope of various quantities. The scope rules in Rust are the strictest among all programming languages.

In Rust, memory is managed through the ownership system, which is unlike garbage collection or manual memory release in other languages. The compiler checks a set of ownership rules at compile time, and if any rule is violated, the program doesn't even compile. The ownership rules are simple:

- Each value has an owner (variable).
- Values have only one owner at any given time.
- When the owner leaves the scope, the value is discarded.

This means that values in Rust are managed by a unique object, and memory is released as soon as they are no longer in use.

Rust's ownership rules make it more memory efficient because it releases unused memory as soon as it is no longer needed, which can then be reused by later variables. In contrast, languages that use garbage collection, such as in Go, which reclaims memory through the tricolor method and could lead to stop the world problems and program lag, or manual memory release may cause memory leaks as unused memory may not be released. The ownership mechanism is achieved through the automatic call of the drop method when the variable leaves the scope.

```

1 fn main() {
2     let long = 10;      <--- long enters scope (main)
3
4     { // this is a new temp scope
5         let short = 5;  <--- short enters scope (temp)

```



```

6      println!("inner short: {}", short);
7
8      let long = 3.14; <--- long enters scope (temp)
9      println!("inner long: {}", long);
10     }                <--- long and short leave scope (temp)
11
12     let long = 'a';    <--- long reassigns to a char
13     println!("outer long: {}", long);
14 }                    <--- long leaves scope (main)

```

The above shows the scope of each variable. The ownership mechanism is achieved by automatically calling the drop method of the variable when it leaves the scope (e.g. `}` in this case). Note that the internal `long` and the external `long` are two different variables, and the internal `long` does not override the external `long`.

To illustrate Rust's ownership mechanism, we can compare it to human society. For example, when a reader buys a book, they own it. If a friend borrows the book, the ownership still belongs to the reader. However, if the reader gives the book to the friend, the ownership transfers to the friend. Rust also has the concepts of "borrow" and "move," which are illustrated in the below example.

```

1 fn main() {
2     let x = "Shieber".to_string(); // create a string
3     let y = x; // move ownership of x to y
4     // println!("{}", x); // x is no longer valid
5 } <--- y calls drop

```

Let `y = x`; make a move, which transfers ownership of `x` to `y`. Some readers may wonder, "Isn't it released only when you leave the scope?", `x` and `y` are still in the `main` scope, how can it report an error? Actually, the second rule of ownership mechanism is very clear: a value can have only one owner at any time. So after the variable `x` is moved, it is immediately released and cannot be used later. Its scope only goes to the line `let y = x`; and not to the line `}`. In addition, this mechanism ensures that only `y`, not `x`, is freed at `}`, avoiding the memory safety problem of secondary freeing. To use both `x` and `y`, you can do something like the following.

```

1 fn main() {
2     let x = "Shieber".to_string(); // create a string
3     let y = &x; // borrow x
4     println!("{}", x); // x is still valid
5 } <--- y calls drop

```

`let y = &x`; is called a borrowing, `&x` is a reference to `x`. A reference is like a pointer (address) from which you can access data stored at that address that belongs to another variable, and a reference is created to be borrowed by someone else.

The book we borrowed cannot, by definition, be altered in any way, i.e. it is immutable. However, there's nothing wrong with a friend sketching over it. Similarly, there are mutable and immutable variables borrowed from Rust, and the `let y = &x`; above shows that `y` is just a non-variable borrowed from `x`. If you want mutable, you need to add the `mut` modifier.

```

1 fn main() {
2     let x = "Shieber".to_string(); // create a string
3     let y = &mut x; // borrow x mutably
4     y.push_str(", handsome!");
5     // let z = &mut x; // cannot borrow x mutably twice
6     println!("{}", x); // x is still valid
7 }

```

On line: `let z = &mut x;`, an error is reported, indicating that there can be only one variable borrow. This is done to avoid data competition, such as writing data at the same time. More than one immutable reference can exist at the same time, because multiple immutable references do not affect the variable; only multiple mutable references cause errors.

Now look at another situation. If the reader buys an electronic version of this book, then you can give a copy to your friend by copying it (Please follow the intellectual property laws), so you both have a copy of the book, and both have ownership of the book. Extending this concept in Rust gives us "copy", "clone".

```
1 fn main() {
2     let x = "Shieber".to_string(); // create a string
3     let y = x.clone(); // clone x
4     println!("{x}, {y}"); // x and y both are valid
5 }
```

The clone function, as the word literally implies, makes a deep copy of the data to y. Borrowing is just getting a valid pointer, which is fast, while cloning requires copying the data, which is less efficient and doubles the memory consumption. If the reader tries to write the following without the clone function, and finds that it compiles and runs without errors, does it not satisfy the ownership rule?

```
1 fn main() {
2     let x = 10; // create an integer on the stack
3     let y = x;
4     println!("{x}, {y}"); // x and y both are valid. paradox?
5 }
```

In fact, `let y = x;` does not give 10 to y, but automatically copies a new 10 to y, so that x and y each have a 10, so there is no conflict with the ownership rule. Because these simple variables are on the stack, Rust implements a uniform trait for this kind of data called Copy, which allows for fast copying without releasing the old variable x. In Rust, values, booleans, characters, etc. all implement the Copy trait, so moving such variables is equivalent to copying. Here you can call clone as well, but it's not necessary.

As mentioned earlier, a reference is a valid pointer. However, invalid pointers are often found, similar to the invalid dangling pointers in other programming languages.

```
1 fn dangle() -> &String {
2     let s = "Shieber".to_string();
3     &s
4 }
5
6 fn main() {
7     let ref_to_nothing = dangle();
8 }
```

The above code will compile with an error message similar to the following (with deletions).

```
error[E0106]: missing lifetime specifier
--> dangle.rs:1:16
1 | fn dangle() -> &String {
  |               ^ expected named lifetime parameter
  | help: function's return type contains a borrowed value,
  | but there is no value for it to be borrowed from
  | help: consider using the 'static lifetime
1 | fn dangle() -> &'static String {
  |               ~~~~~~
```

In fact, the problem can be found by analyzing the code or the error message, the function returned an invalid reference.

```
1 fn dangle() -> &String { <--- return a reference to a String
2     let s = "Shieber".to_string();
3     &s <---- return a reference to s
4 }           <---- s is dropped here, and it's memory goes away
```

According to the ownership analysis, `s` release is satisfied, `&s` is a pointer, returned, and seems to be fine, at most the pointer position is invalid. `s` and `&s` are two different things, the ownership system can only check the data according to the three rules, but it is impossible to know that the address `&s` points to is actually invalid. So why does the compilation error? The error message indicates that the lifetime specifier is missing. You can see that the suspension reference and the lifetime are in conflict, and that's why the error is reported. This means that even if your ownership system passes, but the lifetime does not, the error will be reported.

In fact, every reference in Rust has its own lifetime, i.e., the scope for which the reference remains valid. The ownership system does not guarantee that the data is absolutely valid, and requires a lifetime to ensure validity. Most of the time the lifetime is implicit and can be inferred, just as most of the time the type can be automatically inferred. When there is a reference in the scope, it is necessary to indicate the lifetime to show the interrelationship, so that the reference actually used at runtime is absolutely valid.

```
1 fn main() {
2     let a;                // -----+'a, a lifetime begins
3                           // |
4     {                    // |
5         let b = 10;       // --+'b      b lifetime begins
6                           // |
7         a = &b;           // -+        b lifetime ends
8     }                    // |
9                           // |
10    println!("a: {}", a); // -----+    a lifetime ends
11 }
```

a references `b`, and `a`'s lifetime `'a` is longer than `b`'s lifetime `'b`, so the compiler compiles with an error. For `a` to reference `b` properly, then `b`'s lifetime must be at least as long as the end of `a`'s lifetime. By comparing lifetimes, Rust can find references that don't make sense and thus avoid dangling reference problems.

To use variables legally, Rust requires that all data carry a lifetime specifier. Lifetimes are indicated by single quotes `'` followed by a letter, such as `&'a`, `&mut 't`. References in functions also require lifetime markers.

```
1 fn longest<'a>(x: &'a String, y: &'a String) -> &'a String {
2     if x.len() < y.len() {
3         y
4     } else {
5         x
6     }
7 }
```

The pointed brackets are lifecycle arguments, which are generic arguments that need to be declared in the pointed brackets of the function name and argument list to indicate that both arguments and the returned reference will live as long as they do. Because Rust automatically infers the lifetime, it is recommended to omit the lifetime specifier.

```

1 fn main() {
2     // static lifetime, lives for the entire duration
3     let s: &' static str = "Shieber";
4
5     let x = 10;
6     let y = &x; // can rewrite as let y = &'a x;
7     println!("y: {}", y);
8 }

```

In this section, we learn about ownership system, borrowing, cloning, scope rules, lifetime. The ownership system is a memory management mechanism, borrowing and cloning are ways of using variables that extend the ownership system, and the lifetime is a complement to the ownership system, used to solve problems such as dangling references that the ownership system cannot handle.

1.4.7 Generic, Trait

To add two i64s, you can implement an add function of type i64 called add_i64.

```

1 fn add_i64(x: i64, y: i64) -> i64 {
2     x + y
3 }

```

However, if you were to implement add functions for all numeric types, the code would become lengthy and the naming of functions would become cumbersome, leading to function names like add_i8, add_i16, and so on. The addition operation should be universal for any numeric type, and writing multiple sets of code is unnecessary. To tackle this problem, Rust uses generics. Generics are abstract substitutes for concrete types or properties, allowing developers to express the properties of the generic type without knowing its actual type when writing code.

```

1 // generic function: add
2 fn add<T>(x: T, y: T) -> T {
3     x + y
4 }

```

The add function uses a generic type, and the generic parameter T is placed inside the parentheses immediately after add in the function declaration to indicate that this is a generic function. The generic parameter T represents any numeric type, and the return value type and the type of arguments involved in the function are both T, allowing it to handle all numeric types.

In addition to functions, generics can be used for other program components, such as enums. The Result in Rust is an enum type, which handles errors, uses the generic parameters T and E, where T represents the return value type on success and E represents the return value type on error.

```

1 enum Result<T, E> {
2     Ok(T),
3     Err(E),
4 }

```

Generic types can also be used for commonly used structs

```

1 struct Point<T> {
2     x: T,
3     y: T,
4 }
5

```

```

6 let point_i = Point { x: 3, y: 4 };
7 let point_f = Point { x: 2.1, y: 3.2 };
8 // let point_m = Point { x: 2_i32, y: 3.2_f32 }; Error

```

The generic parameter `T` is used to constrain the input parameters, ensuring that both parameters are of the same type. For instance, for the `Point` struct, it would be incorrect if `x` used `i32` and `y` used `f32`.

Although a generic set of code has been implemented for the `add` function using generics, there are still some issues. The `T` parameter is not guaranteed to be a numeric type, so calling `add` on a type that does not support addition would be incorrect. It would be preferable to restrict the generic argument type of `add` to only numeric types. A trait in Rust is a way to define and restrict the behavior of a generic type, encapsulating the functionality shared by each type. Using traits, the behavior of each type can be well-controlled, and a trait can consist of three parts: method, type, and constant. It defines an abstract interface that can be implemented by a type or directly inherited from its default implementation. For example, if you have two types, `teacher` and `student`, and both have a greeting behavior, you can implement the following trait:

```

1 trait Greete {
2     // default implementation
3     fn say_hello(&self) {
4         println!("Hello!");
5     }
6 }
7
8 // contains different properties
9 struct Student {
10     education: i32, // education years
11 }
12 struct Teacher {
13     education: i32, // education years
14     teaching: i32, // teaching years
15 }
16
17 impl Greete for Student {}
18
19 impl Greete for Teacher {
20     // override default implementation
21     fn say_hello(&self) {
22         println!("Hello, I am teacher Zhang!");
23     }
24 }
25
26 // generic constraint
27 fn outer_say_hello<T: Greete>(t: &T) {
28     t.say_hello();
29 }
30
31 fn main() {
32     let s = Student{ education: 3 };
33     s.say_hello();
34
35     let t = Teacher{ education: 20, teaching: 2 };
36     outer_say_hello(&t);
37 }

```

The `outer_say_hello` function above adds the generic constraint `T: Greete`, known as trait bound, indicating that only a type `T` that implements `Greete` can call the `say_hello` function. the previous `add` function can rewrite as below.

```
1 fn add<T: Addable>(x: T, y: T) -> T {
2     x + y
3 }
```

The `Addable` trait is a constraint that ensures that only types implementing the `Addable` trait can be added together. This allows any type that doesn't meet the constraint to be reported as an error at compile time rather than waiting until runtime.

Another way to write the trait constraint is by using the `impl` keyword, as shown in the following example. Here, `t` must be a reference implementing the `Greete` trait.

```
1 fn outer_say_hello(t: &impl Greete) {
2     t.say_hello();
3 }
```

Multiple trait bounds and multiple argument types can be combined using a comma and a plus sign.

```
1 fn some_func<T: trait1 + trait2, U: trait1> (x: T, y: U) {
2     do_some_work();
3 }
```

To avoid using multiple trait bounds in parentheses, Rust has introduced a `where` syntax that separates the trait bound from the parentheses.

```
1 // where clause
2 fn some_func<T, U> (x: T, y: U)
3     where T: trait1 + trait2,
4           U: trait1,
5 {
6     do_some_work();
7 }
```

The implementation of the `say_hello` method in `Greete` for `Student` and `Teacher` allows each type to encapsulate its own unique properties. While `Student` is more like a direct inheritance from `Greete`, `Teacher` overrides `Greete`. Although they use the same `say_hello` method, `Student` and `Teacher` display different states. The concepts of encapsulation, inheritance, and polymorphism are present here, which seems to imply that the concept of classes has been implemented through `impl`, `trait`, and thus object-oriented programming.

1.4.8 Enum and Match

When designing a questionnaire that requires multiple choice answers for gender, education, marital status, and other options, it is necessary to provide a variety of choices for participants to select. Enumerations in programming languages allow you to define a type by listing all possible members, similar to defining a structure. For instance, you can use an enumeration to represent gender.

```
1 enum Gender {
2     Male,
3     Female,
4     TransGender,
5 }
```

To use an enumeration in Rust, simply use "enum type::enum name".

```
1 let male = Gender::Male;
```

A common enumeration type in Rust is Option, where Some represents a value and None indicates no value.

```
1 enum Option<T> {  
2     Some(T),  
3     None  
4 }
```

Enumerations can also be combined with match for process control. A match expression compares a value to a series of patterns and executes the corresponding code based on the matching pattern. The patterns can include literal values, variables, wildcards, and other content. The matched values pass through each pattern of the match, and when the first matching pattern is encountered, it enters the associated block of code. Match expressions are like a classifier or sieve that filters values based on patterns. In Rust, the matched values can be filtered into more fine-grained classes. For example, an enumeration could represent the different denominations of currency values.

```
1 enum Cash { // US dollar($)  
2     One,  
3     Two,  
4     Five,  
5     Ten,  
6     Twenty,  
7     Fifty,  
8     Hundred,  
9 }  
10  
11 fn cash_value(cash: Cash) -> u8 {  
12     match cash {  
13         Cash::One => 1,  
14         Cash::Two => 2,  
15         Cash::Five => 5,  
16         Cash::Ten => 10,  
17         Cash::Twenty => 20,  
18         Cash::Fifty => 50,  
19         Cash::Hundred => 100,  
20     }  
21 }
```

Additionally, match supports wildcard and _ placeholder matches.

```
1 match cash {  
2     Cash::One => 1,  
3     Cash::Two => 2,  
4     Cash::Five => 5,  
5     Cash::Ten => 10,  
6     other => 0, // _ => 0,  
7 }
```

For instance, "other" can be replaced with _ if you don't need to use the value directly. Here other is used instead of all denominations greater than 10.

The match expression must be exhaustive, which means it should cover all possible matching patterns. However, if you only need to match a specific pattern, using match can be cumbersome. Fortunately, Rust generalizes the matching pattern of match and introduces the if let matching pattern. The if let match counts all bills larger than \$1.

```
1 let mut greater_than_one = 0;
2 if let Cash::One = cash { // only match Cash::One
3     println!("cash is one");
4 } else {
5     greater_than_one += 1;
6 }
```

1.4.9 Functional Programming

Rust prioritizes functional programming instead of class-based problem-solving methods used in object-oriented languages. This programming paradigm involves using functions as parameter values or returning them as the result of other functions. The key components of functional programming are closures and iterators.

Closures are anonymous functions that can be passed as parameters to other functions and capture variables. They can be created in one location and executed in different contexts. Closures allow the capturing of values in the caller's scope, making them particularly useful for defining functions that are only used once.

```
1 // define a function
2 fn function_name(parameters) -> return_types {
3     code_body;
4     return_value
5 }
6
7 // define closure
8 |parameters| {
9     cody_body;
10    return_value
11 }
```

Closures can be parameterless and may or may not return values. Rust infers closure parameter and return value types, eliminating the need to specify them explicitly. To use a closure, it must be assigned to a variable and called like a function. For instance, a closure function that determines odd or even numbers can be defined.

```
1 let is_even = |x| { 0 == x % 2 };
2
3 let num = 10;
4 println!("{num} is even: {}", is_even(num));
```

It's also possible to use external variables in closures.

```
1 let val = 2;
2 let add_val = |x| { x + val };
3
4 let num = 2;
5 let res = add_val(num);
6 println!("{num} + {val} = {res}")
```


The `add_val` function captures the external variable `val`. Closures can capture external variables by taking ownership, reference, or mutable reference, which Rust defines using three function traits: `FnOnce`, `FnMut`, `Fn`.

- The `FnOnce` trait consumes the captured variable from the surrounding scope and moves it into the closure when it is defined. This trait indicates that the closure can only be called once.
- The `FnMut` trait obtains a mutable borrowed value, allowing it to change the external variable.
- The `Fn` trait obtains an immutable borrowed value from its environment.

These traits correspond to the move, mutable reference, and reference implementation of the ownership system. All closures can be called at least once, making them implement the `FnOnce` trait. Closures that use mutable references but do not move variable ownership into the closure implement the `FnMut` trait, while those that do not require mutable access to the variables implement `Fn`.

The `move` keyword is used to force the ownership of an external variable to be moved into a closure.

```
1 let val = 2;
2 let add_val = move |x| { x + val };
3 // println!("{val}"); error: use of moved value: val
```

In Rust, the "for in" loop is used to iterate through an array, which is an iterator that implements the iteration functionality by default. Iterators pass each element in a collection to a processing logic in sequence, allowing specific operations to be performed on the items in a sequence. They are responsible for iterating through each item in the sequence and determining when to stop.

By default, iterators must implement the `Iterator` trait, which has two methods: `iter()` and `next()`.

```
iter(), return a iterator
next(), return the next element
```

The `iter()` method can be divided into three types depending on whether the data can be modified during iteration.

method	return type
<code>iter()</code>	return a immutable iterator whose element type is <code>&T</code>
<code>iter_mut()</code>	return a mutable iterator whose element type is <code>&mut T</code>
<code>into_iter()</code>	return a immutable iterator whose element type is <code>T</code> , original data consummed

In Rust, iterators can be classified into "reentrant" and "non-reentrant" types based on their behavior towards the original data after iteration. A "reentrant" iterator allows the original data to be used again after iteration, while a "non-reentrant" iterator consumes the original data. The `iter()`, `iter_mut()`, and `into_iter()` methods are used to implement iterators for reading values, changing values, and taking ownership of the original data, respectively.

```
1 let nums = vec![1,2,3,4,5,6];
2
3 // 1.iter immutable
4 for num in nums.iter() {
5     println!("num: {num}")
6 };
7 println!("{:?}", nums); // use nums after iter
8
9 // 2.iter_mut mutable
10 for num in nums.iter_mut() {
```

```

11     *num += 1;
12 }
13 println!("{:?}", nums); // use nums after iter_mut
14
15 // 3.into_iter transfer nums into iterator and consume nums
16 for num in nums.into_iter() {
17     println!("num: {num}");
18 }
19 // println!("{:?}", nums); error: use of moved value "nums"

```

In addition to transferring ownership of the original data, iterators can also be consumed or regenerated. Consumers are special operations on an iterator that convert the iterator into a value of another type. Examples of consumers include `sum`, `collect`, `nth`, `find`, `next`, and `fold`. They perform operations on the iterator to obtain the final value. Producers, on the other hand, are adapters that traverse the iterator and generate another iterator. Examples of adapters include `take`, `skip`, `rev`, `filter`, `map`, `zip`, and `enumerate`. In Rust, an iterator itself is considered an adapter.

```

1 // adapter_consumer.rs
2
3 fn main() {
4     let nums = vec![1,2,3,4,5,6];
5     let nums_iter = nums.iter();
6     let total = nums_iter.sum::<i32>(); // sum is a consumer
7
8     let new_nums: Vec<i32> = (0..100).filter(|&n| 0 == n % 2)
9                                     .collect(); // Adapter
10    println!("{:?}", new_nums);
11
12    // calculate the sum of all numbers less than 1000
13    // that are divisible by 3 or 5
14    let sum = (1..1000).filter(|n| n % 3 == 0 || n % 5 == 0)
15                .sum::<u32>();
16                // combine adapter and consumer
17    println!("{sum}");
18 }

```

The following code combines adapters, closures, and consumers to find the sum of all integers less than 1000 that are divisible by 3 or 5, showcasing the power of functional programming. This approach allows for complex calculations to be performed in a concise and efficient manner. In comparison, imperative programming may result in lengthy and difficult-to-understand code. Therefore, it is recommended to use closures with iterators, adapters, and consumers for functional programming.

```

1 fn main() {
2     let mut nums: Vec<u32> = Vec::new();
3     for i in 1..1000 {
4         if i % 3 == 0 || i % 5 == 0 {
5             nums.push(i);
6         }
7     }
8
9     let sum = nums.iter().sum::<u32>();
10    println!("{sum}");
11 }

```

Functional programming is a programming paradigm that is distinct from other paradigms such as imperative programming and declarative programming. Imperative programming is an abstraction of computer hardware, involving variables, assignment statements, expressions, control statements, etc. Structured programming and object-oriented programming, which are commonly used, fall under this paradigm. The main focus is on the steps that the computer executes, instructing it on what to do step by step.

In contrast, declarative programming expresses program logic as data structures, concentrating on what to do rather than how to do it. SQL statements are an example of declarative programming. While functional programming and declarative programming share a common idea of focusing on what to do, functional programming is not limited to declarative programming. It is an abstraction of mathematics, describing computation as the evaluation of expressions. In other words, a functional program is a mathematical expression. The functions in functional programming refer to mathematical functions, which map the independent variable to the dependent variable $y = f(x)$. The output of a function solely depends on the input parameter value, not on any other state. For instance, the `sin()` function in mathematics calculates the sine value of `x`. As long as `x` remains unchanged, the final result will always be the same, no matter how or when it is called.

1.4.10 Smart Pointer

Rust, being a systems language, still requires the use of pointers. Pointers refer to or point to other data by containing a memory address. The most common pointer in Rust is a reference, which is similar to C's concept of pointers but has no other functionality besides referencing data. On the other hand, smart pointers are data structures that behave like pointers and contain metadata. They provide additional functions such as memory management or binding checks, like managing file handles and network connections. In Rust, `Vec` and `String` can be considered smart pointers.

Smart pointers were first introduced in C++ and later adopted by Rust. Rust encapsulates two major traits for smart pointers: `Deref` and `Drop`. When a variable implements `Deref` and `Drop`, it becomes more than an ordinary variable. After implementing `Deref`, the variable overloads the dereference operator `**`, which can be used as an ordinary reference and can be automatically or manually dereferenced when necessary. After implementing `Drop`, the variable is automatically released from the heap when it goes out of scope. Other functions can also be customized, such as releasing files or network connections, similar to some languages' destructor functions. The following are the main features of smart pointers:

1. Smart pointers usually own the data they point to.
2. Smart pointers are data structures that are typically implemented using structures.
3. Smart pointers implement two major traits: `Deref` and `Drop`.

Many common smart pointers are available, and users can also implement the smart pointers they need. The following are frequently used smart pointers or data structures. Note that although `Cell` and `RefCell` share similarities with smart pointers, they are not considered smart pointers according to the above characteristics.

- `Box<T>`: A unique ownership smart pointer that points to data of type `T` stored on the heap.
- `Rc<T>`: A reference counting smart pointer that shares ownership and is used to track the number of references to a value stored on the heap.
- `Arc<T>`: A thread-safe reference counting smart pointer that shares ownership and can be used for multi-threading.
- `Cell<T>`: A container that provides interior mutability and is not a smart pointer. It allows for borrowed mutable data with compile-time checks, requiring `T` to implement the `Copy` trait.
- `RefCell<T>`: A container that provides interior mutability and is not a smart pointer. It allows for borrowed mutable data with run-time checks, with no requirement for `T` to implement the `Copy` trait.
- `Weak<T>`: A weak reference type that corresponds to `Rc` and is used to solve the problem of circular references in `RefCell`.
- `Cow<T>`: A copy-on-write enumerated smart pointer that is primarily used to reduce memory allocation and copying. It is suitable for scenarios with frequent reads and infrequent writes.

In Rust, Deref and Drop are the two most important traits for smart pointers. By implementing Deref and Drop for a custom data type similar to Box, we can better understand the difference between references and smart pointers.

```

1 // define tuple struct
2 struct SBox<T>(T);
3 impl<T> SBox<T> {
4     fn new(x: T) -> Self {
5         Self(x)
6     }
7 }
8
9 fn main() {
10     let x = 10;
11     let y = SBox::new(x);
12     println!("x = {x}");
13     // println!("y = {}", *y); cannot dereference a raw pointer
14 } <--- x, y call drop

```

Here is an example of implementation of Deref and Drop for SBox.

```

1 use std::ops::Deref;
2
3 // implement Deref for SBox
4 impl<T> Deref for SBox<T> {
5     type Target = T; // define associated type,
6                       // which is the type that Deref is
6                       // targeting to
7     fn deref(&self) -> &Self::Target {
8         &self.0 // .0 means the first element of tuple struct
9     }
10 }
11
12 // implement Drop for SBox
13 impl<T> Drop for SBox<T> {
14     fn drop(&mut self) {
15         println("SBox drop itself!"); // just print a message
16     }
17 }

```

If Deref is not implemented, a variable of our custom data type cannot be dereferenced. On the other hand, if Drop is not implemented, the variable will be released from the heap only when it goes out of scope. However, if the variable contains other references or smart pointers, they will not be released until the program exits, resulting in a memory leak. Therefore, it is crucial to implement Drop correctly to avoid memory leaks.

```

1 fn main() {
2     let x = 10;
3     let y = SBox::new(x);
4     println!("x = {x}");
5     println!("y = {}", *y); // *y equals *(y.deref())
6     // y.drop(); error: drop twice
7 } <--- x, y call drop and y prints "SBox drop itself!"

```

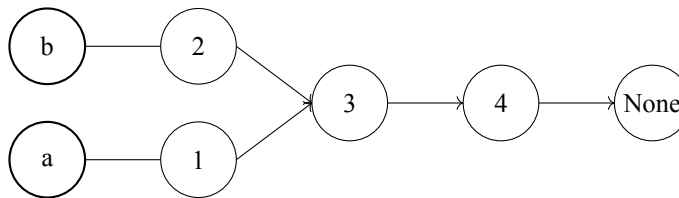
Box stores its data on the heap, and after implementing Deref, it can be automatically dereferenced.

```

1 fn main() {
2     let num = 10; // num stored in stack
3     let n_box = Box::new(num); // n_box stored in heap
4     println!("n_box = {}", n_box); // n_box deref to num
5                                     // automatically
6     println!("{}", 10 == *n_box); // n_box deref to num
7                                     // manually
8 }
```

The ownership system rules state that a value can only have one owner at any given time, but in some scenarios, we need values to have multiple owners. To address this, Rust provides the Rc smart pointer. Rc is a shareable reference counting smart pointer that can produce multiple ownership values. Reference counting means that it determines whether a value is still in use by keeping track of the number of references to it, and if the reference count is zero, the value can be cleaned up.

Rust's ownership system mandates that a value can have only one owner at any given time. However, in some cases, multiple ownership is necessary. To solve this problem, Rust provides the Rc smart pointer, which is a reference counting smart pointer that can produce multiple ownership values. It tracks whether a value is still in use by counting the number of references to it. Once the reference count drops to zero, the value can be cleaned up.



In the example shown, 3 is shared by variables a(1) and b(2), and cloning an Rc increases the reference count. Sharing is like turning off the lights in a classroom, only the last person leaving needs to turn off the lights. Similarly, each user of Rc will only clear the data on the last use. Cloning an Rc will increase the reference count, just like when a new person enters a classroom.

```

1 use std::rc::Rc;
2 fn main() {
3     let one = Rc::new(1);
4     // increase reference count
5     let one_1 = one.clone();
6     // display reference count
7     println!("sc:{}", Rc::strong_count(one_1));
8 }
```

Rc can be used for shared ownership in single-threaded environments only. For multithreaded environments, Rust provides Arc (atomic reference counting), a thread-safe version of Rc. It allocates a T type value with shared ownership on the heap. Cloning an Arc increases the reference count while producing a new Arc instance pointing to the same heap as the source Arc. Arc is immutable by default and requires locking mechanisms like Mutex to modify it between multiple threads.

By default, Rc and Arc do not allow internal data modification. Rust provides two containers with internal mutability, Cell and RefCell, for scenarios where modifying internal values is necessary. Internal mutability is a Rust design pattern that allows data modification while having immutable references. Cell provides methods for getting and changing internal values. For data types that implement Copy, the get method is used to view the internal value, while the take method replaces the internal value with the

default value and returns the replaced value. For all data types, the `replace` method replaces and returns the replaced value, while the `into_inner` method consumes `Cell` and returns the internal value.

```

1 // use_cell.rs
2
3 use std::cell::Cell;
4
5 struct Fields {
6     regular_field: u8,
7     special_field: Cell<u8>,
8 }
9
10 fn main() {
11     let fields = Fields {
12         regular_field: 0,
13         special_field: Cell::new(1),
14     };
15
16     let value = 10;
17     // fields.regular_field = value;
18     // error[E0594]: cannot assign to immutable field
19
20     fields.special_field.set(value);
21     // although fields is immutable,
22     // we can still change the value of special_field
23
24     println!("special: {}", fields.special_field.get());
25 }

```

To enable modification of certain fields internally, `Cell` provides a backdoor to immutable struct `Fields`.

`RefCell` has similar characteristics to `Cell`, but instead of using `get` and `set` methods to modify internal data, it directly obtains a mutable reference to the data. For example, `RefMut<_>` can be used to modify `HashMap` through `RefCell`. Here, `shared_map` obtains a `RefMut<_>` type map directly through `borrow_mut()`, and then adds an element to the map through `insert`, which modifies `shared_map`.

```

1 // use_refcell.rs
2
3 use std::cell::{RefCell, RefMut};
4 use std::collections::HashMap;
5 use std::rc::Rc;
6
7 fn main() {
8     let shared_map: Rc<RefCell<_>> =
9         Rc::new(RefCell::new(HashMap::new()));
10
11     {
12         let mut map: RefMut<_> = shared_map.borrow_mut();
13         map.insert("kew", 1);
14         map.insert("shieber", 2);
15         map.insert("mon", 3);
16         map.insert("hon", 4);
17     }
18 }

```

```

18     let total: i32 = shared_map.borrow().values().sum();
19     println!("{}", total);
20 }

```

While both `Cell` and `RefCell` can modify internal values, they do so differently. `Cell` replaces values directly, while `RefCell` modifies them through mutable references. Since `Cell` needs to replace and move values, it must implement `Copy`, while `RefCell` is suitable for data types that do not implement `Copy`. `Cell` is checked at compile time, while `RefCell` is checked at runtime, and misuse can lead to panics.

Although Rust provides memory safety guarantees, `RefCell` may still cause circular references and memory leaks. To prevent this, Rust provides the Weak smart pointer, which allows for the value to be cleared even when there are circular references. The `upgrade` method can be used to check if the referenced value is still valid. Each time `Rc` is cloned, the strong reference count `strong_count` of the instance is increased, and the instance is only cleared when `strong_count` is 0. `strong_count` will never be 0 in circular references. When using the Weak smart pointer, the value of `strong_count` is not increased, but the value of `weak_count` does. `weak_count` can be cleared without being 0, which solves the problem of circular references.

As Weak smart pointer, to ensure that the referenced value is still valid, its `upgrade` method can be called, which returns `Option<Rc<T>`. If the `Rc<T>` value has not been dropped, the result is `Some`. If dropped, the result is `None`. The following is an example of using Weak to solve circular references. `Car` and `Wheel` have mutual references, and circular references will occur if they are both used with `Rc`.

```

1  // use_weak.rs
2
3  use std::cell::RefCell;
4  use std::rc::{Rc, Weak};
5
6  struct Car {
7      name: String,
8      wheels: RefCell<Vec<Weak<Wheel>>>, // reference to Wheel
9  }
10
11 struct Wheel {
12     id: i32,
13     car: Rc<Car>, // reference to Car
14 }
15
16 fn main() {
17     let car: Rc<Car> = Rc::new(
18         Car {
19             name: "Tesla".to_string(),
20             wheels: RefCell::new(vec![]),
21         }
22     );
23
24     let w1 = Rc::new(Wheel { id:1, car: Rc::clone(&car) });
25     let w2 = Rc::new(Wheel { id:2, car: Rc::clone(&car) });
26
27     let mut wheels = car.wheels.borrow_mut();
28
29     // downgrade to Weak
30     wheels.push(Rc::downgrade(&w1));
31     wheels.push(Rc::downgrade(&w2));
32

```

```

33     for wheel_weak in car.wheels.borrow().iter() {
34         let wl = wheel_weak.upgrade().unwrap(); // Option
35         println!("wheel {} owned by {}", wl.id, wl.car.name);
36     }
37 }

```

Now, let's take a look at Cow (Copy on Write) Smart Pointer.

```

1  pub enum Cow<'a, B>
2      where B: 'a + ToOwned + 'a + ?Sized {
3      Borrowed(&'a B), // wrap a reference
4      Owned(<B as ToOwned>::Owned), // wrap a owned value
5  }

```

The concept of borrowing involves accessing borrowed content in an immutable way, while ownership means cloning the data when mutable borrowing is needed. To filter out all spaces in a string, we can use the following code.

```

1  // use_cow.rs
2  fn delete_spaces(src: &str) -> String {
3      let mut dest = String::with_capacity(src.len());
4      for c in src.chars() {
5          if ' ' != c {
6              dest.push(c);
7          }
8      }
9      dest
10 }

```

While this code works, its efficiency is not optimal. First, we need to decide whether to use `&str` or `String` for the parameter. If we use `String` but input `&str`, we need to clone it before calling the function. If we use `String`, the string will be moved to the interior after the call, and the external cannot use it anymore. In either case, a string generation and copy are performed in the function. If the string does not contain whitespace characters, it is best to return it as is without copying. This is where Cow comes in handy as it reduces copying and improves efficiency.

```

1  // use_cow.rs
2  use std::borrow::Cow;
3  fn delete_spaces2<'a>(src: &'a str) -> Cow<'a, str> {
4      if src.contains(' ') {
5          let mut dest = String::with_capacity(src.len());
6          for c in src.chars() {
7              if ' ' != c { dest.push(c); }
8          }
9          // capture the ownership and drop dest
10         return Cow::Owned(dest);
11     }
12     return Cow::Borrowed(src); // borrow the src
13 }

```

Here is an example that utilizes Cow.

```

1  // use_cow.rs
2  fn main() {
3      let s = "i love you";

```



```

4     let res1 = delete_spaces(s);
5     let res2 = delete_spaces2(s);
6     println!("{res1}, {res2}");
7 }

```

When there are no whitespace characters, the function returns directly, avoiding copying and improving efficiency.

```

1  // more_cow_usage.rs
2
3  use std::borrow::Cow;
4  fn abs_all(input: &mut Cow<[i32]>) {
5      for i in 0..input.len() {
6          let v = input[i];
7          if v < 0 { input.to_mut()[i] = -v; }
8      }
9  }
10 fn main() {
11     // read only, no write, no copy
12     let a = [0, 1, 2];
13     let mut input = Cow::from(&a[..]);
14     abs_all(&mut input);
15     assert_eq!(input, Cow::Borrowed(a.as_ref()));
16     // copy on write, copy when read -1
17     let b = [0, -1, -2];
18     let mut input = Cow::from(&b[..]);
19     abs_all(&mut input);
20     assert_eq!(input, Cow::Owned(vec![0,1,2])) as Cow<[i32]>);
21     // no copy on write, because already owned
22     let mut c = Cow::from(vec![0, -1, -2]);
23     abs_all(&mut c);
24     assert_eq!(c, Cow::Owned(vec![0,1,2])) as Cow<[i32]>);
25 }

```

This section contains a lot of diverse content, and readers are encouraged to study it further in conjunction with other dedicated books. The information provided here is only brief and not deeply discussed.

1.4.11 Exception

Rust provides a unique exception handling mechanism as it does not have try-catch blocks like other languages. This section is titled "Exceptions," but the author has used the term to refer to failures, errors, and exceptions collectively. There are four types of exceptions in Rust: Option, Result, Panic, and Abort.

Option is used to handle possible failure cases, indicating whether the operation succeeded or failed with Some and None. For instance, when getting a value that may not exist, the result may be None, which should not cause an error but needs to be handled appropriately. Failure is different from error, as it is an anticipated outcome that will not cause problems for the program. The definition of Option, which we have already learned, is provided below.

```

1  enum Option<T> {
2      Some(T),
3      None,
4  }

```

Result is used to handle recoverable errors and represents success and failure. An error may not necessarily cause the program to crash, but it needs to be specifically handled to allow the program to continue executing. The definition of Result is as follows:

```
1 enum Result<T,E> {
2     Ok(T),
3     Err(E),
4 }
```

Opening a file that does not exist, accessing a file without permission, or attempting to convert a non-numeric string to a number will result in Err(E).

```
1 use std::fs::File;
2 use std::io::ErrorKind;
3
4 let f = File::open("kw.txt");
5 let f = match f {
6     Ok(file) => file,
7     Err(err) => match err.kind() {
8         ErrorKind::NotFound => match File::create("kw.txt"){
9             Ok(fc) => fc,
10            Err(e) => panic("Error while creating file!"),
11        }
12         ErrorKind::PermissionDenied => panic("No permission!"),
13         other => panic!("Error while opening file"),
14     }
15 }
```

In Rust, handling errors is crucial, and there are various mechanisms to achieve it. One of them is the panic mechanism, which should only be used when no other solution is available. When encountering an unrecoverable error, panic will immediately stop the program's execution to allow the programmer to identify the problem. This mechanism is also useful for cleaning up memory. However, if you do not want to use panic to clean up when encountering such errors, you can use the abort mechanism to let the operating system handle it instead.

Although error handling in Rust may look verbose when using match, there are more concise options available. For example, you can use the unwrap or expect methods to handle errors.

```
1 use std::fs::File;
2 use std::io;
3 use std::io::Read;
4
5 fn main() {
6     let f = File::open("kew.txt").unwrap();
7     let f = File::open("mon.txt").expect("Open file failed!");
8     // expect gives more specific error message than unwrap
9 }
```

If you only want to propagate the error and not handle it, you can use the "?" operator. In this case, the return type is Result, which returns String on success and io::Error on failure.

```
1 fn main() {
2     let s = read_from_file();
3     match s {
4         Err(e) => println!("{}", e),
5         Ok(s) => println!("{}", s),
6     }
7 }
```

```

7  }
8
9  fn read_from_file() -> Result<String, io::Error> {
10     let f = File::open("kew.txt"?); // if error, throw it
11     let mut s = String::new();
12     f.read_to_string(&mut s);
13
14     Ok(s)
15 }

```

When working with errors in Rust, you have to choose between using `Option` or `Result` depending on the type of failure or error you are handling. `Option` is used for handling possible failures and represents success or failure with `Some` or `None`. On the other hand, `Result` is used for handling recoverable errors and represents success and failure with `Ok` or `Err`. It is essential to handle errors appropriately to avoid program crashes. Finally, note that `Option` and `Result` are very similar, as `Option<T>` can be seen as `Result<T, ()>`.

1.4.12 Macro

Rust does not have built-in library functions, meaning that programmers must define everything themselves. However, Rust offers a set of powerful macros, including declarative and procedural macros, that can accomplish many tasks. Unlike C macros, Rust macros are more powerful and widely used. For instance, using the `"derive"` macro, new functionality can be added to structures, and commonly used methods like `"println!"`, `"vec!"`, and `"panic!"` are also macros.

There are two main categories of macros in Rust: declarative macros declared using `"macro_rules!"` and procedural macros, further divided into custom derive macros, function-like macros, and attribute macros. We've already covered declarative and custom derive macros, so this section will still focus on those. Readers can refer to related materials for other macro types.

The format of a declarative macro is `"macro_name!()"`, `"macro_name![]"`, or `"macro_name!"`. These macros all use a macro name followed by an exclamation mark, which is of one of these bracket types `()`, `[]`, `{}`. Different macro purposes require different brackets, with `"vec![]"` not being the same as `"vec!()"`. The latter looks more like a function and is the reason `"println!()"` uses parentheses. However, any of these brackets can be used and different brackets are just to satisfy the unity of meaning and form.

```

1  macro_rules! macro_name {
2      ($matcher) => {
3          $code_body;
4          return_value
5      };
6  }

```

In the macro definition, `"$matcher"` marks syntax elements like empty, identifiers, literals, keywords, symbols, patterns, and regular expressions. The `"$"` symbol captures values, which are then used in `"$code_body"` to process and potentially return a value. For example, the following macro can be used to calculate the left and right child nodes of a binary tree parent node `"p"`, with the left and right child node indices being `"2p"` and `"2p+1"`, respectively.

```

1  macro_rules! left_child {
2      ($parent:ident) => {
3          $parent << 1
4      };
5  }
6  macro_rules! right_child {

```

```

7      ($parent:ident) => {
8          ($parent << 1) + 1
9      };
10 }
```

When calculating left and right child nodes, macros such as `"left_child!(p)"` and `"right_child!(p)"` can be used instead of expressions like `"2 * p"` and `"2 * p + 1"`. This simplifies the code and clarifies its meaning. The `"indent"` and colon in the macro are metavariable specifiers. `"indent"` is an attribute of `"parent"`, indicating that `"parent"` is a value. Rust's macros rely on these specifiers to function correctly.

The second form of macros is called procedural macros because they are more like a process or function. They take code as input and produce other code as output. Derive macros, a type of procedural macro, add new functionality to code directly. The following example uses the `"derive"` macro to add the `"Clone"` method to `"Student"`, so that `"Student"` can directly call `"clone()"` to achieve copying. This is a simplified writing of `"impl Clone for Student"`.

```

1 #[derive(Clone)]
2 struct Student;
```

Multiple traits can also be placed in `"derive"` simultaneously, allowing for implementation of various functions.

```

1 #[derive(Debug, Clone, Copy)]
2 struct Student;
```

Macros are a complex topic and should only be used when necessary and when they simplify the code. It's not recommended to use macros for their own sake.

1.4.13 Code Organization and Dependency

In Rust, there are several concepts to understand for organizing code, including packages, crates, libraries, and modules. When using the `"cargo new"` command, Rust generates a package that contains multiple directories, each of which is a crate. After compilation, a crate can either be a binary executable or a library that can be used by other functions. Within a crate, there are typically multiple `.rs` files, which are called modules and are accessed using the `"use"` keyword. The table below provides a detailed overview of these concepts and their corresponding terms in Rust.

package	--> crates (dirs)	a package have multiple crates
crate	--> modules (lib/EFL)	a crate have multiple modules which can be compiled into a library or executable file
module	--> file.rs (file)	a module can contain one or more .rs files
package	<-- package name	
-- Cargo.toml		
-- src	<-- crate	
-- main.rs	<-- main module	
-- lib.rs	<-- module, library module	
-- math	<-- module, math module	
- mod.rs	<-- module, import add/sub from modules	
- add.rs	<-- module, implement add of math module	
- sub.rs	<-- module, implement sub of math module	
-- file	<-- crate	
-- core	<-- module, file operation module	
-- clear	<-- module, clear module	

To organize a large project, it is recommended to follow this approach when using Rust libraries. The **Tikv** codebase is a good example to follow.

Rust also offers many standard libraries for solving general programming tasks. By studying these libraries, developers can deepen their understanding of Rust and find inspiration for solving real-world problems. The standard libraries in Rust include:

alloc	env	i64	pin	task
any	error	i128	prelude	thread
array	f32	io	primitive	time
ascii	f64	isize	process	u8
borrow	ffi	iter	raw	u16
boxed	fmt	marker	mem	u32
cell	fs	net	ptr	u64
char	future	num	rc	u128
clone	hash	ops	result	usize
cmp	hint	option	slice	vec
collections	i8	os	str	backtrace
convert	i16	panic	string	intrinsics
default	i32	path	sync	lazy

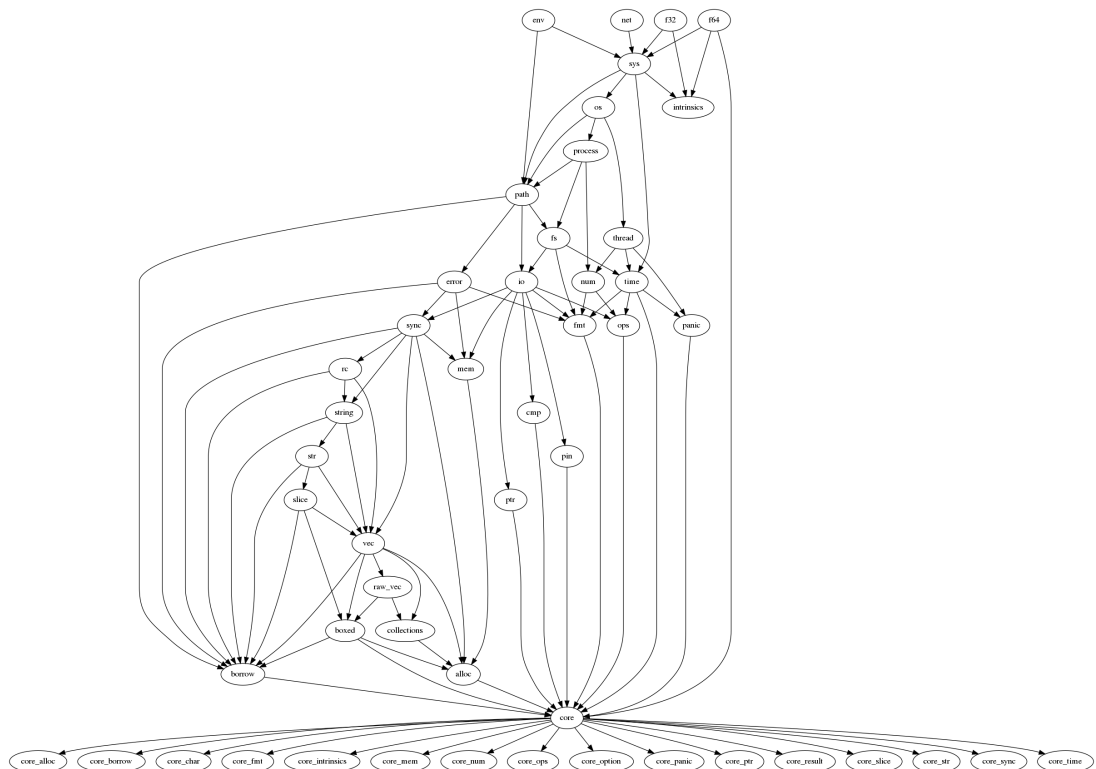


Figure 1.1: Rust package dependency

The image shown here is a simplified dependency graph of Rust libraries. It was created to illustrate how Rust as a language is built and how the language's libraries are interrelated. These libraries are considered the de facto standards of Rust, with some being fundamental and serving as dependencies for other libraries.

An analysis of Rust’s standard libraries revealed that they can be divided into three layers: core, alloc, and std. Each layer builds upon the previous one, with std being the top layer, alloc in the middle, and core serving as the most critical layer. Core is responsible for defining and implementing various core concepts in Rust’s basic syntax, such as variables, numbers, characters, and Boolean types. These are the foundational concepts that beginners learn when studying Rust’s basics.

1.4.14 Project: a password generator

In the previous sections, we covered many basic Rust concepts. Now, let’s apply them to a comprehensive project. In this project, readers will learn how to organize Rust project code, import modules, use various annotations, write tests, and build command-line tools.

In the age of the internet, people have numerous accounts and passwords to remember. Unfortunately, it’s challenging to remember unique passwords for each account, and people often set similar or identical passwords, leaving themselves vulnerable to hacking. Although password management software can help, relying on someone else’s software to manage your own passwords can make you uneasy. To solve this problem, the author created PasswdGenerator, a command-line tool that generates passwords that are at least 16 characters long and appear like random strings, such as 9KbAM4QWMCcyXAar.

PasswdGenerator can generate complex passwords of default length 16 by controlling the seed parameter, and the length parameter can adjust password length. This tool makes generating different passwords for all accounts easy, allowing users to input a digit or character that only they know to generate complex passwords. Using PasswdGenerator, the author generated different passwords for all his accounts and, because he thought of the seed himself, he didn’t have to worry about forgetting it. PasswdGenerator has been used for several years, and the author doesn’t even know his passwords for WeChat, Alipay, Twitter, Youtube, Github accounts etc.

PasswdGenerator is a command-line tool, and its usage method is similar to that of command-line tools in Unix-like systems. Therefore, its first usage is “PasswdGenerator -h” or “PasswdGenerator -help”.

```
shieber@kew $ PasswdGenerator -h
PasswdGenerator 0.1.0
A simple password generator for any account

USAGE:
    PasswdGenerator [OPTIONS] --seed <SEED>

OPTIONS:
    -h, --help            Prints help information
    -l, --length <LENGTH> Length of password [default: 16]
    -s, --seed <SEED>     Seed to generate password
    -V, --version          Prints version information
```

The USAGE section provides instructions on how to use the tool, while the OPTIONS section displays the control parameters.

```
shieber@kew $ PasswdGenerator -s wechat
wechat: GQnaoXobRwrgW21A
```

The output of the PasswdGenerator tool begins with a short seed, such as “wechat,” which can be customized with prefixes or suffixes to associate it with a specific account. The seed serves to indicate that the password was generated specifically for that account.

```
shieber@kew $ PasswdGenerator -s wechat -l 20
wechat: GQnaoXobRwrgW21Ac2Pb
```

The length of the password is controlled by the "-l" parameter. If this parameter is not included, the password defaults to a length of 16, which is sufficient for all accounts.

To handle command-line arguments, the clap library can be used, as shown below.

```
use clap::Parser;

/// A simple password generator for any account
#[derive(Parser, Debug)]
#[clap(version, about, long_about= None)]
struct Args {
    /// Seed to generate password
    #[clap(short, long)]
    seed: String,

    /// Length of password
    #[clap(short, long, default_value_t = 16)]
    length: usize,
}
```

The "seed" and "length" are the two command-line arguments, including their short and long forms, with "length" having a default value of 16. The command-line arguments can be obtained by calling `Args::parse()`.

Once the command-line arguments are obtained, the next step is to generate the password. My approach was to use a combination of a hash algorithm and base64 encoding. While there are many types of hash algorithms and various libraries available for generating hashes, it is practical to write a hash algorithm oneself. Here, I chose to use Mersenne primes to generate hash values.

To generate the hash value, the ASCII value of each character in the seed is multiplied by its corresponding index value, and the result is taken modulo the Mersenne prime 127. The remainder is then raised to the third power to obtain the hash value of the seed.

Mersenne primes are prime numbers of the form $M_n = 2^n - 1$, where n is also a prime number. While $2^n - 1$ may not always be a prime number, as in the case of primes such as 11, 23, and 29, Mersenne primes have the property that both n and M_n must be prime, making them suitable for calculating the hash value here. Some examples of Mersenne primes include 3, 7, 31, and 127 are shown below, in which 127 is the fourth Mersenne prime.

No	n	Mn
1	2	3
2	3	7
3	5	31
4	7	127
5	13	8191
6	17	131071
7	19	524287
8	31	2147483647
9	61	2305843009213693951
10	89	618970019642690137449562111

One method to convert a hash value to a string is by character replacement. The hash value is first modulo-ed with a fixed-length string (password), which results in a remainder used as an index to obtain a character. Repeating this process cyclically generates a string of characters that can be concatenated to form the password. For more complexity, the seed can be split and concatenated with the generated password.

```
// combine seed and passwd

let interval = passwd.clone();
for c in seed.chars() {
    passwd.push(c);
    passwd += &interval;
}
```

To make the password suitable for use, any unsuitable characters in the password string can be converted to Base64 encoding, which uses 64 visible characters. Additionally, "+" and "/" in Base64 can be replaced with "*" in the password.

```
// encode passwd (base64)
passwd = encode(passwd);
passwd = passwd.replace("+", "*").replace("/", "*");
```

If the password is still not long enough, it can be cyclically written into itself and wrapped with the seed and password using format, then returned.

```
let interval = passwd.clone();
while passwd.len() < length {
    passwd += &interval;
}

format!("{}", seed, &passwd[..length])
```

The tool uses three functions: "hash" crate to calculate the Mersenne hash value, "encryptor" crate to generate the password, and the main module to obtain command-line arguments. The main module calls the "generate_password" function from "encryptor" crate, which in turn calls "mersenne_hash" function from "hash" crate to calculate the hash value.

To organize the code, we can use Cargo's workspace mechanism to generate corresponding crates by specifying their names. The first step is to create and enter a directory.

```
shieber@kew $ mkdir PasswdGenerator && cd PasswdGenerator
```

Then, we need to write a "Cargo.toml" file that defines the names of the crates we want to generate.

```
1 [workspace]
2 members = [
3     "hash",
4     "encryptor",
5     "main",
6 ]
```

After that, we can use "cargo" to generate the three crates. Note that different parameters should be used to generate the library and the main program.

```
shieber@kew $ cargo new hash --lib
    Created library hash package
shieber@kew $ cargo new encryptor --lib
    Created library encryptor package
shieber@kew $ cargo new main
    Created binary (application) main package
```


Once generated, the directory structure should look like the following table.

```

.
|-- Cargo.toml
|-- encryptor
|   |-- Cargo.toml
|   |-- src
|       |-- lib.rs
|-- hash
|   |-- Cargo.toml
|   |-- src
|       |-- lib.rs
|-- main
|   |-- Cargo.toml
|   |-- src
|       |-- main.rs

```

6 directories, 7 files

In each crate, the modules need to be exported for use by the main program. For example, in the "hash" library, we can implement the "mersenne_hash" function in "merhash.rs" and export it in "lib.rs". Similarly, in the "encryptor" library, we can implement the function for generating passwords in "password.rs" and export it in "lib.rs". The "lib.rs" files of each crate should only be used to export functions. For mersenne_hash, it can be encapsulated in merhash.rs, while the function for generating passwords can be encapsulated in password.rs. The final directory looks like this:

```

|-- Cargo.toml
|-- encryptor
|   |-- Cargo.toml
|   |-- src
|       |-- lib.rs
|       |-- password.rs
|-- hash
|   |-- Cargo.toml
|   |-- src
|       |-- lib.rs
|       |-- merhash.rs
|-- main
|   |-- Cargo.toml
|   |-- src
|       |-- main.rs

```

6 directories, 9 files

It is important to note that the "hash" library does not depend on the external libraries and can be implemented first.

```

1 // hash/src/lib.rs
2
3 pub mod merhash; // export module merhash
4
5 #[cfg(test)] // test module
6 mod tests {

```

```

7     use crate::merhash::mersenne_hash;
8
9     #[test]
10    fn mersenne_hash_works() {
11        let seed = String::from("jdxjp");
12        let hash = mersenne_hash(&seed);
13        assert_eq!(2000375, hash);
14    }
15 }

```

Then, we implement merhash.

```

1  // hash/src/merhash.rs
2
3  ///! Mersenne Hash
4  ///!
5  ///! calculate hash value with mersenne prime 127
6  ///!
7  ///! # Example
8  ///! use hash::merhash::mersenne_hash;
9  ///!
10 ///! let seed = "jdxjp";
11 ///! let hash = mersenne_hash(&seed);
12 ///! assert_eq!(2000375, hash);
13 pub fn mersenne_hash(seed: &str) -> usize {
14     let mut hash: usize = 0;
15
16     for (i, c) in seed.chars().enumerate() {
17         hash += (i + 1) * (c as usize);
18     }
19
20     (hash % 127).pow(3) - 1
21 }

```

To ensure the code is well-documented and easy to maintain, document comments and module comments are used throughout the project. Tests are also included in the document comments, which is very convenient for testing and subsequent document maintenance.

To complete the "encryptor" library, we need to rely only on the external "base64" library. In addition, to handle error situations, the "anyhow" library is also introduced. To do this, we open the "Cargo.toml" file under the "encryptor" directory and add the necessary dependencies under the "[dependencies]" section, including the "anyhow" and "base64" libraries, as well as the "hash" library that was previously written.

```

1  [package]
2  name = "encryptor"
3  authors = ["shieber"]
4  version = "0.1.0"
5  edition = "2021"
6
7  [dependencies]
8  anyhow = "1.0.56"
9  base64 = "0.13.0"
10 hash = { path = "../hash" }

```

The implementation of the "password.rs" module is shown below.

```

1 // encryptor/src/lib.rs
2
3 pub mod password; // export module password
4
5 // encryptor/src/password.rs
6 use anyhow::{bail, Error, Result};
7 use base64::encode;
8 use hash::merhash::mersenne_hash;
9
10 /// crypto(length=100), you can exchange/add/delete characters
11 const CRYPTO: &str = "!pqHr$*+ST1Vst_uv:?wWS%X&Y-/Z01_2.34<AB\
12 9EC0|x#yDE^F{GHEI[]JK>LM#NOBWPQ:RaKU@}cde56R7=8f/9gIhi,jkzmn";
13
14 /// Password generator with hash value, use hash value's high
15 /// power to get character from CRYPTO
16 ///
17 /// #Example
18 /// use encryptor::password::generate_password;
19 /// let seed = "jdwnp";
20 /// let length = 16;
21 /// let passwd = generate_password(seed, length);
22 /// match passwd {
23 ///     Ok(val) => println!("{:?}", val),
24 ///     Err(err) => println!("{:?}", err),
25 /// }
26 pub fn generate_password(seed: &str, length: usize)
27     -> Result<String, Error>
28 {
29     // check password length
30     if length < 6 {
31         bail!("length must >= 6"); // return error
32     }
33
34     // calculate mer_hash
35     let p = match length {
36         6..=10 => 1,
37         11..=15 => 2,
38         16..=20 => 3,
39         _ => 3,
40     };
41     let mut mer_hash = mersenne_hash(seed).pow(p);
42
43     // calculate password by mer_hash
44     let mut passwd = String::new();
45     let crypto_len = CRYPTO.len();
46     while mer_hash > 9 {
47         let loc = mer_hash % crypto_len;
48         let nthc = CRYPTO.chars()
49             .nth(loc)
50             .expect("Error while getting char!");

```

```

51         passwd.push(nthc);
52         mer_hash /= crypto_len;
53     }
54
55     // combine seed and passwd
56     let interval = passwd.clone();
57     for c in seed.chars() {
58         passwd.push(c);
59         passwd += &interval;
60     }
61
62     // encode passwd to base64
63     passwd = encode(passwd);
64     passwd = passwd.replace("+", "*").replace("/", "*");
65
66     // length is not enough, use interval to fill
67     let interval = passwd.clone();
68     while passwd.len() < length {
69         passwd += &interval;
70     }
71
72     // return first length characters as password
73     Ok(format!("{}", seed, &passwd[..length]))
74 }

```

In the "Cargo.toml" of the main module, we introduce the necessary libraries, and note that the name is "PasswdGenerator".

```

1 [package]
2 name = "PasswdGenerator"
3 authors = ["shieber"]
4 version = "0.1.0"
5 edition = "2021"
6
7 [dependencies]
8 anyhow = "1.0.56"
9 clap = { version = "3.1.6", features = ["derive"] }
10 encryptor = { path = "../encryptor" }

```

Finally, in the main function, we call the "encryptor" and command line structure to generate passwords and return them.

```

1 // passwdgenerate/src/main.rs
2
3 use anyhow::{bail, Result};
4 use clap::Parser;
5 use encryptor::password::generate_password;
6
7 /// A simple password generator for any account
8 #[derive(Parser, Debug)]
9 #[clap(version, about, long_about = None)]
10 struct Args {
11     /// Seed to generate password

```

```

12     #[clap(short, long)]
13     seed: String,
14
15     /// Length of password
16     #[clap(short, long, default_value_t = 16)]
17     length: usize,
18 }
19
20 fn main() -> Result<()> {
21     let args = Args::parse();
22
23     // seed must be longer than 4 characters
24     if args.seed.len() < 4 {
25         bail!("seed {} length must >= 4", &args.seed);
26     }
27
28     let (seed, length) = (args.seed, args.length);
29     let passwd = generate_password(&seed[..], length);
30     match passwd {
31         Ok(val) => println!("{}", val),
32         Err(err) => println!("{}", err),
33     }
34
35     Ok(())
36 }

```

Once the program is complete, compile it with "cargo build --release" to get the "PasswdGenerator" command line tool in the "target/release/" directory. You can put it in the "/usr/local/bin" directory so that you can use it anywhere on the system. Of course, using "cargo doc" can also generate a "doc" directory, which contains detailed documentation about the project.

Through this small project, readers have become familiar with Rust code organization, module usage, testing methods, and coding styles. Although this password generator is simple, it is completely sufficient for personal use.

1.5 Summary

In this chapter, we provide an overview of Rust fundamentals. Firstly, we begin by introducing the installation of the Rust toolchain and summarizing some learning resources. We then review Rust basics, covering variables, functions, ownership, lifetimes, generics, traits, smart pointers, error handling, and macro systems.

After that, we present a comprehensive project that implements a command-line tool using Rust. Although the content of this chapter is relatively basic, it is essential to understand these concepts to build more complex projects in Rust. Readers who need further clarification are recommended to refer to the Rust documentation.

By the end of this chapter, readers will have a general understanding of Rust. In the following chapters, we will delve into learning data structures and algorithms.

Chapter 2

Computer Science

2.1 Objectives

- Understand the ideas of computer science.
- Understand the concept of abstract data types.
- Review the basics of the Rust programming language.

2.2 Getting Started

In the realm of computer technology, acquiring a solid understanding of the fundamentals is crucial to effectively solving problems. However, even with this foundation, tackling some problems can still be challenging. The complexity of a problem and its corresponding solution can often obscure the key ideas needed to reach a resolution. Additionally, since problems can have multiple solutions, each with its own constraints and logic, trying to combine the structure of one problem with the solution logic of another can lead to trouble.

This chapter focuses on computer science, algorithms, and data structures, specifically examining why studying these topics is essential. By understanding the structure and logic of problem statements, we can better approach solving problems.

2.3 What is Computer Science?

Computer science is a challenging field to define, possibly due to the word "computer" in its name. However, computer science is not limited to the study of computers alone, as it encompasses the study of problems, solutions, and the process of generating solutions. The objective of computer scientists is to create an algorithm, a set of instructions, to solve a given problem instance. Following this algorithm allows similar problems to be solved in a finite amount of time. Although computer science can be regarded as the study of algorithms, it is important to recognize that some problems do not have algorithms to solve them, such as NP-complete problems^[9]. Although these problems cannot currently be solved, researching them is essential because solving them could result in technological advancements. Like the "Goldbach Conjecture^[10]", research into it has developed many tools. Therefore, computer science can be defined as the study of solvable problem solutions and unsolvable problem ideas.

When describing problems and solutions, a computable problem is one that has an algorithm to solve it. In contrast, computer science can be defined as "the study of whether there is a solution to problems that are computable and non-computable." It is noteworthy that this definition does not mention the word "computer." Solutions are independent of the machine and consist of ideas, regardless of whether a computer is employed.

Computer science focuses on the art of problem-solving, which is achieved through abstraction. Abstraction allows us to consider problems and their solutions from a logical perspective, rather than being bogged down by physical details. An everyday example of this is driving a car. As a driver, you interact with the car to reach your destination, using functions provided by the car’s designer, such as the key, engine, gears, brakes, accelerator, and steering wheel. These functions are interfaces that simplify the driving experience. In contrast, a car mechanic has a different perspective. They understand the inner workings of the car, including the engine, transmission, temperature control, and windshield wipers. These details operate at the physical level and happen “under the hood.”

Similarly, users interact with computers from a logical perspective, using programs such as email, document editors, and media players. In contrast, computer scientists, programmers, and system administrators work at a lower level, understanding how the operating system works, how to configure network protocols, and how to write control function scripts. The key concept behind both of these examples is abstraction. By hiding the complex details of the system, an interface simplifies the user’s interaction with the computer. Rust’s mathematical calculation functions are a good example of this.

```
1 // sin_cos_function.rs
2
3 fn main() {
4     let x: f32 = 1.0;
5     let y = x.sin();
6     let z = x.cos();
7     println!("sin(1) = {y}, cos(1) = {z}");
8     // sin(1) = 0.84147096, cos(1) = 0.5403023
9 }
```

We may not know how to calculate the sine and cosine values, but as long as we understand how to use the function, that’s all we need. Someone has implemented the algorithm for calculating sine and cosine, and we can trust that the function will return the correct result. This is the idea of a “black box.” The interface describes the function’s input and output, while the details of the algorithm are hidden within.

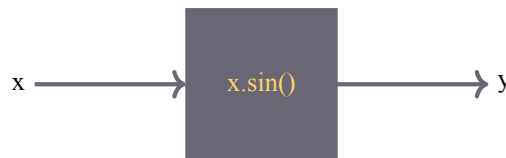


Figure 2.1: calculate the sine value

2.4 What is Programming?

Programming is the process of converting algorithms into computer instructions. An algorithm is a set of steps that produces a solution based on data and expected results. It is essential to have a solution before writing a program. Although computer science is not solely about programming, programming is a crucial skill for computer scientists. When programming, one organizes the problem statement structure in their mind and expresses it clearly for efficient computer processing.

To represent processes and data, programming languages must provide control methods and various data types. Control methods allow for algorithms to be expressed clearly and unambiguously, with sequential processing, selection, and iterative repetition being the minimum requirements.

All data items in a computer are represented in binary form, and data types provide an interpretation and presentation of this binary data. Data types are abstractions of the physical world used to represent

entities in the problem, with underlying primitive data types such as integer and floating-point data types being the foundation for algorithm development.

The operations that can be performed on data are also described in the data type. For example, the most fundamental operations for numbers are addition, subtraction, multiplication, and division. However, programming languages' simple structures and data types may not always suffice to represent complex solutions. To control this complexity, more reasonable data management methods (data structures) and operation processes (algorithms) are required.

2.5 Why Study Data Structures and Abstract Data Types?

Computer scientists use abstraction to manage the complexity of problems and obtain concrete steps to solve them. Abstraction allows them to focus on the big picture without getting lost in the details. Creating a domain model enables computers to solve problems more effectively, and algorithms can be described in a more consistent manner.

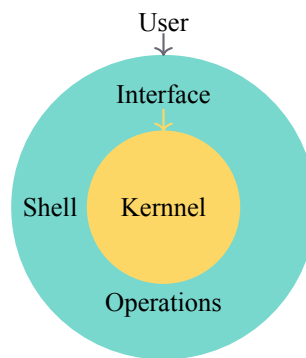


Figure 2.2: system level diagram

An Abstract Data Type (ADT) is a logical description of how to view and operate on data. It provides a level of abstraction where only the meaning of the data is of concern, not its final storage form. By wrapping the data in this way, implementation details can be hidden. The user's interaction with the interface is an abstract operation, while the user and shell are abstract data types. Although the specific operations are not known, understanding their interaction mechanism is still possible. This is the benefit of abstract data types for algorithm design.

To implement abstract data types, primitive data types are used to build new data types from a physical view, known as data structures. There are usually many different ways to implement abstract data types, but different implementations should have the same physical view. This allows programmers to change implementation details without altering the interaction method, and users to continue to focus on the problem.

The common logic for abstract data types includes creating new data types, retrieving data, adding, deleting, searching, modifying, checking for empty data, and computing size. For example, to implement a queue, its abstract data type logic should include at least: `new()`, `is_empty()`, `len()`, `clear()`, `enqueue()`, and `dequeue()`. Once the specific operational logic is known, implementation becomes simple, and there are various ways to implement it as long as these abstract logic exists.

2.6 Why Study Algorithms?

In the computer world, the goal is to use algorithms that are efficient in terms of speed and memory usage. However, some problems may be difficult to handle, and no algorithm can provide an answer within an expected time frame. In such cases, it is important to distinguish between problems that have

solutions and those that do not, as well as problems that have solutions but require significant time or resources. As computer scientists, we need to evaluate and compare different solutions to determine the most appropriate one.

Learning from the problem-solving approaches of others is an efficient way to develop our skills. By examining different solutions, we can gain insights into how various algorithm designs help us tackle challenging problems. By studying various algorithms, we can uncover their core principles and develop a universal algorithm that can be applied to similar problems in the future. However, for the same problem, different people may provide different algorithm implementations. As we saw earlier in the example of computing sine and cosine, there can be many different implementation versions. If one algorithm can achieve the same result as another algorithm in less time, it is obviously the better option.

2.7 Summary

This chapter delves into the concepts of computer science thinking and abstract data types, providing clear definitions and highlighting the roles of algorithms and data structures. By abstracting data types, specific implementation and operation logic is removed, which clarifies the boundaries between data structures and algorithms and significantly simplifies the process of algorithm design. In the forthcoming chapters, we will continue to utilize abstract data types to design various data structures.

Chapter 3

Algorithm Analysis

3.1 Objectives

- Understanding the Importance of Algorithm Analysis.
- Learning to perform performance benchmark tests on Rust programs.
- Being able to use the Big O notation to analyze the complexity of algorithms.
- Understanding the Big O analysis results for Rust data structures such as arrays.
- Understanding how Rust data structure implementations affect algorithm analysis.

3.2 What is Algorithm Analysis?

In Chapter 2, we defined an algorithm as a set of instructions that solves a specific problem by producing an expected result given a specific input. A program is an algorithm written in a programming language. As programmers use different programming languages and have varying levels of knowledge, different programs may describe the same algorithm.

New computer science students may compare their programs with others. To explore the differences between programs that solve the same problem, let's consider the functions "sum_of_n" and "tik_tok," which both calculate the sum of the first n integers. The first function uses an accumulator variable initialized to 0 and iterates over the n integers, adding each value to the accumulator. The "tik_tok" function does the same thing but uses unclear identifier names and an unnecessary assignment.

```
1 // sum_of_n.rs
2 fn sum_of_n(n: i32) -> i32 {
3     let mut sum: i32 = 0;
4     for i in 1..=n { sum += i; }
5     sum
6 }
7
8 // tiktok.rs
9 fn tiktok(tik: i32) -> i32 {
10    let mut tok = 0;
11    for k in 1..=tik {
12        let ggg = k;
13        tok = tok + ggg;
14    }
15    tok
16 }
```

When evaluating the two functions, the answer depends on the reader's criteria. If readability is the concern, then "sum_of_n" is better. However, in this book, we focus more on the statements of the algorithms themselves, and clean coding is not part of the discussion.

Algorithm analysis compares the amount of resources used by algorithms. An algorithm is better than another algorithm if it uses fewer resources or is more efficient in using resources. When comparing the "sum_of_n" and "tik_tok" functions, they appear very similar in terms of resource usage. Identifying the resources used for computation is crucial and requires considering both time and space perspectives.

- The space used by an algorithm is determined by its memory consumption, which is typically influenced by the size and nature of the problem. However, some algorithms have special space requirements.

- Algorithm execution time refers to the time it takes for an algorithm to execute all its steps.

For instance, the execution time of the "sum_of_n" function can be analyzed through benchmark testing. In Rust, the execution time of code can be calculated by recording the system time before and after function execution. The `SystemTime` function, which is available in `std::time`, returns the system time when called and gives the elapsed time afterward. By calling this function at the beginning and end of function execution, we can obtain the function execution time.

```

1 // static_func_call.rs
2
3 use std::time::SystemTime;
4
5 fn sum_of_n(n: i64) -> i64 {
6     let mut sum: i64 = 0;
7     for i in 1..=n {
8         sum += i;
9     }
10    sum
11 }
12
13 fn main() {
14     for _i in 0..5 {
15         let now = SystemTime::now();
16         let _sum = sum_of_n(500000);
17         let duration = now.elapsed().unwrap();
18         let time = duration.as_millis();
19         println!("func used {time} ms");
20     }
21 }

```

To illustrate this, we executed the "sum_of_n" function five times, each time calculating the sum of the first 500,000 integers and 1000,000 integers. The results were as follows:

```

func used 10 ms
func used 6 ms
func used 6 ms
func used 6 ms
func used 6 ms

```

```

func used 17 ms
func used 12 ms
func used 12 ms
func used 12 ms
func used 12 ms

```

After comparing the results of two executions, it was found that the time taken by the function to execute is basically the same. On average, the function takes 6 milliseconds to execute. The first execution takes longer at 10 milliseconds because the function requires initialization and preparation. The subsequent four executions do not require initialization, and their times are more accurate, hence why multiple executions are necessary. When calculating the sum of the first 1,000,000 integers, it can be observed that the first execution takes longer, and the subsequent executions take exactly twice the time taken to calculate the sum of the first 500,000 integers. This indicates that the execution time of the algorithm is proportional to the size of the calculation.

To further explore this idea, consider the following function, which also calculates the sum of the first n integers, but uses the mathematical formula $\sum_{i=0}^n = \frac{n(n+1)}{2}$ to compute it.

```
1 // static_func_call2.rs
2 fn sum_of_n2(n: i64) -> i64 {
3     n * (n + 1) / 2
4 }
```

The `sum_of_n` function in `static_func_call.rs` is modified to use this formula, and a benchmark test is performed using three different values of n (100,000, 500,000, 1,000,000). Each calculation is repeated five times, and the average execution time is recorded. The results are as follows:

```
func used 1396 ns
func used 1313 ns
func used 1341 ns
```

There are two important points to note in the output. First, the recorded execution time is in nanoseconds, which is significantly shorter than the previous examples. The execution times for all three calculations are approximately 0.0013 milliseconds, which is orders of magnitude shorter than the previous 6 milliseconds. Second, the execution time is independent of n . As n increases, the calculation time remains the same, indicating that the calculation is almost unaffected by n .

This benchmark test indicates that the iterative solution `sum_of_n` performs more work because some program steps are repeated, which is why it takes longer. Moreover, the execution time of the iterative solution increases with n . Additionally, it should be noted that running similar functions on different computers or using different programming languages may yield different results. The calculated value of 1341 nanoseconds mentioned in the text refers to the computer used in this study, which is Lenovo Legion R7000P with a 16-core AMD R7-4800H CPU. Older computers may take longer to execute `sum_of_n2`.

It is necessary to find a better way to describe the execution time of algorithms. Although benchmark tests measure the actual time taken by a program to execute, they do not provide a useful measure because the execution time depends on the specific machine, and there are also magnitude conversions between milliseconds and nanoseconds. We require a measure that is independent of the program or computer used, and can be used to compare the efficiency of different algorithm implementations. Big O notation is a good method for measuring algorithmic efficiency in the field of algorithm analysis, analogous to acceleration measuring the change in speed per second.

3.3 Big-O Notation Analysis

To evaluate the performance of an algorithm, we need to quantify the number of operation steps and storage space required independently of any specific program or computer. For the summing algorithm, counting the execution statements is a good measure. In the function `sum_of_n`, there is one assignment statement (the `_sum = 0`), and n addition statements (the `_sum += i`), making the statement count for the function equal to $n + 1$. The storage space used is two variables, n and `sum`, so the count is 2.

We use the function T to represent the total number of executions to measure the algorithm's time complexity, where $T(n) = 1 + n$. The parameter n represents the problem size, and $T(n)$ is the time spent solving a problem with a size greater than or equal to n . Using n to represent the problem size makes sense for the summing function, as summing 100,000 integers is a larger problem than summing 1000 integers, and thus takes longer. Our goal is to demonstrate how the algorithm's execution time changes relative to the problem size.

For measuring the algorithm's space complexity, we use the function S to represent the total memory consumption, where $S(n) = 2$. The parameter n still represents the problem size, but $S(n)$ is independent of n . To analyze the performance of an algorithm, it is crucial to determine the primary components of its time complexity function $T(n)$ and space complexity function $S(n)$, rather than merely counting the number of operation steps and space usage. As the problem size grows, some components of these functions become dominant. The fastest-growing parts of these functions with n are represented by the big O notation, denoted as $O(f(n))$. The function $f(n)$ signifies the primary part of $T(n)$ or $S(n)$.

For instance, $T(n) = n + 1$ in the previous example. As n increases, the constant 1 becomes less significant. Thus, we can approximate the running time of $T(n)$ as $O(T(n)) = O(n + 1) = O(n)$. While the constant 1 is undoubtedly important for $T(n)$, its contribution becomes negligible when n is large. However, when $T(n) = n^3 + 1$, ($n = 1$), discarding the constant 1 is unreasonable since it would mean discarding a significant portion of the running time. In this case, using $O(T(n)) = O(n^3 + 1)$ is appropriate for small values of n . But as n increases, the 1 becomes less significant, and we can approximate $T(n)$ as $O(T(n)) = O(n^3)$.

For $S(n)$, since it is a constant, $O(S(n)) = O(2) = O(1)$. The big O notation only represents the order of magnitude. Therefore, although the actual complexity of $S(n)$ is $O(2)$, we use $O(1)$ instead.

Assuming an algorithm has a number of operational steps determined by $T(n) = 6n^2 + 37n + 996$. When n is small, such as 1 or 2, the constant term 996 may seem to be the main part of the function. However, as n grows larger, the significance of the n^2 term increases. In fact, as n becomes very large, the other two terms become negligible in the final result. Thus, we can approximate $T(n)$ by ignoring the other terms and focusing only on $6n^2$. Moreover, the coefficient 6 also becomes insignificant as n grows larger. Thus, we say that $T(n)$ has a complexity order of n^2 , or $O(n^2)$.

However, the complexity of some algorithms depends on the exact values of the data, rather than the size of the problem. For such algorithms, their performance needs to be characterized based on best case, worst case, or average case scenarios. Worst case refers to a specific dataset that leads to particularly poor algorithm performance, while the same algorithm may have significantly different performance under different datasets. In most cases, the efficiency of algorithm execution lies between the two extremes of worst and best cases, i.e., the average case. Thus, it is important for programmers to understand these differences and avoid being misled by a particular case.

In the study of algorithms, some common order functions appear repeatedly, as shown in the table and figure below. To determine which of these functions is the main part, we need to observe how they relate to each other as n grows larger.

Table 3.1: Complexity functions

$O(1)$	$O(\log n)$	$O(n)$	$O(n \log n)$	$O(n^2)$	$O(n^3)$	$O(2^n)$
constants	logarithmic	linear	linear logarithmic	quadratic	cubic	power exponential.

The figure below illustrates the growth of various functions. At small values of n , it can be difficult to discern which function dominates the others. However, as n increases, the relative sizes of the functions become clear. It is worth noting that at $n = 10$, 2^n exceeds n^3 , though this case is not shown on the graph. The figure also highlights the differences between different orders of magnitude. Generally, for $n > 10$, we have $O(2^n) > O(n^3) > O(n^2) > O(n \log n) > O(n) > O(\log n) > O(1)$. This information is useful in algorithm design, as it allows us to calculate the complexity of an algorithm. If we determine that an algorithm has a complexity similar to $O(2^n)$, for example, we know that it is not practical and can look to optimize it for better performance.

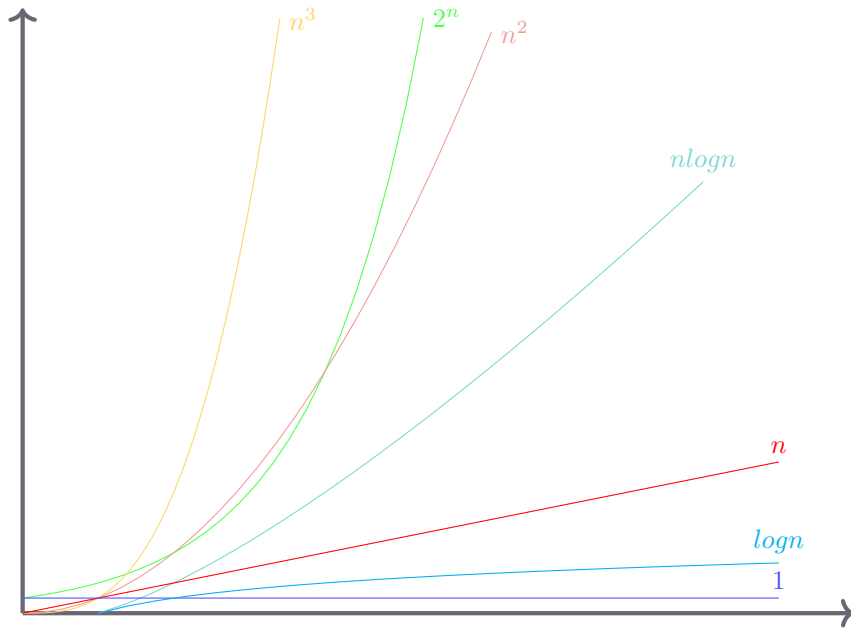


Figure 3.1: Complexity curve

The above analysis examines time and space complexity, but often time complexity is the primary concern because space is typically difficult to optimize. For example, the space complexity of an input array is limited from the outset, but different algorithms can result in significant differences in the time required to run on the array. Moreover, as storage becomes cheaper and more abundant, time becomes the most important factor because it cannot be increased. Therefore, most of the complexity analysis in this book focuses on time complexity, unless otherwise specified.

The following code only employs addition, subtraction, multiplication, and division. We can use it to apply our newly acquired understanding of algorithm complexity analysis.

```

1 // big_o_analysis.rs
2 fn main() {
3     let a = 1; let b = 2;
4     let c = 3; let n = 1000000;
5
6     for i in 0..n {
7         for j in 0..n {
8             let x = i * i;
9             let y = j * j;
10            let z = i * j;
11        }
12    }
13
14    for k in 0..n {
15        let w = a*b + 45;
16        let v = b*b;
17    }
18
19    let d = 996;
20 }
```

The code above has a time complexity of $O(n^2)$, which can be obtained by analyzing the code. Firstly, the time to allocate operations a , b , c , and n is a constant 4. Then, the second term is $3n^2$, which comes from the execution of three statements n^2 times due to nested iteration. The third term is $2n$, as two statements are iteratively executed n times. Finally, the fourth term is a constant 1, representing the final assignment statement $d = 996$. Thus, the function that determines the number of operational steps is $T(n) = 4 + 3n^2 + 2n + 1 = 3n^2 + 2n + 5$. By examining the exponent, we can see that the n^2 term is the most significant, indicating that the time complexity of this code is $O(n^2)$.

3.4 Anagram Detection

The problem of checking if two strings are permutations of each other serves as an example to demonstrate different levels of complexity. A permutation is defined as one string, $s1$, being a rearrangement of another string, $s2$. For instance, "heart" and "earth" are permutations of each other, as are "rust" and "trus". Let us assume that both strings have the same length and consist only of 26 lowercase letters. Our goal is to write a function that takes two strings as inputs and returns whether or not they are permutations of each other.

3.4.1 Brute Force

The most naive solution to this problem is to use brute force and check all possible permutations of $s1$. This method is highly resource-intensive, consuming both time and memory. When generating all possible strings of $s1$, there are n possibilities for the first position, $n-1$ for the second position, $n-2$ for the third position, and so on, resulting in $n! = n \times (n-1) \times (n-2) \times \dots \times 3 \times 2 \times 1$ total possibilities. Although some strings may be duplicates, the program cannot know this in advance, and thus it will generate $n!$ strings.

For example, if $s1$ has a length of 20, there will be $20! = 2,432,902,008,176,640,000$ possible strings. If we process one possible string per second, it would take 77,146,816,596 years to exhaust the entire list. In fact, $n!$ grows even faster than n^2 , making brute force an infeasible solution for any learning or real software project. Nonetheless, being aware of its existence and making an effort to avoid such situations is essential.

3.4.2 Checking Off

A more efficient solution to the problem involves checking if each character in the first string is present in the second string. If all characters are found, then the two strings are permutations of each other. We can use the `' '` character to replace a character once it has been checked.

```

1 // anagram_solution2.rs
2
3 fn anagram_solution2(s1: &str, s2: &str) -> bool {
4     if s1.len() != s2.len() {
5         return false;
6     }
7
8     // put the characters of s1 and s2 into
9     // vec_a and vec_b respectively
10    let mut vec_a = Vec::new();
11    let mut vec_b = Vec::new();
12    for c in s1.chars() {
13        vec_a.push(c);
14    }
15    for c in s2.chars() {

```

```

16         vec_b.push(c);
17     }
18
19     // index: pos1, pos2
20     let mut pos1: usize = 0;
21     let mut pos2: usize;
22
23     // loop controlling
24     let mut is_anagram = true;
25     let mut found: bool;
26     while pos1 < s1.len() && is_anagram {
27         pos2 = 0;
28         found = false;
29         while pos2 < vec_b.len() && !found {
30             if vec_a[pos1] == vec_b[pos2] {
31                 found = true;
32             } else {
33                 pos2 += 1;
34             }
35         }
36
37         // replace a character with ' '
38         if found {
39             vec_b[pos2] = ' ';
40         } else {
41             is_anagram = false;
42         }
43
44         // tackle the next character of s1
45         pos1 += 1;
46     }
47
48     is_anagram
49 }
50
51 fn main() {
52     let s1 = "rust";
53     let s2 = "trus";
54     let result = anagram_solution2(s1, s2);
55     println!("s1 and s2 is anagram: {result}");
56     // s1 and s2 is anagram: true
57 }

```

Upon analyzing this algorithm, we observe that each character in $s1$ is compared to all characters in $s2$, which leads to a maximum of n comparisons for each character in $s1$. As a result, the n positions in $blist$ will be visited once to match the characters from $s1$. The total number of visits can be expressed as the sum of integers from 1 to n .

$$1 + 2 + \cdots + n = \sum_{i=1}^n i = \frac{n(n+1)}{2} = \frac{1}{2}n^2 + \frac{1}{2}n \quad (3.1)$$

The time complexity of this algorithm is $O(n^2)$, as the term n^2 dominates as n grows. This is much better than the brute force algorithm that has a complexity of $O(n!)$.

3.4.3 Sort and Compare

An alternative solution for scrambled strings is to take advantage of the fact that even if `s1` and `s2` are different, they consist of exactly the same characters. Therefore, we can sort each string in alphabetical order from `a` to `z`, and if the two sorted strings are the same, then they are scrambled strings.

```
1 // anagram_solution3.rs
2
3 fn anagram_solution3(s1: &str, s2: &str) -> bool {
4     if s1.len() != s2.len() {
5         return false;
6     }
7
8     // put the characters of s1 and s2 into
9     // vec_a and vec_b respectively then sorting
10    let mut vec_a = Vec::new();
11    let mut vec_b = Vec::new();
12    for c in s1.chars() {
13        vec_a.push(c);
14    }
15    for c in s2.chars() {
16        vec_b.push(c);
17    }
18    vec_a.sort();
19    vec_b.sort();
20
21    // compare vec_a and vec_b, if there is any mismatch of
22    // any character then break the loop
23    let mut pos: usize = 0;
24    let mut is_anagram = true;
25    while pos < vec_a.len() && is_anagram {
26        if vec_a[pos] == vec_b[pos] {
27            pos += 1;
28        } else {
29            is_anagram = false;
30        }
31    }
32
33    is_anagram
34 }
35
36 fn main() {
37     let s1 = "rust";
38     let s2 = "trus";
39     let result = anagram_solution3(s1, s2);
40     println!("s1 and s2 is anagram: {result}");
41     // s1 and s2 is anagram: true
42 }
```

At first glance, the time complexity of the algorithm that checks for scrambled strings seems to be $O(n)$ because it has only one while loop. However, the complexity of calling the `sort()` function is not negligible, and it is usually $O(n^2)$ or $O(n \log n)$, which makes the algorithm's complexity of the same order of magnitude as sorting algorithms.

3.4.4 Count and Compare

The previous methods always create `vec_a` and `vec_b`, which can be memory-intensive when `s1` and `s2` are long. To solve this problem, we can use two lists with a length of 26 to count the frequency of each character in `s1` and `s2`. This approach involves multiple iterations, but the iterations are non-nested. Moreover, the third iteration that compares the two frequency lists only needs 26 iterations because there are only 26 lowercase letters. Thus, the time complexity of this algorithm is $O(n)$, where n is the length of `s1` or `s2`. This is a linear time complexity algorithm, and both its space and time complexities are relatively efficient.

```

1 // anagram_solution4.rs
2
3 fn anagram_solution4(s1: &str, s2: &str) -> bool {
4     if s1.len() != s2.len() { return false; }
5
6     // with the size of 26, c1 and c2 are storage
7     // for saving ASCII charaters
8     let mut c1 = [0; 26];
9     let mut c2 = [0; 26];
10    for c in s1.chars() {
11        let pos = (c as usize) - 97; // 97: ASCII value of a
12        c1[pos] += 1;
13    }
14    for c in s2.chars() {
15        let pos = (c as usize) - 97;
16        c2[pos] += 1;
17    }
18
19    // compare the value of ASCII
20    let mut pos = 0;
21    let mut is_anagram = true;
22    while pos < 26 && is_anagram {
23        if c1[pos] == c2[pos] {
24            pos += 1;
25        } else {
26            is_anagram = false;
27        }
28    }
29
30    is_anagram
31 }
32
33 fn main() {
34     let s1 = "rust";
35     let s2 = "trus";
36     let result = anagram_solution4(s1, s2);
37     println!("s1 and s2 is anagram: {result}");
38     // s1 and s2 is anagram: true
39 }

```

However, if `s1` and `s2` are short and do not use all 26 characters, this algorithm sacrifices some storage space. When deciding which algorithm to use, it is essential to make a trade-off between space and time and consider the real scenarios that your algorithm will face.

3.5 Performance of Rust Data Structures

3.5.1 Scalar and Complex Data Structures

In this section, we will explore the big O performance of Rust's basic data types. Understanding the performance of these data structures is crucial because they are the core building blocks in Rust, and more complex data structures are constructed from them. Rust's basic data types are divided into two major categories: scalar and composite types. Scalar types represent a single value, while composite types are combinations of scalar types.

Rust has four basic scalar types: integer, floating-point, boolean, and character. On the other hand, Rust has two composite types: tuple and array. Scalar types are the most basic and memory-coupled primitive types, with computational efficiency that is typically $O(1)$. Meanwhile, composite types are more complex, and their complexity varies with their data scale.

Let's take a look at some examples of using basic data types.

```
1 let a: i8 = -2;
2 let b: f32 = 2.34;
3 let c: bool = true;
4 let d: char = 'a';
5
6 // tuple with various types
7 let x: (i32, f64, u8) = (200, 5.32, 1);
8 let xi32 = x.0;
9 let xf64 = x.1;
10 let xu8 = x.2;
```

The tuple is a composite data structure that combines multiple values of various types, with a fixed length. Once declared, its length cannot be increased or decreased. Tuple indexing starts from 0, and its values can be directly accessed using the "." (dot notation). Arrays, on the other hand, must have the same type for each element, and once declared, their length cannot be increased or decreased.

```
1 let months = ["January", "February", "March", "April",
2               "May", "June", "July", "August", "September",
3               "October", "November", "December"];
4
5
6 let first_month = months[0]
7 let halfyear = &months[..6];
8
9 let mut monthsv = Vec::new();
10 for month in months { monthsv.push(month); }
```

Rust offers various data types, composed of scalar and composite types, such as Vec and HashMap. Vec is a collection type similar to an array, which allows for adding, shrinking, and limiting its length. It provides greater scalability and can be used wherever an array can be used.

3.5.2 Collection Data Structures

Rust's collection types are divided into linear and non-linear types, constructed based on scalar and composite types. The linear collection types include String, Vec, VecDeque, and LinkedList, while the non-linear collection types include HashMap, BTreeMap, HashSet, BTreeSet, and BinaryHeap. These collection types involve indexing, adding, and deleting operations, and their respective complexities are often $O(1)$, $O(n)$, etc.

Rust's String type is based on Vec, which allows String to be modified like Vec. To use a portion of the String, one can use &str, which is a slice of the String type that is easy to index. However, since &str is based on String, it cannot be changed because modifying the slice would change the data in String, which may be used elsewhere. In Rust, mutable strings use String, while immutable strings use &str. Vec is similar to lists in other languages, and VecDeque extends Vec to support inserting data at both ends of the sequence, making it a double-ended queue. LinkedList is a linked list and can be used when an unknown size Vec is desired.

The HashMap in Rust is similar to dictionaries in other languages, while BTreeMap is a B-tree whose nodes contain data and pointers. It is often used to implement databases, file systems, and other content storage applications. The HashSet and BTreeSet in Rust are similar to the Set collection in other languages, which are used to record a single value that has appeared once. HashSet is implemented using HashMap as its underlying data structure, while BTreeSet is implemented using BTreeMap. BinaryHeap is a priority queue, storing a set of elements that can extract the maximum value at any time.

Rust's collection data types are highly efficient, as can be seen from the tables below that show the performance of various data structures. The highest complexity of these data structures is $O(n)$.

Table 3.2: Time complexity of linear composite types

Type	get	insert	remove	append	split_off
Vec	$O(1)$	$O(n - i)$	$O(n - i)$	$O(m)$	$O(n - i)$
VecDeque	$O(1)$	$O(\min(i, n - i))$	$O(\min(i, n - i))$	$O(m)$	$O(\min(i, n - i))$
LinkedList	$O(\min(i, n - i))$	$O(\min(i, n - i))$	$O(\min(i, n - i))$	$O(1)$	$O(\min(i, n - i))$

Table 3.3: Time complexity of non-linear composite types

Type	get	insert	remove	predecessor	append
HashMap	$O(1)$	$O(1)$	$O(1)$	N/A	N/A
HashSet	$O(1)$	$O(1)$	$O(1)$	N/A	N/A
BTreeMap	$O(\log n)$	$O(\log n)$	$O(\log n)$	$O(\log n)$	$O(n + m)$
BTreeSet	$O(\log n)$	$O(\log n)$	$O(\log n)$	$O(\log n)$	$O(n + m)$
Type	push	pop	peek	peek_mut	append
BinaryHeap	$O(1)$	$O(\log n)$	$O(1)$	$O(1)$	$O(n + m)$

3.6 Summary

This chapter introduced the big O notation analysis method for evaluating algorithm complexity, which involves calculating the number of code execution steps and determining the maximum quantity level. We then examined the complexity of Rust's basic data types and collection data types. Upon comparison and analysis, it becomes clear that Rust's built-in scalar, composite, and collection data types are highly efficient. Additionally, custom data structures implemented based on these collection types can also achieve efficient.

Chapter 4

Basic Data Structures

4.1 Objectives

- Understanding Abstract Data Types Vec, Stack, Queue, Deque, Linked List
- Being able to implement stack, queue, deque, and linked list using Rust
- Understanding the performance (complexity) of basic linear data structures
- Understanding prefix, infix, and postfix expression formats
- Using a stack to implement postfix expression and calculate the value
- Using a stack to convert infix expressions to postfix expressions
- Being able to identify whether to use a stack, queue, deque, or linked list for a given problem
- Being able to implement abstract data types as linked lists using nodes and references
- Being able to compare the performance of self-implemented Vec with Rust's built-in Vec.

4.2 Linear Structures

Data structures are used to store data in a container, and there are several types of data structures available such as arrays, stacks, queues, dequeues, and linked lists. These data structures are called linear data structures because they maintain the order in which the data items are added or removed. Linear data structures have two ends, which can be named as the left and right or the front and back. The names are not important, but they indicate that the organization of data is linear, and this linear characteristic is related to memory, which is linear hardware. This relationship between software and hardware is important to understand.

It is important to note that the term "linear" in linear data structures does not refer to the way data is stored but rather to the way it is accessed. In a linked list, for example, data items may not be adjacent in memory, but the access is still linear.

To differentiate between linear data structures, we need to look at the way items are added and removed, particularly their positions. Some data structures only allow adding items from one end, while others only allow removing items from the other end. Some data structures allow manipulation of data items from both ends. These variations and combinations have led to the creation of many useful data structures in computer science, which are used in various algorithms to perform important and practical tasks.

4.3 Stack

A stack is a linear data structure that has numerous applications, such as in function calls and webpage data recording. It is an ordered collection of data items where new items are added or removed from

the top, while the bottom refers to the opposite end. The bottom is significant because the items closest to it have been stored for the longest time, and the most recently added item will be the first one to be removed. This is known as the Last In First Out (LIFO) or First In Last Out (FILO) principle, which means that newer items are closer to the top, while older items are closer to the bottom.

Stacks are ubiquitous in everyday life, like bricks on a construction site, books on a desk, and plates in a restaurant. To access the bottom brick, book, or plate, we need to remove the items on top first. The schematic diagram of a stack is shown below with some conceptual names.

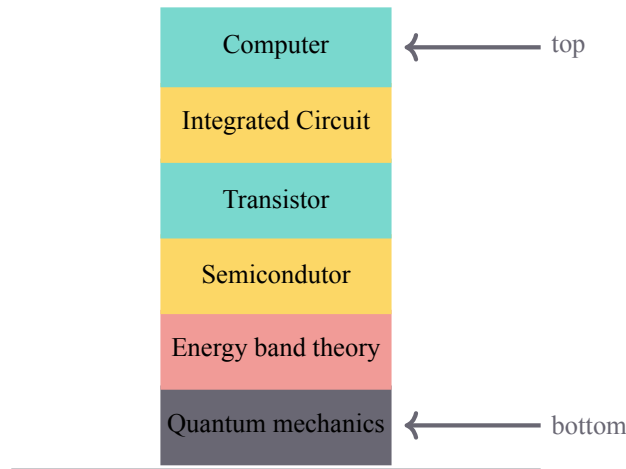


Figure 4.1: Stack

To understand the function of stacks, it is best to observe how they are formed and emptied. Suppose we stack books on a clean desktop one by one; we are creating a stack. Now, if we remove a book, the order of removal is opposite to the order in which the books were placed. The significance of stacks lies in their ability to reverse the order of items, inserting and deleting in the opposite order. The following diagram shows the process of creating and deleting data objects, with particular attention to the order of the data.

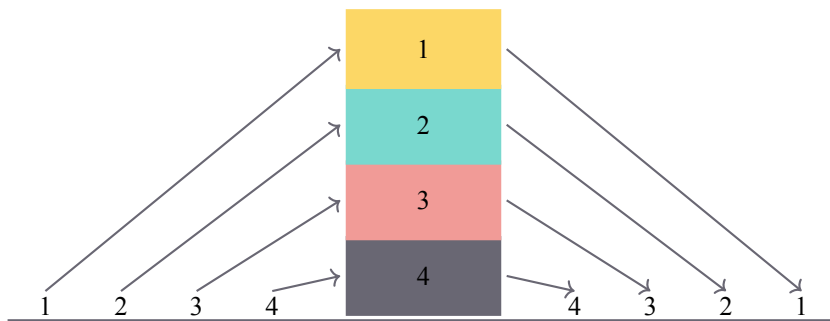


Figure 4.2: Stack reverse order

The ability to reverse the order of items makes the reversal property of stacks particularly useful, as demonstrated in various computer applications. For instance, when browsing news on a web browser, the back function is implemented using a stack. As the user browses web pages, they are stacked, with the current page at the top and the first page at the bottom. Pressing the back button takes the user to the previous page in the opposite order. Without the power of stacks, designing this back function would be nearly impossible. This example highlights the importance of data structures in simplifying certain functions, as choosing the right data structure can make a significant difference.

4.3.1 The Stack Abstract Data Type

A stack is an abstract data type (ADT) that is defined by its structure and operations. It is an ordered collection of items where new items are added and removed from the top. The following operations can be performed on a stack:

- `new()`: creates an empty stack with no arguments and returns an empty stack.
- `push(item)`: adds the data item 'item' to the top of the stack, taking 'item' as an argument and returning nothing.
- `pop()`: removes the top data item from the stack, taking no arguments and returning the data item. The stack is modified.
- `peek()`: returns the top data item from the stack without removing it, taking no arguments and not modifying the stack.
- `is_empty()`: tests if the stack is empty, taking no arguments and returning a Boolean value.
- `size()`: returns the number of data items in the stack as a 'usize' integer, taking no arguments.
- `iter()`: returns an immutable iteration form of the stack, where the stack is not modified and takes no arguments.
- `iter_mut()`: returns a mutable iteration form of the stack, where the stack can be modified and takes no arguments.
- `into_iter()`: changes the stack into an iterable form, consuming the stack and taking no arguments.

To illustrate the stack operations, assume that 's' is an already created empty stack (represented as '[]' here), and the table below shows the results of stack operations, with the stack top on the right.

Table 4.1: Stack Operations

Operation	Value	Return Value
<code>s.is_empty()</code>	[]	true
<code>s.push(1)</code>	[1]	
<code>s.push(2)</code>	[1,2]	
<code>s.peek()</code>	[1,2]	2
<code>s.push(3)</code>	[1,2,3]	
<code>s.size()</code>	[1,2,3]	3
<code>s.is_empty()</code>	[1,2,3]	false
<code>s.push(4)</code>	[1,2,3,4]	
<code>s.push(5)</code>	[1,2,3,4,5]	
<code>s.size()</code>	[1,2,3,4,5]	5
<code>s.pop()</code>	[1,2,3,4]	5
<code>s.push(6)</code>	[1,2,3,4,6]	
<code>s.peek()</code>	[1,2,3,4,6]	6

4.3.2 Implementing a Stack in Rust

The abstract data type of a stack has been defined above, and now we will use Rust to implement it as a data structure. In Rust, abstract data types are implemented by creating a new struct and implementing the operations as functions of the struct. Using the basic data structures provided by Rust to implement a stack and its operations is very helpful.

We will use the `Vec` collection container as the underlying implementation of the stack because it provides an ordered collection mechanism and a set of operation methods. To implement other operations, we only need to choose which end of the `Vec` is the top of the stack. The implementation assumes that the tail of `Vec` will save the top element, and as the stack grows, new items will be added to the end of `Vec`. Since the type of data to be inserted is unknown, we use the generic data type `T`. We have also added three structs, `IntoIter`, `Iter`, and `IterMut`, to complete the three types of iteration.

```
1 // stack.rs
2
3 #[derive(Debug)]
4 struct Stack<T> {
5     size: usize, // stack size
6     data: Vec<T>, // stack data
7 }
8
9 impl<T> Stack<T> {
10     // initialize a stack
11     fn new() -> Self {
12         Self {
13             size: 0,
14             data: Vec::new()
15         }
16     }
17
18     fn is_empty(&self) -> bool {
19         0 == self.size
20     }
21
22     fn len(&self) -> usize {
23         self.size
24     }
25
26     // clear stack
27     fn clear(&mut self) {
28         self.size = 0;
29         self.data.clear();
30     }
31
32     // put the item into the tail of Vec
33     fn push(&mut self, val: T) {
34         self.data.push(val);
35         self.size += 1;
36     }
37
38     // size decrease by 1 and then return the value
39     fn pop(&mut self) -> Option<T> {
40         if 0 == self.size {
41             return None;
42         }
43         self.size -= 1;
44         self.data.pop()
45     }
46
47     // return reference to the top value
48     fn peek(&self) -> Option<&T> {
49         if 0 == self.size {
50             return None;
51         }
52     }
```



```

52
53     self.data.get(self.size - 1)
54 }
55
56 fn peek_mut(&mut self) -> Option<&mut T> {
57     if 0 == self.size {
58         return None;
59     }
60
61     self.data.get_mut(self.size - 1)
62 }
63
64 // Implementing iteration for stack
65 // into_iter: stack modified and becomes a iterator
66 // iter: stack unmodified and get a unmutable iterator
67 // iter_mut: stack unmodified and get a mutable iterator
68 fn into_iter(self) -> IntoIter<T> {
69     IntoIter(self)
70 }
71
72 fn iter(&self) -> Iter<T> {
73     let mut iterator = Iter { stack: Vec::new() };
74     for item in self.data.iter() {
75         iterator.stack.push(item);
76     }
77
78     iterator
79 }
80
81 fn iter_mut(&mut self) -> IterMut<T> {
82     let mut iterator = IterMut { stack: Vec::new() };
83     for item in self.data.iter_mut() {
84         iterator.stack.push(item);
85     }
86
87     iterator
88 }
89 }
90
91 // Implementation of 3 iterations
92 struct IntoIter<T>(Stack<T>);
93 impl<T: Clone> Iterator for IntoIter<T> {
94     type Item = T;
95     fn next(&mut self) -> Option<Self::Item> {
96         if !self.0.is_empty() {
97             self.0.size -= 1;
98             self.0.data.pop()
99         } else {
100             None
101         }
102     }
103 }

```

```
104
105 struct Iter<'a, T: 'a> { stack: Vec<&'a T>, }
106 impl<'a, T> Iterator for Iter<'a, T> {
107     type Item = &'a T;
108     fn next(&mut self) -> Option<Self::Item> {
109         self.stack.pop()
110     }
111 }
112
113 struct IterMut<'a, T: 'a> { stack: Vec<&'a mut T> }
114 impl<'a, T> Iterator for IterMut<'a, T> {
115     type Item = &'a mut T;
116     fn next(&mut self) -> Option<Self::Item> {
117         self.stack.pop()
118     }
119 }
120
121 fn main() {
122     basic();
123     peek();
124     iter();
125
126     fn basic() {
127         let mut s = Stack::new();
128         s.push(1); s.push(2); s.push(3);
129
130         println!("size: {}, {:?}", s.len(), s);
131         println!("pop {:?}, size {}", s.pop().unwrap(), s.len());
132         ;
133         println!("empty: {}, {:?}", s.is_empty(), s);
134
135         s.clear();
136         println!("{:?}", s);
137     }
138
139     fn peek() {
140         let mut s = Stack::new();
141         s.push(1); s.push(2); s.push(3);
142
143         println!("{:?}", s);
144         let peek_mut = s.peek_mut();
145         if let Some(top) = peek_mut {
146             *top = 4;
147         }
148
149         println!("top {:?}", s.peek().unwrap());
150         println!("{:?}", s);
151     }
152
153     fn iter() {
154         let mut s = Stack::new();
155         s.push(1); s.push(2); s.push(3);
```

```

155
156     let sum1 = s.iter().sum::<i32>();
157     let mut addend = 0;
158     for item in s.iter_mut() {
159         *item += 1;
160         addend += 1;
161     }
162
163     let sum2 = s.iter().sum::<i32>();
164     println!("{sum1} + {addend} = {sum2}");
165     assert_eq!(9, s.into_iter().sum::<i32>());
166 }
167 }

```

After executing the code, we obtain the following results.

```

size: 3, Stack { size: 3, data: [1, 2, 3] }
pop 3, size 2
empty: false, Stack { size: 2, data: [1, 2] }
Stack { size: 0, data: [] }
Stack { size: 3, data: [1, 2, 3] }
top 4
Stack { size: 3, data: [1, 2, 4] }
6 + 3 = 9

```

4.3.3 Simple Balanced Parentheses

Previously, we implemented the stack data structure and now we will use it to solve the parenthesis matching problem. Arithmetic expressions that evaluate values contain parentheses as shown below.

$$(5 + 6) \times (7 + 8) / (4 + 3)$$

Similarly, in the Lisp language, the multiply function shown below also has parentheses.

```

1 (defun multiply(n)
2   (* n n))

```

The focus here is on parentheses, as they alter the order of operations and constrain the language semantics. It is essential to ensure that the parentheses are complete; otherwise, the entire expression will be incorrect. Humans can quickly identify incomplete parenthesis, but how can a computer do this? Clearly, the computer must verify whether the parentheses match and report an error if they don't.

In both the examples above, the parentheses must appear in pairs. Parenthesis matching means that each opening symbol has a corresponding closing symbol, and the parentheses are correctly nested so that the computer can process them correctly.

Consider the following correctly matched parenthesis strings and these mismatched parenthesis.

```

((()())())
(((()))
(()((()))())

```

```

(((((((
)))
(()()()

```

To solve the parenthesis matching problem, a deeper understanding of brackets and their matching is needed. When processing symbols from left to right, the nearest left starting bracket '(' must match the next right closing symbol ')' (as shown in Figure 4.3). In addition, the first left starting bracket processed must wait until it matches the last right closing bracket. Ending brackets match starting brackets in the opposite order, from inside to outside. This is a problem that can be solved using a stack.

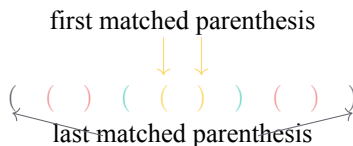


Figure 4.3: Parenthesis matching

Matching parenthesis is crucial in computer programs since it determines the next operation. The challenge is to write an algorithm that can read a string of symbols from left to right and determine whether the brackets match. Parenthesis and their matching are essential for computer programs since bracket and nesting are prevalent in programs.

The implementation of the algorithm for matching parentheses using a stack is quite simple since it only involves push, pop, and judgment operations on the stack. To start, process the bracket string from left to right with an empty stack. If the symbol is a left starting symbol, push it onto the stack; if it is an ending symbol, pop the top element of the stack and match these two symbols. If they match, continue to process the next bracket until the string is completely processed. At the end of the processing, the stack should be empty; if not, it indicates that there are unmatched brackets. Here is a Rust implementation of the bracket matching program.

```

1 // par_checker1.rs
2
3 fn par_checker1(par: &str) -> bool {
4     // adding characters into Vec
5     let mut char_list = Vec::new();
6     for c in par.chars() {
7         char_list.push(c);
8     }
9
10    let mut index = 0;
11    let mut balance = true; // determine if balanced
12    let mut stack = Stack::new();
13    while index < char_list.len() && balance {
14        let c = char_list[index];
15
16        if '(' == c { // if is '(', put data into stack
17            stack.push(c);
18        } else { // if is ')', determine if stack is empty
19            if stack.is_empty() { // empty stack, matched
20                balance = false;
21            } else {
22                let _r = stack.pop();
23            }
24        }
25
26        index += 1;
27    }

```

```

28
29     // parenthesis matched: balanced and empty stack
30     balance && stack.is_empty()
31 }
32
33 fn main() {
34     let sa = "()()()";
35     let sb = "()((((";
36     let res1 = par_checker1(sa);
37     let res2 = par_checker1(sb);
38     println!("{sa} balanced:{res1}, {sb} balanced:{res2}");
39     // ()()() balanced:true, ()((( balanced:false
40 }

```

While the above example only involves matching the parentheses '()', there are actually three commonly used types of parentheses: '()', '[]', and '{}'. These different types of left and right parentheses are often nested together, as shown in Rust, where square brackets '[]' are used for indexing, curly brackets '{}' are used for formatting output, and parentheses '()' are used for function parameters, tuples, mathematical expressions, and more. As long as each symbol maintains its own left starting and right ending relationship, mixed nesting symbols can be used.

```

{ { ( [ ] [ ] ) } ( ) }
[ [ { { ( ( ) ) } } ] ]
[ ] [ ] [ ] ( ) { }

```

All of the parentheses shown above are matched. On the contrary, the following expression is not matched.

```

( } [ ]
( ( ( ) ] ) )
[ { ( ) ]

```

To handle three types of parentheses, the previous parentheses checking program, `par_checker1` (only able to detect '()' parentheses), needs to be extended. However, the algorithm process remains the same. Each left starting parenthesis is pushed onto the stack, waiting for the matching right ending parenthesis to appear. When an ending parenthesis appears, the program checks whether the types of parentheses match. If the two parentheses do not match, then the string does not match. If the entire string has been processed and the stack is empty, then the parentheses expression matches.

To detect whether the types of parentheses match, a new function called `par_match()` has been added. This function can detect the three commonly used types of parentheses. The detection principle is very simple: the program arranges the parentheses in order and checks whether their indices match.

```

1 // par_checker2.rs
2
3 // check if parentheses match of various symbols
4 fn par_match(open: char, close: char) -> bool {
5     let opens = "([{";
6     let closers = ")]}";
7     opens.find(open) == closers.find(close)
8 }
9
10 fn par_checker2(par: &str) -> bool {
11     let mut char_list = Vec::new();

```

```

12     for c in par.chars() {
13         char_list.push(c);
14     }
15
16     let mut index = 0;
17     let mut balance = true;
18     let mut stack = Stack::new();
19     while index < char_list.len() && balance {
20         let c = char_list[index];
21         // check 3 open symbols simultaneously
22         if '(' == c || '[' == c || '{' == c {
23             stack.push(c);
24         } else {
25             if stack.is_empty() {
26                 balance = false;
27             } else {
28                 // determine if match
29                 let top = stack.pop().unwrap();
30                 if !par_match(top, c) {
31                     balance = false;
32                 }
33             }
34         }
35         index += 1;
36     }
37     balance && stack.is_empty()
38 }
39
40 fn main() {
41     let sa = "(){}[]";
42     let sb = "({})[]";
43     let res1 = par_checker2(sa);
44     let res2 = par_checker2(sb);
45     println!("sa balanced:{res1}, sb balanced:{res2}");
46     // (){}[] balanced:true, ({}[]) balanced:false
47 }

```

The current implementation can handle different types of bracket matching problems. However, if the input expression contains non-bracket characters, the program will fail to work properly.

```
(a+b)(c*d)func()
```

The apparent complexity of the problem can be deceiving because the actual issue is still related to detecting matching brackets. Hence, non-bracket characters can be skipped during processing. In the case of the given example, the non-bracket characters can be ignored, leaving only the brackets, resulting in the string: `(){}()`. The problem remains the same as before, and the code can be modified to detect matching brackets even in strings containing non-bracket characters. The following implementation is the modified code based on `par_checker2.rs`.

```

1 // par_checker3.rs
2
3 fn par_checker3(par: &str) -> bool {
4     let mut char_list = Vec::new();

```

```

5     for c in par.chars() { char_list.push(c); }
6
7     let mut index = 0;
8     let mut balance = true;
9     let mut stack = Stack::new();
10    while index < char_list.len() && balance {
11        let c = char_list[index];
12        // open mark, push
13        if '(' == c || '[' == c || '{' == c {
14            stack.push(c);
15        }
16        // close mark, determine if balanced
17        if ')' == c || ']' == c || '}' == c {
18            if stack.is_empty() {
19                balance = false;
20            } else {
21                let top = stack.pop().unwrap();
22                if !par_match(top, c) { balance = false; }
23            }
24        }
25        // skip if not the bracket
26        index += 1;
27    }
28    balance && stack.is_empty()
29 }
30
31 fn main() {
32     let sa = "(2+3){func}[abc]"; let sb = "(2+3)*(3-1";
33     let res1 = par_checker3(sa); let res2 = par_checker3(sb);
34     println!("sa balanced:{res1}, sb balanced:{res2}");
35     // (2+3){func}[abc] balanced:true, (2+3)*(3-1 balanced:false
36 }

```

4.3.4 Converting Decimal Numbers to Binary Numbers

Binary is a fundamental concept in the computer world, as it is the universal data format at the low-level of computing. All values stored in computers are in the form of 0s and 1s, making it critical to convert between binary numbers and ordinary characters for successful interaction with computers. Integer values are commonly used in computer programs and calculations, but we typically learn them in decimal form. For instance, decimal 233(10) can also be represented in binary as 11101001(2).

$$\begin{aligned}
 &2 \times 10^2 + 3 \times 10^1 + 3 \times 10^0 = 233 \\
 &1 \times 2^7 + 1 \times 2^6 + 1 \times 2^5 + 0 \times 2^4 + 1 \times 2^3 + 0 \times 2^2 + 0 \times 2^1 + 1 \times 2^0 = 233
 \end{aligned} \tag{4.1}$$

To convert an integer to binary, the "divide by 2" algorithm (as shown in Figure 4.4) is a simple and effective method that involves using a stack to track the binary result's digits. Starting with a positive integer greater than 0, the algorithm iteratively divides the decimal number by 2, keeping track of the remainders. The first remainder after dividing by 2 determines whether the value is even or odd, with an even number producing a remainder of 0 and an odd number resulting in a remainder of 1. By recording these remainders during the iteration and reversing the sequence, the binary digit sequence is obtained, with the first remainder (i.e., the last digit in the sequence) placed at the bottom of the stack. Popping all the digits from the stack results in the binary representation of the original decimal number.

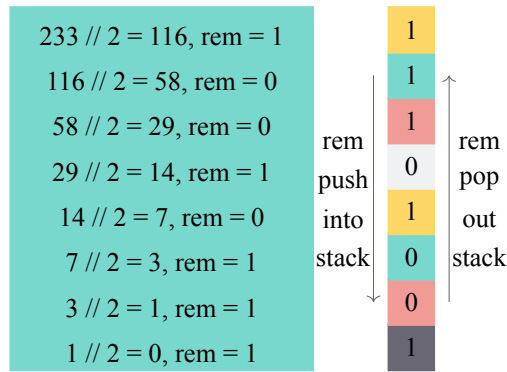


Figure 4.4: Divided by 2

The function shown below takes a decimal parameter and uses Rust's modulus operator (%) to extract the remainder, which is then pushed onto a stack. The parameter is repeatedly divided by 2 until it reaches 0, and the program constructs the binary representation of the original value.

```

1 // divide_by_two.rs
2
3 fn divide_by_two(mut dec_num: u32) -> String {
4     // save the remainder in a stack
5     let mut rem_stack = Stack::new();
6
7     // push rem into the stack
8     while dec_num > 0 {
9         let rem = dec_num % 2;
10        rem_stack.push(rem);
11        dec_num /= 2;
12    }
13
14    // pop out elems from the stack to form a string
15    let mut bin_str = "".to_string();
16    while !rem_stack.is_empty() {
17        let rem = rem_stack.pop().unwrap().to_string();
18        bin_str += &rem;
19    }
20    bin_str
21 }
22
23 fn main() {
24     let num = 10;
25     let bin_str: String = divided_by_two(num);
26     println!("{num} = b{bin_str}");
27     // 10 = b1010
28 }

```

This algorithm for converting from decimal to binary can be extended to perform conversions between any two bases commonly used in computer science, such as binary, octal, and hexadecimal. For example, the decimal number 233(10) corresponds to 351(8) in octal and e9(16) in hexadecimal.

To make the function more general, it can be modified to accept a predetermined conversion base,

replacing the concept of dividing by 2 with dividing by the conversion base. The remainders are still pushed onto a stack until the value being converted reaches 0. However, for bases greater than 10, remainders greater than 10 will inevitably occur, so it is best to represent them as a single character. In the `base_converter` function, we choose to use A-F to represent 10-15, but lowercase letters such as a-f or other character sequences can also be used.

```

1 // base_converter.rs
2
3 fn base_converter(mut dec_num: u32, base: u32) -> String {
4     // digits is the string form of integers(especially for 10
5     // -15)
6     let digits = ['0', '1', '2', '3', '4', '5', '6', '7',
7                 '8', '9', 'A', 'B', 'C', 'D', 'E', 'F'];
8     let mut rem_stack = Stack::new();
9
10    // push rem into the stack
11    while dec_num > 0 {
12        let rem = dec_num % base;
13        rem_stack.push(rem);
14        dec_num /= base;
15    }
16
17    // pop out elems from the stack to form a string
18    let mut base_str = "".to_string();
19    while !rem_stack.is_empty() {
20        let rem = rem_stack.pop().unwrap() as usize;
21        base_str += &digits[rem].to_string();
22    }
23    base_str
24 }
25
26 fn main() {
27     let num1 = 10;
28     let num2 = 43;
29     let bin_str: String = base_converter(num1, 2);
30     let hex_str: String = base_converter(num2, 16);
31     println!("{num1} = b{bin_str}, {num2} = x{hex_str}");
32     // 10 = b1010, 43 = x2B
33 }

```

4.3.5 Prefix, Infix, Postfix Expressions

In mathematics, an arithmetic expression, like $B * C$, has a specific format that makes it easy to understand. For instance, B multiplied by C , with the multiplication operator `*` in between two operands, forms an infix expression. This type of expression is naturally easy to read and follow since it maintains the order of the expression.

However, an infix expression like $A + B * C$, with the `+` and `*` operators sandwiched between operands, requires an understanding of operator priority to determine the order of operations. When operators of equal priority are present, they evaluate from left to right. Parentheses can alter the order of the operation, and arithmetic operator priority places multiplication and division before addition and subtraction.

While humans can easily solve such expressions by mentally adding parentheses, computers require

specific algorithms to perform the same task. A computer can avoid confusion by using fully parenthesized expressions, where each operator has a pair of parentheses, indicating the order of the operation. However, processing these expressions can be difficult since computers work from left to right. Therefore, even simple tasks that seem easy for humans can be complex for computers, as they lack intelligence and the ability to understand expressions intuitively.

To ensure that computers do not confuse the order of operations, a fully parenthesized expression can be used, where each operator is enclosed in a pair of parentheses indicating the order of operations. However, since computers process data from left to right, they may have difficulty jumping between inner and outer parentheses to calculate expressions like $(A + (B * C))$. While humans can do this intuitively, it is challenging for computers since they are rigid and require explicit instructions.

A more intuitive method for computers is to separate the operator and operands by moving the operator outside the parentheses. This creates prefix and postfix expressions, which can be distinguished from the original infix expression. In prefix expressions, operators come before the operands, and in postfix expressions, operators come after the corresponding operands. Calculations are performed by taking the operator and operands, calculating the result as the current value, and then proceeding to subsequent operations until the expression is fully calculated.

For example, the infix expression $A + B$ can be written as $+ A B$ in prefix notation or $A B +$ in postfix notation. By following the rules for each notation, readers can calculate more complex expressions in prefix or postfix notation and confirm the correct result.

Table 4.2: Prefix-Infix-Postfix Expression

prefix	infix	postfix
$A + B$	$+ A B$	$A B +$
$A + B * C$	$+ A * B C$	$A B C * +$
$(A + B) * C$	$* + A B C$	$A B + C *$
$A + B - C$	$- + A B C$	$A B + C -$
$A * B + C$	$+ * A B C$	$A B * C +$
$A * B / C$	$/ * A B C$	$A B * C /$

To analyze complex expressions, readers must understand the importance of parentheses in expressing the order of operations. Even small changes in the placement of parentheses can result in vastly different expressions. While infix expressions require parentheses to disambiguate the order of operations, prefix and postfix expressions do not need them. This is because the order of operations is entirely determined by the position of operators in these notations. As a result, prefix and postfix expressions can clearly express the calculation logic without ambiguity, ensuring that calculations based on them are error-free. The table below shows some examples of complex expressions in different notations.

Table 4.3: Complex Pre-In-Postfix Expressions

infix	prefix	postfix
$A + B * C - D$	$- + A * B C D$	$A B C * + D -$
$A * B - C / D$	$- * A B / C D$	$A B * C D / -$
$A + B + C + D$	$+++ A B C D$	$A B + C + D +$
$(A + B) * (C - D)$	$* + A B - C D$	$A B + C D - *$

Prefix and postfix expressions may appear more complicated to compute at first. For instance, the prefix expression $+ A * B C$ for $A + B * C$ requires calculating $B * C$ before adding it to A . However, the multiplication sign is still inside the addition operator, making it impossible to calculate. To resolve this issue, stacks can be used to reverse the order of operations. By pushing the operands and operators onto separate stacks from left to right, the order of operations can be reversed, as shown in Figure (4.5). This approach enables prefix and postfix expressions to be computed efficiently and accurately.

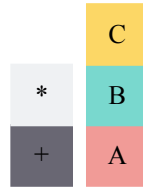


Figure 4.5: Expressions on a stack

To calculate expressions in prefix notation, the operator and operands are pushed onto separate stacks from left to right. To compute the expression, the top operator is popped from the operator stack, and the two top operands are popped from the operand stack. The operator is applied to these operands to obtain a result, which is then pushed back onto the operand stack. This process is repeated until the operator stack is empty. Finally, the top value in the operand stack is the result of the expression. This method is as efficient as fully bracketed expressions since the computer only needs to push and pop from the stacks, and does not need to handle parentheses, as shown in Figure (4.6).

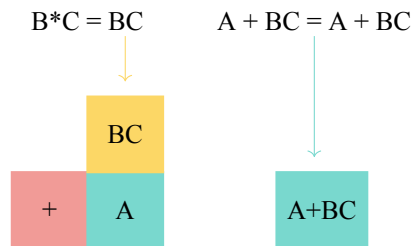


Figure 4.6: calculating expression by the stack

Similarly, postfix expressions can also be calculated using stacks. Only one stack is required to perform the calculation. For example, the postfix expression for $A + B * C$, which is $A B C * +$, is calculated by pushing A , B , and C onto the stack. When the $*$ operator is encountered, the top two operands, B and C , are popped, multiplied, and pushed back onto the stack. Then, when the $+$ operator is encountered, the top two operands, A and BC , are popped, added, and pushed back onto the stack, resulting in the final result of $A + BC$. This approach demonstrates the efficiency of stack-based computations, which can handle expressions in any notation without the need for parentheses.

4.3.6 Conversion of Infix Expressions to Prefix and Postfix

The calculation process demonstrated that converting infix expressions into prefix or postfix expressions is necessary to achieve efficiency. The first step in this process is to obtain prefix or postfix expressions, and a method to achieve this is through the use of fully bracketed expressions.

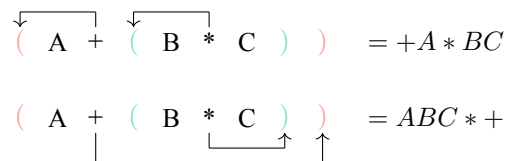


Figure 4.7: infix to prefix and postfix

For instance, $A + B * C$ can be expressed as $(A + (B * C))$ to indicate that multiplication has a higher priority than addition. It is noteworthy that each pair of brackets represent the start and end of

an operand pair. The internal expression of $(A + (B * C))$ is $(B * C)$, and by moving the multiplication symbol to the left bracket position and deleting the left and matching right brackets, the subexpression can be converted to a prefix expression. Similarly, by moving the addition operator to its corresponding left bracket position and deleting the matching right bracket, the complete prefix expression $+ A * B C$ can be obtained. By repeating this operation for all operators, the complete postfix expression can be obtained.

To convert an expression, regardless of whether it is a prefix or postfix expression, the expression must first be converted into a fully bracketed expression based on the order of operations. Once the expression is fully bracketed, the operator inside the brackets can be moved to the position of the left or right bracket to achieve the desired notation. For example, the more complex expression $(A + B) * C - (D + E) / (F + G)$ can be converted to a prefix or postfix expression using this method. Although the result may be complex for humans, computers can handle this process with ease.

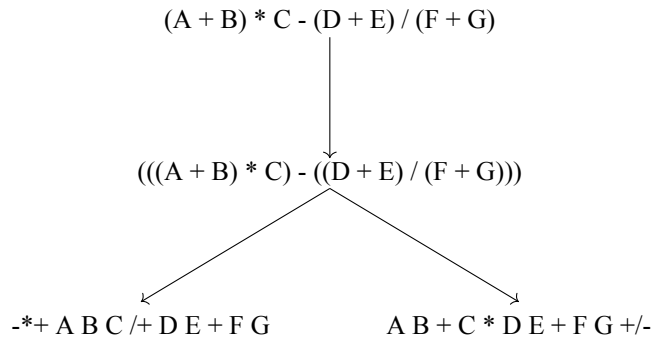


Figure 4.8: Infix to Prefix and Postfix

Obtaining a fully parenthesized expression is a difficult task, and it requires modifying the string by moving and deleting characters. Therefore, this method is not universal enough. A more convenient approach is to handle operators separately. When converting an infix expression, if the operands are not considered, they maintain their original relative positions, and only the operators change position. Therefore, it is not necessary to change the position of operands when encountering them; instead, only operators need to be handled when they are encountered. However, operators have priorities and often reverse the order. The characteristic of reversing the order is inherent to the stack data structure, which can be used to store operators.

To convert an infix expression to a postfix expression, we can use a stack to store operators while scanning the expression from left to right. When an operand is encountered, it is added to the postfix expression. If an operator is encountered, we compare its priority with the operator at the top of the stack. If the current operator has higher priority, it is pushed onto the stack. If the current operator has lower or equal priority, we pop operators from the stack and add them to the postfix expression until we reach an operator with lower priority or an opening parenthesis. When we encounter a closing parenthesis, we pop operators from the stack and add them to the postfix expression until we reach the corresponding opening parenthesis, which is then discarded. The result is a postfix expression that can be evaluated efficiently.

For example, to convert the infix expression $(A + B) * C$, we scan it from left to right. We first encounter the operand A , which is added to the postfix expression. Next, we encounter the operator $+$, which has lower priority than the opening parenthesis, so we push it onto the stack. We then encounter the operand B , which is added to the postfix expression. Finally, we encounter the closing parenthesis, so we pop the operator $+$ from the stack and add it to the postfix expression, followed by the operator $*$. The resulting postfix expression is $A B + C *$.

In the same way, we can convert the infix expression $A * B + C * D$ to the postfix expression $A B * C D * +$, as shown in the figure below.

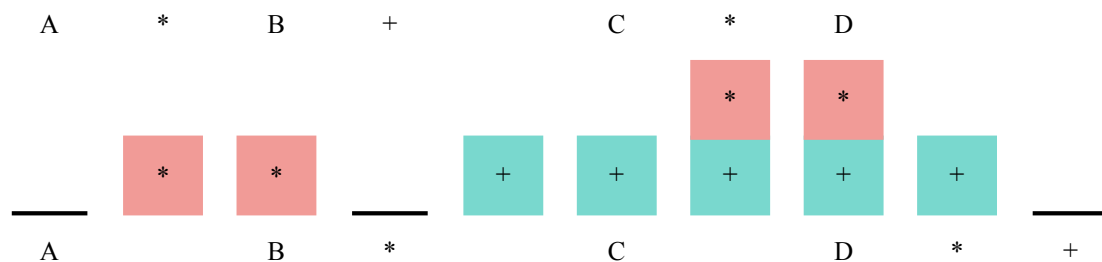


Figure 4.9: Postfix expression by a stack

To convert an infix expression to postfix notation, follow these steps:

1. Create an empty stack `op_stack` to hold operators, an empty list `postfix` to store the output.
2. Split the input infix string into tokens and store them in a list called `src_str`.
3. Iterate through the tokens in `src_str` from left to right:

If the token is an operand, append it to the end of the postfix list.

If the token is a left parenthesis, push it onto the `op_stack`.

If the token is a right parenthesis, pop operators from `op_stack` and append them to postfix until the corresponding left parenthesis is removed.

If the token is an operator (+, -, *, or /), push it onto the `op_stack`. However, first pop any higher or equal priority operators from `op_stack` and append them to postfix.

4. After processing the input, check the `op_stack`. Pop any remaining operators from the `op_stack` and append them to postfix. The final postfix list is the result of the conversion.

Algorithm 4.1: Algorithm: expression conversion from infix to postfix

Input: infix expression

Output: postfix expression

```

1 Create an empty stack op_stack, an empty list postfix
2 Convert the input infix expression into a token list(src_str)
3 for c  $\in$  src_str do
4   if c  $\in$  'A – Z' then
5     | postfix.append(c)
6   else if c == '(' then
7     | op_stack.push(c)
8   else if c  $\in$  '+ – * /' then
9     | while op_stack.peek() prior to c do
10      | postfix.append(op_stack.pop())
11     end
12     | op_stack.push(c)
13   else if c == ')' then
14     | while op_stack.peek() != '(' do
15      | postfix.append(op_stack.pop())
16     end
17   end
18 end
19 while !op_stack.is_empty() do
20   | postfix.append(op_stack.pop())
21 end
22 return ' '.join(postfix)

```

To convert an infix expression to postfix expression, we can use a `HashMap` named "prec" to store the priority of operators. This `HashMap` maps each operator to an integer that is used to compare the priority of operators. The priority of parentheses is assigned the lowest value so that any operator compared with them has higher priority. Operators are limited to "+-*/", while operands are defined as uppercase letters `A - Z` or digits `0 - 9`.

```

1 // infix_to_postfix.rs
2 use std::collections::HashMap;
3
4 fn infix_to_postfix(infix: &str) -> Option<String> {
5     // check parenthesis
6     if !par_checker3(infix) {
7         return None;
8     }
9
10    // set priority of all symbols
11    let mut prec = HashMap::new();
12    prec.insert("(", 1); prec.insert(")", 1);
13    prec.insert("+", 2); prec.insert("-", 2);
14    prec.insert("*", 3); prec.insert("/", 3);
15
16    // ops: svaе operators, postfix: svaе postfix expression
17    let mut ops = Stack::new();
18    let mut postfix = Vec::new();
19    for token in infix.split_whitespace() {
20        // characters range from 0 - 9 and A-Z can
21        // be pushed onto the stack
22        if ("A" <= token && token <= "Z") ||
23            ("0" <= token && token <= "9") {
24            postfix.push(token);
25        } else if "(" == token {
26            // open mark, push onto the stack
27            ops.push(token);
28        } else if ")" == token {
29            // close mark, pop out from stack
30            let mut top = ops.pop().unwrap();
31            while top != "(" {
32                postfix.push(top);
33                top = ops.pop().unwrap();
34            }
35        } else {
36            // check the priority of operators
37            while (!ops.is_empty()) &&
38                (prec[ops.peek().unwrap()]
39                 >= prec[token]) {
40                postfix.push(ops.pop().unwrap());
41            }
42            ops.push(token);
43        }
44    }
45
46    // push the remaining operators onto the stack

```

```

47     while !ops.is_empty() {
48         postfix.push(ops.pop().unwrap())
49     }
50     // pop out operators and create the postfix expression
51     let mut postfix_str = "".to_string();
52     for c in postfix {
53         postfix_str += &c.to_string();
54         postfix_str += " ";
55     }
56
57     Some(postfix_str)
58 }
59
60 fn main() {
61     let infix = "( A + B ) * ( C + D )";
62     let postfix = infix_to_postfix(infix);
63     match postfix {
64         Some(val) => { println!("{infix} -> {val}"); },
65         None => {
66             println!("{infix} isn't a correct infix string");
67         },
68     }
69     // ( A + B ) * ( C + D ) -> A B + C D + *
70 }

```

When calculating postfix expressions, special attention must be given to the "-" and "/" operators. Unlike the "+" and "*" operators, the order of operands matters for "-" and "/". For example, A / B and B / A , $A - B$ and $B - A$ are completely different and cannot be handled like "+", and "*". Assuming the postfix expression is a space-separated string with operators "+-*/" and operands as integers, the output is also an integer. The following are the algorithm steps for calculating postfix expressions:

1. Create an empty stack named `op_stack`.
2. Split the string into a list of symbols.
3. Scan the symbol list from left to right. If the symbol is an operand, convert it from a character to an integer and push the value onto `op_stack`. If the symbol is an operator, pop `op_stack` twice. The second pop is the first operand, and the first pop is the second operand. Perform the arithmetic operation and push the result back onto the operand stack.
4. When the entire input expression has been processed, the result is on the stack. Pop `op_stack` to get the final result of the operation.

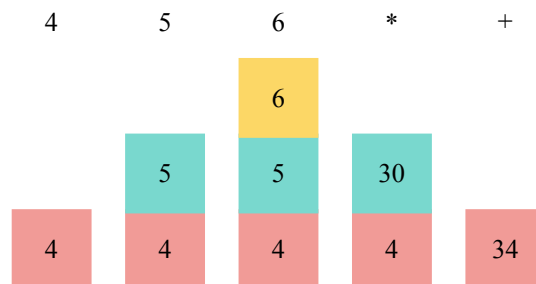


Figure 4.10: Evaluation of Postfix Expression

The specific steps for evaluating postfix expressions are shown in the above figure, and below is the implementation code.

```

1 // postfix_eval.rs
2
3 fn postfix_eval(postfix: &str) -> Option<i32> {
4     // the expression needs at least two operands and
5     // one operator, and two spaces to separate them.
6     if postfix.len() < 5 { return None; }
7
8     let mut ops = Stack::new();
9     for token in postfix.split_whitespace() {
10         // Strings can be compared directly
11         if "0" <= token && token <= "9" {
12             ops.push(token.parse::<i32>().unwrap());
13         } else {
14             // For subtraction and division, the order matters
15             let op2 = ops.pop().unwrap();
16             let op1 = ops.pop().unwrap();
17             let res = do_calc(token, op1, op2);
18             ops.push(res);
19         }
20     }
21     // The value remained in the stack is the result
22     Some(ops.pop().unwrap())
23 }
24
25 // Perform arithmetic operations
26 fn do_calc(op: &str, op1: i32, op2: i32) -> i32 {
27     if "+" == op {
28         op1 + op2
29     } else if "-" == op {
30         op1 - op2
31     } else if "*" == op {
32         op1 * op2
33     } else if "/" == op {
34         if 0 == op2 {
35             panic!("ZeroDivisionError: Invalid operation!");
36         }
37         op1 / op2
38     } else {
39         panic!("OperatorError: Invalid operator: {:?}", op);
40     }
41 }
42
43 fn main() {
44     let postfix = "1 2 + 1 2 + *";
45     let res = postfix_eval(postfix);
46     match res {
47         Some(val) => println!("res = {val}"),
48         None => println!("{postfix} isn't a valid postfix"),
49     }
50     // res = 9
51 }

```


4.4 Queue

A queue is an ordered collection of items that has a front and a rear end. New items are added at the rear, and items are removed from the front. Each element added to the queue moves towards the front until it becomes the next item to be removed. This type of ordering is known as First In First Out (FIFO). A stack, in contrast, uses Last In First Out (LIFO) ordering.

Queues are commonly used in everyday situations such as long lines of people waiting to board a train or bus, or at self-service restaurants. Queues have limited behavior because they have only one entrance and one exit. It is not possible to cut in line or leave early; one must wait for their turn. While it is true that real-life queues and queue data structures may allow cutting in line, this discussion does not consider that possibility.

Operating systems use queues as a data structure to control processes. Multiple different queues are used in scheduling algorithms to prioritize executing programs as quickly as possible and service as many users as possible. When typing on a keyboard, a delay may occur before characters appear on the screen. This is because the keystrokes are placed in a buffer similar to a queue.



Figure 4.11: Queue

4.4.1 The Queue Abstract Data Type

A queue is a data structure that maintains the FIFO sorting property. It is defined by the following structure and operations:

- `new()` creates a new queue, takes no parameters, and returns an empty queue.
- `enqueue(item)` adds a new item to the end of the queue, takes `item` as a parameter, no return.
- `dequeue()` removes an item from the front of the queue, takes no parameters, returns the removed item, and modifies the queue.
- `is_empty()` checks whether the queue is empty, takes no parameters, and returns a Boolean value.
- `size()` returns the number of items in the queue, takes no parameters, and returns a `usize` integer.
- `iter()` returns an immutable iteration of the queue, the queue remains unchanged, and takes no parameters.
- `iter_mut()` returns a mutable iteration of the queue, the queue is mutable, and takes no parameters.
- `into_iter()` changes the queue into an iteration, consumes the queue, and takes no parameters.

Assuming an empty queue `q` has already been created, the table below shows the expected results of various queue operations, with the front of the queue on the left:

Table 4.4: Queue Operations

operation	value	return
<code>q.is_empty()</code>	<code>[]</code>	<code>true</code>
<code>q.enqueue(1)</code>	<code>[1]</code>	
<code>q.enqueue(2)</code>	<code>[1,2]</code>	
<code>q.enqueue(3)</code>	<code>[1,2,3]</code>	
<code>q.dequeue()</code>	<code>[2,3]</code>	<code>1</code>
<code>q.enqueue(4)</code>	<code>[2,3,4]</code>	
<code>q.enqueue(5)</code>	<code>[2,3,4,5]</code>	
<code>q.dequeue()</code>	<code>[3,4,5]</code>	<code>2</code>
<code>q.size()</code>	<code>[3,4,5]</code>	<code>3</code>
<code>q.is_empty()</code>	<code>[3,4,5]</code>	<code>false</code>

4.4.2 Implementing a Queue in Rust

To implement a queue in Rust, we can use a `Vec` with the left end as the rear end of the queue and the right end as the front end. Adding an element to the queue has a complexity of $O(n)$, while removing an element has a complexity of $O(1)$. To prevent the queue from growing infinitely, we can set a `cap` parameter to limit its length.

```

1 // queue.rs
2
3 #[derive(Debug)]
4 struct Queue<T> {
5     cap: usize,    // capacity
6     data: Vec<T>, // store elements
7 }
8
9 impl<T> Queue<T> {
10     fn new(size: usize) -> Self {
11         Self { cap: size, data: Vec::with_capacity(size) }
12     }
13
14     fn is_empty(&self) -> bool { 0 == Self::len(&self) }
15
16     fn is_full(&self) -> bool { self.len() == self.cap }
17
18     fn len(&self) -> usize { self.data.len() }
19
20     fn clear(&mut self) {
21         self.data = Vec::with_capacity(self.cap);
22     }
23
24     fn enqueue(&mut self, val: T) -> Result<(), String> {
25         if self.len() == self.cap { // check left space
26             return Err("No space available".to_string());
27         }
28         self.data.insert(0, val);
29         Ok(())
30     }
31
32     // pop out values
33     fn dequeue(&mut self) -> Option<T> {
34         if self.len() > 0 {
35             self.data.pop()
36         } else {
37             None
38         }
39     }
40
41     // Implemntion of iterations for a queue
42     // into_iter: queue modified and became a iterator
43     // iter: queue unmodified and get a unmutable iterator
44     // iter_mut: queue unmodified and get a mutable iterator
45     fn into_iter(self) -> IntoIter<T> {
46         IntoIter(self)

```

```

47     }
48
49     fn iter(&self) -> Iter<T> {
50         let mut iterator = Iter { stack: Vec::new() };
51         for item in self.data.iter() {
52             iterator.stack.push(item);
53         }
54         iterator
55     }
56
57     fn iter_mut(&mut self) -> IterMut<T> {
58         let mut iterator = IterMut { stack: Vec::new() };
59         for item in self.data.iter_mut() {
60             iterator.stack.push(item);
61         }
62         iterator
63     }
64 }
65
66 // Implementation of 3 iterations
67 struct IntoIter<T>(Queue<T>);
68 impl<T: Clone> Iterator for IntoIter<T> {
69     type Item = T;
70     fn next(&mut self) -> Option<Self::Item> {
71         if !self.0.is_empty() {
72             Some(self.0.data.remove(0))
73         } else {
74             None
75         }
76     }
77 }
78
79 struct Iter<'a, T: 'a> { stack: Vec<&'a T>, }
80 impl<'a, T> Iterator for Iter<'a, T> {
81     type Item = &'a T;
82     fn next(&mut self) -> Option<Self::Item> {
83         if 0 != self.stack.len() {
84             Some(self.stack.remove(0))
85         } else {
86             None
87         }
88     }
89 }
90
91 struct IterMut<'a, T: 'a> { stack: Vec<&'a mut T> }
92 impl<'a, T> Iterator for IterMut<'a, T> {
93     type Item = &'a mut T;
94     fn next(&mut self) -> Option<Self::Item> {
95         if 0 != self.stack.len() {
96             Some(self.stack.remove(0))
97         } else {
98             None

```

```

99         }
100    }
101 }

```

```

1  fn main() {
2      basic();
3      iter();
4      fn basic() {
5          let mut q = Queue::new(4);
6          let _r1 = q.enqueue(1); let _r2 = q.enqueue(2);
7          let _r3 = q.enqueue(3); let _r4 = q.enqueue(4);
8          if let Err(error) = q.enqueue(5) {
9              println!("Enqueue error: {error}");
10         }
11         if let Some(data) = q.dequeue() {
12             println!("dequeue data: {data}");
13         } else {
14             println!("empty queue");
15         }
16         println!("empty: {}, len: {}", q.is_empty(), q.len());
17         println!("full: {}", q.is_full());
18         println!("q: {:?}", q);
19         q.clear();
20         println!("{:?}", q);
21     }
22
23     fn iter() {
24         let mut q = Queue::new(4);
25         let _r1 = q.enqueue(1); let _r2 = q.enqueue(2);
26         let _r3 = q.enqueue(3); let _r4 = q.enqueue(4);
27         let sum1 = q.iter().sum::<i32>();
28         let mut addend = 0;
29         for item in q.iter_mut() {
30             *item += 1;
31             addend += 1;
32         }
33         let sum2 = q.iter().sum::<i32>();
34         println!("{sum1} + {addend} = {sum2}");
35         println!("sum = {}", q.into_iter().sum::<i32>());
36     }
37 }

```

Here is the result of the execution.

```

Enqueue error: No space available
dequeue data: 1
empty: false, len: 3
full: false
q: Queue { cap: 4, data: [4, 3, 2] }
Queue { cap: 4, data: [] }
10 + 4 = 14
sum = 14

```

4.4.3 Hot Potato

A queue is commonly used to simulate real-world scenarios where data is processed in a FIFO (first-in, first-out) manner. For instance, the game of hot potato involves children forming a circle and passing a potato quickly to the child next to them, with the objective of keeping the potato moving as if it were hot (see Figure 4.12). When the passing action stops, the child holding the potato is removed from the circle. The game continues until only one child remains. The game is similar to musical chairs, where people perform an action such as speaking, dancing, or leaving their seat when the music stops.

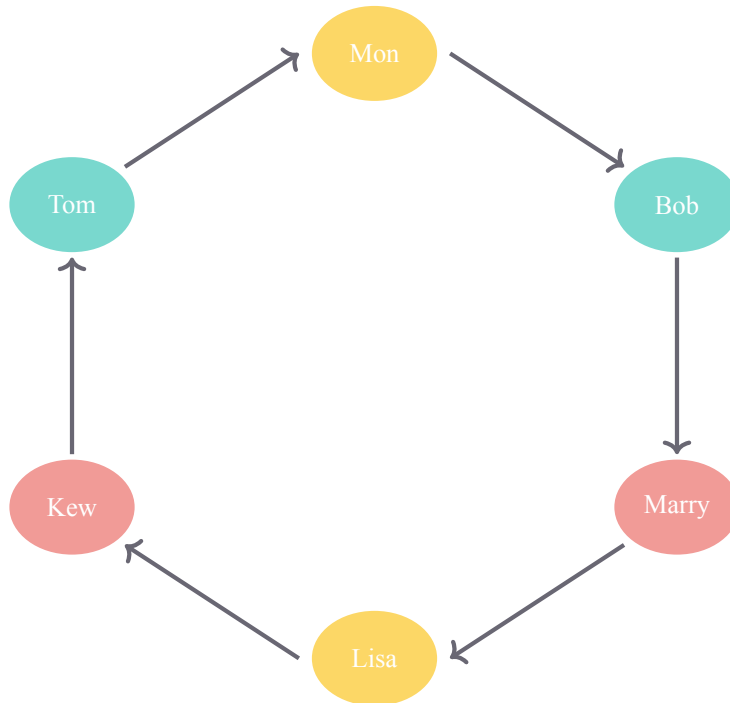


Figure 4.12: hot potato

The game of hot potato is comparable to the Josephus problem, a legendary story recounted by the historian Flavius Josephus. Josephus and his comrades were trapped by the Roman army in a cave and chose to die instead of becoming Roman slaves. They formed a circle and counted clockwise to the eighth person, who was then killed. This continued until only one person was left alive. As a mathematician, Josephus figured out where he should sit to be the last person alive and ultimately joined the Roman side. While there are variations of this story, the fundamental ideas of both the hot potato game and the Josephus problem can be simulated using a queue.



Figure 4.13: Simulation of game of hot potato

To implement the hot potato game, the program accepts multiple names of children and a constant num to set the count of how many children to skip before removing one. Assuming the child holding the hot potato is always at the front of the queue, they leave the queue and re-enter at the back, effectively passing the potato to the next child who must be at the front of the queue. After num rounds of dequeuing

and enqueueing, the child at the front of the queue is permanently removed. The process repeats until only one name remains.

The queue model for the Hot Potato game is clearly illustrated in the two figures above. Here is the implementation of the game based on this model.

```

1 // hot_potato.rs
2
3 fn hot_potato(names: Vec<&str>, num: usize) -> &str {
4     // Initialize the queue and enqueue the names
5     let mut q = Queue::new(names.len());
6     for name in names {
7         let _nm = q.enqueue(name);
8     }
9
10    while q.size() > 1 {
11        // Dequeue and enqueue the names,
12        // which simulates passing the potato
13        for _i in 0..num {
14            let name = q.dequeue().unwrap();
15            let _rm = q.enqueue(name);
16        }
17
18        // After num dequeue/enqueue cycles
19        // remove one person
20        let _rm = q.dequeue();
21    }
22
23    q.dequeue().unwrap()
24 }
```

```

1 fn main() {
2     let name = vec!["Mon","Tom","Kew","Lisa","Marry","Bob"];
3     let survivor = hot_potato(name, 8);
4     println!("The survival person is {survivor}");
5     // The survival person is Marry
6 }
```

It is worth noting that the counting value used in the implementation is 8, which is greater than the number of people in the queue (which is 6). However, this is not a problem because the queue functions like a circle that returns to the head after reaching the tail, and continues until the count is reached. Therefore, there will always be someone dequeued in the end.

4.5 Deque

A deque also is a linear data structure with two ends: the front and the rear. Unlike a queue, a deque allows items to be added and removed from both ends. This flexibility makes it a hybrid linear structure that can function as both a stack and a queue.

Although a deque shares similarities with both a stack and a queue, it does not enforce LIFO or FIFO ordering. The order of adding and removing data determines whether it acts like a stack or a queue. It is important to note that a deque should not be used as a replacement for either a stack or a queue, as each data structure has its own unique properties and is designed for specific computational purposes. The image below provides an example of Deque.



Figure 4.14: Deque

4.5.1 The Deque Abstract Data Type

The deque data structure is an ordered collection of items that allows adding and removing items from either end. It is defined by the following operations:

- `new()` creates a new deque with no arguments and returns an empty deque.
- `add_front(item)` adds item to the front of the deque, requiring item as a parameter, and does not return any content.
- `add_rear(item)` adds item to the back of the deque, requiring item as a parameter, and does not return any content.
- `remove_front()` removes the front item from the deque, requiring no arguments, returns item, and modifies the deque.
- `remove_rear()` removes the rear item from the deque, requiring no arguments, returns item, and modifies the deque.
- `is_empty()` tests whether the deque is empty, requires no arguments, and returns a boolean value.
- `size()` returns the number of items in the deque, requires no arguments, and returns a usize integer.
- `iter()` returns an immutable iteration form of the deque, with the deque unchanged and no arguments required.
- `iter_mut()` returns a mutable iteration form of the deque, with the deque mutable and no arguments required.
- `into_iter()` changes the deque into an iterable form, with the deque consumed and no arguments required.

Assuming that `d` is an already created empty deque, the following table shows the results of a series of operations. Note that the front is on the right-hand side. When moving items in and out, it is important to keep track of the front and back contents, as modifying both ends can make the result look somewhat messy.

Table 4.5: Deque Operations

operations	value	return
<code>d.is_empty()</code>	<code>[]</code>	<code>true</code>
<code>d.add_rear(1)</code>	<code>[1]</code>	
<code>d.add_rear(2)</code>	<code>[2,1]</code>	
<code>d.add_front(3)</code>	<code>[2,1,3]</code>	
<code>d.add_front(4)</code>	<code>[2,1,3,4]</code>	
<code>d.remove_rear()</code>	<code>[1,3,4]</code>	2
<code>d.remove_front()</code>	<code>[1,3]</code>	4
<code>d.size()</code>	<code>[1,3]</code>	2
<code>d.is_empty()</code>	<code>[1,3]</code>	<code>false</code>
<code>d.add_front(2)</code>	<code>[1,3,2]</code>	
<code>d.add_rear(4)</code>	<code>[4,1,3,2]</code>	
<code>d.size()</code>	<code>[4,1,3,2]</code>	4
<code>d.is_empty()</code>	<code>[4,1,3,2]</code>	<code>false</code>
<code>d.add_rear(5)</code>	<code>[5,4,1,3,2]</code>	
<code>d.add_front(10)</code>	<code>[5,4,1,3,2,10]</code>	

4.5.2 Implementing a Deque in Rust

To implement the deque abstract data type in Rust, we can use a `Vec` as the underlying data structure. The left end of the `Vec` serves as the rear of the deque, while the right end serves as the front. To prevent the deque from growing indefinitely, a capacity parameter is added to control the maximum length of the deque.

```
1 // deque.rs
2
3 // deque definition
4 #[derive(Debug)]
5 struct Deque<T> {
6     cap: usize,    // capacity
7     data: Vec<T>, // save elements in data
8 }
9
10 impl<T> Deque<T> {
11     fn new(cap: usize) -> Self {
12         Self {
13             cap: cap,
14             data: Vec::with_capacity(cap),
15         }
16     }
17
18     fn is_empty(&self) -> bool {
19         0 == self.len()
20     }
21
22     fn is_full(&self) -> bool {
23         self.len() == self.cap
24     }
25
26     fn len(&self) -> usize {
27         self.data.len()
28     }
29
30     fn clear(&mut self) {
31         self.data = Vec::with_capacity(self.cap);
32     }
33
34     // use the tail of a Vec as the start of the deque
35     fn add_front(&mut self, val: T) -> Result<(), String> {
36         if self.len() == self.cap {
37             return Err("No space available".to_string());
38         }
39         self.data.push(val);
40
41         Ok(())
42     }
43
44     // the head of the Vec is the tail of the deque
45     fn add_rear(&mut self, val: T) -> Result<(), String> {
46         if self.len() == self.cap {
```



```

47         return Err("No space available".to_string());
48     }
49     self.data.insert(0, val);
50
51     Ok(())
52 }
53
54 // remove data from queue head
55 fn remove_front(&mut self) -> Option<T> {
56     if self.len() > 0 {
57         self.data.pop()
58     } else {
59         None
60     }
61 }
62
63 // remove data from queue tail
64 fn remove_rear(&mut self) -> Option<T> {
65     if self.len() > 0 {
66         Some(self.data.remove(0))
67     } else {
68         None
69     }
70 }
71
72 // Implementation of iteration for the deque
73 // into_iter: deque modified and became a iterator
74 // iter: deque unmodified and get a unmutable iterator
75 // iter_mut: deque unmodified and get a mutable iterator
76 fn into_iter(self) -> IntoIter<T> {
77     IntoIter(self)
78 }
79
80 fn iter(&self) -> Iter<T> {
81     let mut iterator = Iter { stack: Vec::new() };
82     for item in self.data.iter() {
83         iterator.stack.push(item);
84     }
85
86     iterator
87 }
88
89 fn iter_mut(&mut self) -> IterMut<T> {
90     let mut iterator = IterMut { stack: Vec::new() };
91     for item in self.data.iter_mut() {
92         iterator.stack.push(item);
93     }
94
95     iterator
96 }
97 }
98

```

```

99 // Implementing 3 iterations
100 struct IntoIter<T>(Deque<T>);
101 impl<T: Clone> Iterator for IntoIter<T> {
102     type Item = T;
103     fn next(&mut self) -> Option<Self::Item> {
104         // first element of a tuple is not empty
105         if !self.0.is_empty() {
106             Some(self.0.data.remove(0))
107         } else {
108             None
109         }
110     }
111 }
112
113 struct Iter<'a, T: 'a> { stack: Vec<&'a T>, }
114 impl<'a, T> Iterator for Iter<'a, T> {
115     type Item = &'a T;
116     fn next(&mut self) -> Option<Self::Item> {
117         if 0 != self.stack.len() {
118             Some(self.stack.remove(0))
119         } else {
120             None
121         }
122     }
123 }
124
125 struct IterMut<'a, T: 'a> { stack: Vec<&'a mut T> }
126 impl<'a, T> Iterator for IterMut<'a, T> {
127     type Item = &'a mut T;
128     fn next(&mut self) -> Option<Self::Item> {
129         if 0 != self.stack.len() {
130             Some(self.stack.remove(0))
131         } else {
132             None
133         }
134     }
135 }
136
137 fn main() {
138     basic();
139     iter();
140
141     fn basic() {
142         let mut d = Deque::new(4);
143         let _r1 = d.add_front(1);
144         let _r2 = d.add_front(2);
145         let _r3 = d.add_rear(3);
146         let _r4 = d.add_rear(4);
147
148         if let Err(error) = d.add_front(5) {
149             println!("add_front error: {error}");
150         }

```

```

151         println!("{:?}", d);
152
153         match d.remove_rear() {
154             Some(data) => println!("remove rear data {data}"),
155             None => println!("empty deque"),
156         }
157
158         match d.remove_front() {
159             Some(data) => println!("remove front data {data}"),
160             None => println!("empty deque"),
161         }
162         println!("empty: {}, len: {}", d.is_empty(), d.len());
163         println!("full: {}, {:?}", d.is_full(), d);
164
165         d.clear();
166         println!("{:?}", d);
167     }
168
169     fn iter() {
170         let mut d = Deque::new(4);
171         let _r1 = d.add_front(1);
172         let _r2 = d.add_front(2);
173         let _r3 = d.add_rear(3);
174         let _r4 = d.add_rear(4);
175
176         let sum1 = d.iter().sum::<i32>();
177         let mut addend = 0;
178         for item in d.iter_mut() {
179             *item += 1;
180             addend += 1;
181         }
182
183         let sum2 = d.iter().sum::<i32>();
184         println!("{sum1} + {addend} = {sum2}");
185         assert_eq!(14, d.into_iter().sum::<i32>());
186     }
187 }

```

Here is an example output of an execution.

```

add_front error: No space available
Deque { cap: 4, data: [4, 3, 1, 2] }
remove rear data 4
remove front data 2
empty: false, len: 2
full: false, Deque { cap: 4, data: [3, 1] }
Deque { cap: 4, data: [] }
10 + 4 = 14

```

As we can see from the output, a deque has similarities to both a queue and a stack, and can be thought of as a combination of the two. The specific implementation of a deque can vary depending on the situation, and which end is considered the front and which end is considered the rear should be determined based on the context.

4.5.3 Palindrome Checker

Palindromes are strings in which the characters at the same position from both ends are the same, such as "radar", "sos", and "rustsur". In this section, we will present an algorithm for checking whether a given string is a palindrome.

One approach is to use a queue to enqueue the characters of the string and then dequeue the characters to compare them with the reversed original string. This method is simple, but it requires a lot of memory. A more efficient approach is to use a deque.



Figure 4.15: Palindrome Checker

To check for palindromes using a deque, we start by processing the input string from left to right and adding each character to the rear of the deque. At this point, the front of the deque holds the first character of the string, while the rear holds the last character. We can then use the deque's feature of dequeuing from both ends to compare the front and rear characters. If they match, we continue dequeuing the front and rear characters until either all the characters are used up, leaving an empty deque, or a deque of size 1 is left. In both cases, the string is a palindrome. Any other situation indicates that the string is not a palindrome. Below is the code implementation of palindrome detection using a deque.

```

1 // palindrome_checker.rs
2 fn palindrome_checker(pal: &str) -> bool {
3     let mut d = Deque::new(pal.len());
4     for c in pal.chars() {
5         let _r = d.add_rear(c);
6     }
7
8     let mut is_pal = true;
9     while d.size() > 1 && is_pal {
10         let head = d.remove_front();
11         let tail = d.remove_rear();
12         if head != tail {
13             is_pal = false;
14         }
15     }
16     is_pal
17 }
18
19 fn main() {
20     let pal = "rustsur";
21     let is_pal = palindrome_checker(pal);
22     println!("{pal} is palindrome string: {is_pal}");
23     // rustsur is palindrome string: true
24
25     let pal = "panda";
26     let is_pal = palindrome_checker(pal);
27     println!("{pal} is palindrome string: {is_pal}");
28     // panda is palindrome string: false
29 }

```

4.6 LinkedList

An ordered collection of data items is important to maintain the relative position of data and efficient indexing. Arrays and linked lists both allow data to be collected in an ordered manner and saved in relative positions, making them suitable for implementing ordered data types. For example, Rust's default implementation of `Vec` uses an array. This section focuses on linked lists, which offer some unique advantages.

Unlike arrays, linked lists do not require elements to be stored in contiguous memory. Each item in the collection has a reference to the next item, so the items can be randomly placed without the need to allocate a block of memory, resulting in higher efficiency. To use a linked list, we need to specify the position of the first item explicitly. Once we know the position of the first item, we can determine the position of the second item and so on until the end of the entire linked list.

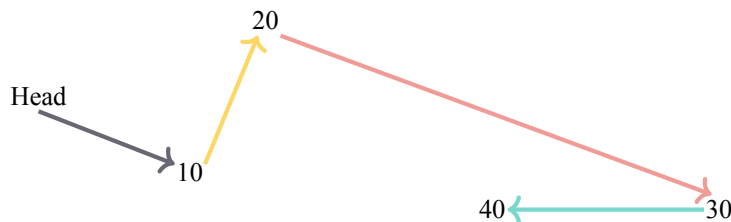


Figure 4.16: Linked list

Usually, linked lists provide a reference to the head of the list, and the last item must be set to null or an empty next item, also known as the tail. With this setup, we can easily traverse the linked list from the head to the tail, or vice versa, by following the references between items. This makes linked lists an efficient data structure for implementing other ordered data types.

4.6.1 The LinkedList Abstract Data Type

The abstract data type of a linked list consists of its structure and operations. As previously mentioned, a linked list is an ordered collection of nodes, and the entire list can be traversed from the head node. The structure of a linked list is simple, with each node containing an element and a reference to the next node. The following definitions and operations apply to a linked list:

- `new()`: Creates a new head node that points to a `Node`, takes no arguments, and returns a pointer.
- `push(item)`: Adds a new `Node`, takes an item parameter, and returns `None`.
- `pop()`: Deletes the head node of the linked list, takes no arguments, and returns a `Node`.
- `peek()`: Returns a referent to the head node of the linked list, takes no arguments.
- `peek_mut()`: Returns the head node of the linked list, takes no arguments, and returns a mutable reference to the node.
- `is_empty()`: Returns a Boolean value indicating whether the current linked list is empty, takes no arguments.
- `size()`: Returns the length of the linked list as a usize integer value, takes no arguments.
- `iter()`: Returns an immutable iteration of the linked list without modifying the linked list, takes no arguments.
- `iter_mut()`: Returns a mutable iteration of the linked list with the linked list mutable, takes no arguments.
- `into_iter()`: Consumes the linked list and returns an iterable form of the linked list, takes no arguments.

Assuming that `l` is an empty linked list, the table below shows the result of a sequence of linked list operations, with `Link<num>` representing the address pointing to the node containing `num`, and the left end of the list being the head node.

Table 4.6: LinkedList Operations

Operation	Value	Return
<code>l.is_empty()</code>	<code>[None->None]</code>	<code>true</code>
<code>l.push(1)</code>	<code>[1->None]</code>	
<code>l.push(2)</code>	<code>[2->1->None]</code>	
<code>l.push(3)</code>	<code>[3->2->1->None]</code>	
<code>l.peek()</code>	<code>[3->2->1->None]</code>	<code>Link<3></code>
<code>l.pop()</code>	<code>[2->1->None]</code>	<code>3</code>
<code>l.size()</code>	<code>[2->1->None]</code>	<code>2</code>
<code>l.push(4)</code>	<code>[4->2->1->None]</code>	
<code>l.peek_mut()</code>	<code>[4->2->1->None]</code>	<code>mut Link<4></code>
<code>l.iter()</code>	<code>[4->2->1->None]</code>	<code>[4,2,1]</code>
<code>l.is_empty()</code>	<code>[4->2->1->None]</code>	<code>false</code>
<code>l.size()</code>	<code>[4->2->1->None]</code>	<code>3</code>
<code>l.into_iter()</code>	<code>[None->None]</code>	<code>[4,2,1]</code>

4.6.2 Implementing a LinkedList in Rust

To represent an item in a linked list, we use a node abstraction that includes the data item and the address of the next item. The node also provides methods to access and modify the data item. As shown in the figure below, a Node consists of data and the address of the next node.

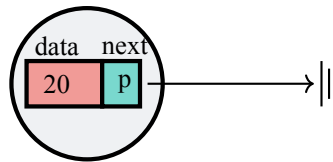


Figure 4.17: Node

In Rust, `None` is used to represent the absence of a next node in both the `Node` and the linked list. The new function initializes a grounded node with its next field set to `None`. Explicitly assigning `None` to the next field is a good practice that helps avoid dangling pointers, a common issue in languages like C/C++.

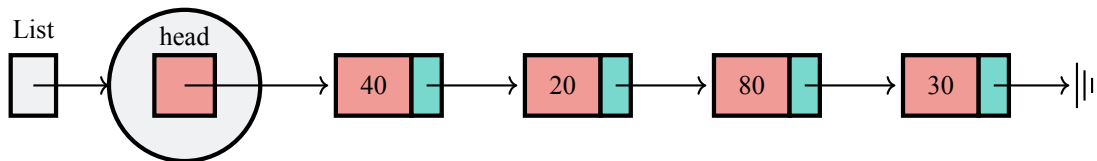


Figure 4.18: LinkedList

The following code shows the implementation of the linked list, with the node link defined as `Link` for code clarity.

```

1 // linked_list.rs
2
3 // The node link uses a Box pointer (size determined),
4 // because only with a determined size can memory be allocated.
5 type Link<T> = Option<Box<Node<T>>>;
6

```

```

7 // Linked list definition
8 struct List<T> {
9     size: usize,    // Number of nodes in the linked list
10    head: Link<T>, // Head node
11 }
12
13 // Linked list node
14 struct Node<T> {
15     elem: T,        // data
16     next: Link<T>, // Link to the next node
17 }
18
19 impl<T> List <T> {
20     fn new() -> Self {
21         Self {
22             size: 0,
23             head: None
24         }
25     }
26
27     fn is_empty(&self) -> bool { 0 == self.size }
28
29     fn len(&self) -> usize { self.size }
30
31     fn clear(&mut self) {
32         self.size = 0;
33         self.head = None;
34     }
35
36     // add a new node ahead of the head node
37     fn push(&mut self, elem: T) {
38         let node = Box::new(Node {
39             elem: elem,
40             next: self.head.take(),
41         });
42         self.head = Some(node);
43         self.size += 1;
44     }
45
46     // take will move data out from node and left a vacuum
47     fn pop(&mut self) -> Option<T> {
48         self.head.take().map(|node| {
49             self.head = node.next;
50             self.size -= 1;
51             node.elem
52         })
53     }
54
55     // peek get a unmutable reference
56     fn peek(&self) -> Option<&T> {
57         self.head.as_ref().map(|node| &node.elem )
58     }

```

```

59
60 // peek_mut get a mutable reference
61 fn peek_mut(&mut self) -> Option<&mut T> {
62     self.head.as_mut().map(|node| &mut node.elem )
63 }
64
65 // Implementation of iteration for the linked list.
66 // into_iter: makes the linked list an iterator
67 //             by consuming it
68 // iter: returns an immutable iterator without modifying
69 //       the linked list
70 // iter_mut: returns a mutable iterator without modifying
71 //           the linked list
72 fn into_iter(self) -> IntoIter<T> {
73     IntoIter(self)
74 }
75
76 fn iter(&self) -> Iter<T> {
77     Iter { next: self.head.as_deref() }
78 }
79
80 fn iter_mut(&mut self) -> IterMut<T> {
81     IterMut { next: self.head.as_deref_mut() }
82 }
83 }
84
85 // Implementation of three iterations
86 struct IntoIter<T>(List<T>);
87 impl<T> Iterator for IntoIter<T> {
88     type Item = T;
89     fn next(&mut self) -> Option<Self::Item> {
90         // (List<T>) tuple's 0th item
91         self.0.pop()
92     }
93 }
94
95 struct Iter<'a, T: 'a> { next: Option<&'a Node<T>> }
96 impl<'a, T> Iterator for Iter<'a, T> {
97     type Item = &'a T;
98     fn next(&mut self) -> Option<Self::Item> {
99         self.next.map(|node| {
100             self.next = node.next.as_deref();
101             &node.elem
102         })
103     }
104 }
105
106 struct IterMut<'a, T: 'a> { next: Option<&'a mut Node<T>> }
107 impl<'a, T> Iterator for IterMut<'a, T> {
108     type Item = &'a mut T;
109     fn next(&mut self) -> Option<Self::Item> {
110         self.next.take().map(|node| {

```



```

111         self.next = node.next.as_deref_mut();
112         &mut node.elem
113     })
114 }
115 }
116
117 // Custom implementation of Drop for the linked list
118 impl<T> Drop for List<T> {
119     fn drop(&mut self) {
120         let mut link = self.head.take();
121         while let Some(mut node) = link {
122             link = node.next.take();
123         }
124     }
125 }
126
127 fn main() {
128     basic_test();
129     into_iter_test();
130     iter_test();
131     iter_mut_test();
132
133     fn basic_test() {
134         let mut list = List::new();
135         list.push(1); list.push(2); list.push(3);
136
137         assert_eq!(list.len(), 3);
138         assert_eq!(list.is_empty(), false);
139         assert_eq!(list.pop(), Some(3));
140         assert_eq!(list.peek(), Some(&2));
141         assert_eq!(list.peek_mut(), Some(&mut 2));
142
143         list.peek_mut().map(|val| {
144             *val = 4;
145         });
146
147         assert_eq!(list.peek(), Some(&4));
148         list.clear();
149         println!("basics test Ok!");
150     }
151
152     fn into_iter_test() {
153         let mut list = List::new();
154         list.push(1); list.push(2); list.push(3);
155
156         let mut iter = list.into_iter();
157         assert_eq!(iter.next(), Some(3));
158         assert_eq!(iter.next(), Some(2));
159         assert_eq!(iter.next(), Some(1));
160         assert_eq!(iter.next(), None);
161
162         println!("into_iter test Ok!");

```

```

163     }
164
165     fn iter_test() {
166         let mut list = List::new();
167         list.push(1); list.push(2); list.push(3);
168
169         let mut iter = list.iter();
170         assert_eq!(iter.next(), Some(&3));
171         assert_eq!(iter.next(), Some(&2));
172         assert_eq!(iter.next(), Some(&1));
173         assert_eq!(iter.next(), None);
174         println!("iter test Ok!");
175     }
176
177     fn iter_mut_test() {
178         let mut list = List::new();
179         list.push(1); list.push(2); list.push(3);
180
181         let mut iter = list.iter_mut();
182         assert_eq!(iter.next(), Some(&mut 3));
183         assert_eq!(iter.next(), Some(&mut 2));
184         assert_eq!(iter.next(), Some(&mut 1));
185         assert_eq!(iter.next(), None);
186         println!("iter_mut test Ok!");
187     }
188 }

```

4.6.3 LinkedList Stack

To implement a stack, we can also use a linked list, which is another type of linear data structure. In this case, we can use the head of the linked list to store the top element of the stack. As new items are added to the stack, they will be inserted at the head of the linked list. The implementation for this approach is shown below. Note that since the push and pop functions modify the linked list nodes, we use the take function to extract the node value.

```

1  // list_stack.rs
2
3  // Linked List Node
4  #[derive(Debug, Clone)]
5  struct Node<T> {
6      data: T,
7      next: Link<T>,
8  }
9
10 // Node Self-contained Reference
11 type Link<T> = Option<Box<Node<T>>>;
12
13 impl<T> Node<T> {
14     fn new(data: T) -> Self {
15         Self {
16             data: data,
17             next: None // No next link when initialized

```

```

18     }
19 }
20 }
21
22 // Linked list stack
23 #[derive(Debug, Clone)]
24 struct LStack<T> {
25     size: usize,
26     top: Link<T>, // Top controls the entire stack
27 }
28
29 impl<T: Clone> LStack<T> {
30     fn new() -> Self {
31         Self {
32             size: 0,
33             top: None
34         }
35     }
36
37     fn is_empty(&self) -> bool {
38         0 == self.size
39     }
40
41     fn len(&self) -> usize {
42         self.size
43     }
44
45     fn clear(&mut self) {
46         self.size = 0;
47         self.top = None;
48     }
49
50     // take out the node on the top, leaving an empty space
51     // that could be filled later
52     fn push(&mut self, val: T) {
53         let mut node = Node::new(val);
54         node.next = self.top.take();
55         self.top = Some(Box::new(node));
56         self.size += 1;
57     }
58
59     fn pop(&mut self) -> Option<T> {
60         self.top.take().map(|node| {
61             let node = *node;
62             self.top = node.next;
63             self.size -= 1;
64             node.data
65         })
66     }
67
68     // Return a reference to the data in the linked list stack
69     fn peek(&self) -> Option<&T> {

```

```

70         self.top.as_ref().map(|node| &node.data)
71     }
72
73     fn peek_mut(&mut self) -> Option<&mut T> {
74         self.top.as_deref_mut().map(|node| &mut node.data)
75     }
76
77     // into_iter: the linked list stack changes
78     // iter: the linked list stack remains unchanged
79     // iter_mut: the linked list stack remains unchanged
80     fn into_iter(self) -> IntoIter<T> {
81         IntoIter(self)
82     }
83
84     fn iter(&self) -> Iter<T> {
85         Iter { next: self.top.as_deref() }
86     }
87
88     fn iter_mut(&mut self) -> IterMut<T> {
89         IterMut { next: self.top.as_deref_mut() }
90     }
91 }
92
93 // Implement three iterations
94 struct IntoIter<T: Clone>(LStack<T>);
95 impl<T: Clone> Iterator for IntoIter<T> {
96     type Item = T;
97     fn next(&mut self) -> Option<Self::Item> {
98         self.0.pop()
99     }
100 }
101
102 struct Iter<'a, T: 'a> { next: Option<&'a Node<T>> }
103 impl<'a, T> Iterator for Iter<'a, T> {
104     type Item = &'a T;
105     fn next(&mut self) -> Option<Self::Item> {
106         self.next.map(|node| {
107             self.next = node.next.as_deref();
108             &node.data
109         })
110     }
111 }
112
113 struct IterMut<'a, T: 'a> { next: Option<&'a mut Node<T>> }
114 impl<'a, T> Iterator for IterMut<'a, T> {
115     type Item = &'a mut T;
116     fn next(&mut self) -> Option<Self::Item> {
117         self.next.take().map(|node| {
118             self.next = node.next.as_deref_mut();
119             &mut node.data
120         })
121     }

```

```

122 }
123
124 fn main() {
125     basic();
126     iter();
127
128     fn basic() {
129         let mut s = LStack::new();
130         s.push(1); s.push(2); s.push(3);
131
132         println!("empty: {:?}", s.is_empty());
133         println!("top: {:?}", size: {}, s.peek(), s.len());
134         println!("pop: {:?}", size: {}, s.pop(), s.len());
135
136         let peek_mut = s.peek_mut();
137         if let Some(data) = peek_mut {
138             *data = 4
139         }
140         println!("top {:?}", size {}, s.peek(), s.len());
141
142         println!("{:?}", s);
143         s.clear();
144         println!("{:?}", s);
145     }
146
147     fn iter() {
148         let mut s = LStack::new();
149         s.push(1); s.push(2); s.push(3);
150
151         let sum1 = s.iter().sum::<i32>();
152         let mut addend = 0;
153         for item in s.iter_mut() {
154             *item += 1;
155             addend += 1;
156         }
157         let sum2 = s.iter().sum::<i32>();
158         println!("{sum1} + {addend} = {sum2}");
159
160         assert_eq!(9, s.into_iter().sum::<i32>());
161     }
162 }

```

The output of the code will depend on how the stack is used and what functions are called.

```

empty: false
top: Some(3), size: 3
pop: Some(3), size: 2
top Some(4), size 2
LStack { size: 2, top: Some(Node { data: 4,
    next: Some(Node { data: 1, next: None }) }) }
LStack { size: 0, top: None }
6 + 3 = 9

```

4.7 Vec

In this chapter, we have demonstrated how the basic data type `Vec` can be used to implement various abstract data types, including stacks, queues, dequeues, and linked lists. `Vec` is a powerful yet simple data container that offers mechanisms for data collection and various operations, which is why we repeatedly use it as the underlying data structure for implementing other data types. It is similar to Python's `List` and is very convenient to use. However, not all programming languages include `Vec`, or not all data types may be suitable for it. In some cases, `Vec` or similar data containers must be implemented separately by programmers.

4.7.1 The Vec Abstract Data Type

`Vec` is a collection of items that maintain a relative position to other items. Here are the various operations of the `Vec` abstract data type:

- `new()`: Creates a new empty `Vec`, takes no arguments.
- `push(item)`: Adds the item to the end of the `Vec`, takes `item` as a parameter.
- `pop()`: Deletes the last item in the `Vec` and returns it, takes no arguments.
- `insert(pos, item)`: Inserts the item at position `pos` in the `Vec`, takes `pos` and `item` parameters.
- `remove(idx)`: Removes the item at index `idx` from the `Vec` and returns it, takes a parameter.
- `find(item)`: Checks if the item exists in the `Vec` and returns a boolean value, takes an item parameter.
- `is_empty()`: Checks if the `Vec` is empty and returns a boolean value, takes no arguments.
- `size()`: Calculates the number of items in the `Vec` and returns a `usize` integer, takes no arguments.
- `iter()`: Returns an immutable iteration form of the `Vec`.
- `iter_mut()`: Returns a mutable iteration form of the `Vec`, takes no arguments.
- `into_iter()`: Consumes the `Vec` and returns an iterable form of the `Vec`, takes no arguments.

Assuming `v` is an empty `Vec` that has been created, the table below shows the `Vec` after different operations, with the left side being the head.

Table 4.7: Operations on a `Vec`

operation	value	return
<code>v.is_empty()</code>	<code>[]</code>	<code>true</code>
<code>v.push(1)</code>	<code>[1]</code>	
<code>v.push(2)</code>	<code>[1,2]</code>	
<code>v.push(3)</code>	<code>[1,2,3]</code>	
<code>v.size()</code>	<code>[1,2,3]</code>	<code>3</code>
<code>v.pop()</code>	<code>[1,2]</code>	<code>3</code>
<code>v.push(5)</code>	<code>[1,2,5]</code>	
<code>v.find(4)</code>	<code>[1,2,5]</code>	<code>false</code>
<code>v.insert(0,8)</code>	<code>[8,1,2,5]</code>	
<code>v.pop()</code>	<code>[8,1,2]</code>	<code>5</code>
<code>v.remove(0)</code>	<code>[1,2]</code>	<code>8</code>
<code>v.size()</code>	<code>[1,2]</code>	<code>2</code>

Finally, it is worth noting that while `Vec` is a powerful data structure, it may not always be the best choice for a particular use case. Programmers should carefully consider the requirements of their project and choose the appropriate data structure accordingly.

4.7.2 Implementing a Vec in Rust

`Vec` is composed of linked list nodes, where each node contains a data item and an explicit reference to the next node. Given the location of the first node, subsequent items can be accessed by following the next link.

Since references play a crucial role in Vec, it is necessary to maintain a reference to the first node. The linked list is created with None indicating that the list does not reference any content, as shown in Figure (4.19). The head of the linked list refers to the first node of the list, which stores the address of the next node. It is important to note that the Vec itself does not contain any node objects but only a reference to the first node in the linked list structure.

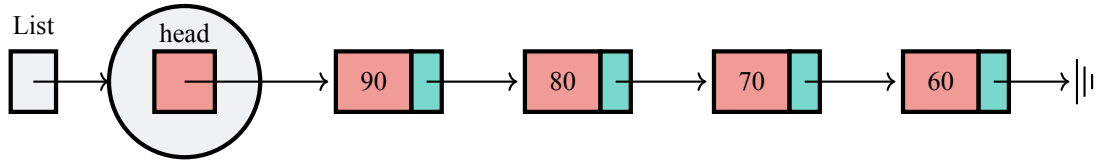


Figure 4.19: LinkedList Formed Vec

To add a new item to the linked list, the only entry point is through the head of the list. As all other nodes can only be accessed by following the next link from the first node, the most efficient way to add a new node is to add it to the head of the linked list. This approach adds the new item as the first element, with the existing items linked to it.

While the implementation of Vec presented here is unordered, it is possible to implement an ordered Vec using a data comparison function with total or partial orders. The following LVec provides only a portion of the functionality of the standard library Vec, and print_lvec is used to print its data items.

```

1 // lvec.rs
2
3 use std::fmt::Debug;
4
5 #[derive(Debug)]
6 struct Node<T> {
7     elem: T,
8     next: Link<T>,
9 }
10
11 type Link<T> = Option<Box<Node<T>>>;
12
13 impl<T> Node<T> {
14     fn new(elem: T) -> Self {
15         Self {
16             elem: elem,
17             next: None
18         }
19     }
20 }
21
22 // LinkedList Vec definition
23 #[derive(Debug)]
24 struct LVec<T> {
25     size: usize,
26     head: Link<T>,
27 }
28
29 impl<T: Copy + Debug> LVec<T> {
30     fn new() -> Self {
31         Self { size: 0, head: None }
32     }
33 }

```

```

32     }
33
34     fn is_empty(&self) -> bool {
35         0 == self.size
36     }
37
38     fn len(&self) -> usize {
39         self.size
40     }
41
42     fn clear(&mut self) {
43         self.size = 0;
44         self.head = None;
45     }
46
47     fn push(&mut self, elem: T) {
48         let node = Node::new(elem);
49         if self.is_empty() {
50             self.head = Some(Box::new(node));
51         } else {
52             let mut curr = self.head.as_mut().unwrap();
53
54             // find the last node in the list
55             for _i in 0..self.size-1 {
56                 curr = curr.next.as_mut().unwrap();
57             }
58
59             // insert the new data after the last node
60             curr.next = Some(Box::new(node));
61         }
62         self.size += 1;
63     }
64
65     // add a new LVec to the end of the stack
66     fn append(&mut self, other: &mut Self) {
67         while let Some(node) = other.head.as_mut().take() {
68             self.push(node.elem);
69             other.head = node.next.take();
70         }
71         other.clear();
72     }
73
74     fn insert(&mut self, mut index: usize, elem: T) {
75         if index >= self.size { index = self.size; }
76
77         // three cases for inserting a new node
78         let mut node = Node::new(elem);
79         if self.is_empty() { // LVec is empty
80             self.head = Some(Box::new(node));
81         } else if index == 0 { // insert at the beginning of
            the list
82             node.next = self.head.take();

```



```

83         self.head = Some(Box::new(node));
84     } else { // insert int the middle of the list
85         let mut curr = self.head.as_mut().unwrap();
86         for _i in 0..index - 1 { // find the right insert
            position
87             curr = curr.next.as_mut().unwrap();
88         }
89         node.next = curr.next.take();
90         curr.next = Some(Box::new(node));
91     }
92     self.size += 1;
93 }
94
95 fn pop(&mut self) -> Option<T> {
96     if self.size < 1 {
97         return None;
98     } else {
99         self.remove(self.size - 1)
100     }
101 }
102
103 fn remove(&mut self, index: usize) -> Option<T> {
104     if index >= self.size { return None; }
105
106     // two cases for deleting a node
107     let mut node;
108     if 0 == index {
109         node = self.head.take().unwrap();
110         self.head = node.next.take();
111     } else { // find the position which will be deleteed
        and arrange the links properly
112         let mut curr = self.head.as_mut().unwrap();
113         for _i in 0..index - 1 {
114             curr = curr.next.as_mut().unwrap();
115         }
116         node = curr.next.take().unwrap();
117         curr.next = node.next.take();
118     }
119     self.size -= 1;
120
121     Some(node.elem)
122 }
123
124 fn into_iter(self) -> IntoIter<T> {
125     IntoIter(self)
126 }
127
128 fn iter(&self) -> Iter<T> {
129     Iter { next: self.head.as_deref() }
130 }
131
132 fn iter_mut(&mut self) -> IterMut<T> {

```

```

133         IterMut { next: self.head.as_deref_mut() }
134     }
135
136     // print LVec
137     fn print_lvec(&self) {
138         if 0 == self.size {
139             println!("Empty lvec");
140         }
141         for item in self.iter() {
142             println!("{:?}", item);
143         }
144     }
145 }
146
147 // Implemion of three iterations
148 struct IntoIter<T: Copy + Debug>(LVec<T>);
149 impl<T: Copy + Debug> Iterator for IntoIter<T> {
150     type Item = T;
151     fn next(&mut self) -> Option<Self::Item> {
152         self.0.pop()
153     }
154 }
155
156 struct Iter<'a, T: 'a> { next: Option<&'a Node<T>> }
157 impl<'a, T> Iterator for Iter<'a, T> {
158     type Item = &'a T;
159     fn next(&mut self) -> Option<Self::Item> {
160         self.next.map(|node| {
161             self.next = node.next.as_deref();
162             &node.elem
163         })
164     }
165 }
166
167 struct IterMut<'a, T: 'a> { next: Option<&'a mut Node<T>> }
168 impl<'a, T> Iterator for IterMut<'a, T> {
169     type Item = &'a mut T;
170     fn next(&mut self) -> Option<Self::Item> {
171         self.next.take().map(|node| {
172             self.next = node.next.as_deref_mut();
173             &mut node.elem
174         })
175     }
176 }
177
178 fn main() {
179     basic();
180     iter();
181
182     fn basic() {
183         let mut lvec1: LVec<i32> = LVec::new();
184         lvec1.push(10); lvec1.push(11);

```

```

185         lvec1.push(12); lvec1.push(13);
186         lvec1.insert(0,9);
187
188         lvec1.print_lvec();
189
190         let mut lvec2: LVec<i32> = LVec::new();
191         lvec2.insert(0, 8);
192         lvec2.append(&mut lvec1);
193
194         println!("len: {}", lvec2.len());
195         println!("pop {:?}", lvec2.pop().unwrap());
196         println!("remove {:?}", lvec2.remove(0).unwrap());
197
198         lvec2.print_lvec();
199         lvec2.clear();
200         lvec2.print_lvec();
201     }
202
203     fn iter() {
204         let mut lvec: LVec<i32> = LVec::new();
205         lvec.push(10); lvec.push(11);
206         lvec.push(12); lvec.push(13);
207
208         let sum1 = lvec.iter().sum::<i32>();
209         let mut addend = 0;
210         for item in lvec.iter_mut() {
211             *item += 1;
212             addend += 1;
213         }
214         let sum2 = lvec.iter().sum::<i32>();
215         println!("{sum1} + {addend} = {sum2}");
216
217         assert_eq!(50, lvec.into_iter().sum::<i32>());
218     }
219 }

```

Here is the execution result.

```

9
10
11
12
13
len: 6
pop 13
remove 8
9
10
11
12
Empty lvec
46 + 4 = 50

```

It's important to note that `LVec` is a linked list with n nodes, and any method that requires traversing nodes, such as `insert`, `push`, `pop`, `remove`, etc., has a time complexity of $O(n)$. Although on average it may only require traversing half as many nodes, in the worst case, every node in the list must be processed.

4.8 Summary

This chapter focuses on several linear data structures, including `stack`, `queue`, `double-ended queue`, `LinkedList`, and `Vec`.

A `stack` is a type of data structure that maintains last-in-first-out (LIFO) ordering and has basic operations such as `push`, `pop`, and `is_empty`. Stacks are useful for designing algorithms that compute parsed expressions and can provide inversion properties that are useful in implementing operating system function calls, website save functions, and more. Prefix, infix, and postfix expressions can all be handled with a `stack`, but computers do not typically use infix expressions.

A `queue` is a simple data structure that maintains a first-in-first-out (FIFO) sorting property and has basic operations such as `enqueue`, `dequeue`, and `is_empty`. Queues are very useful in scheduling system tasks and can help build timed task emulations.

A `double-ended queue` (`Deque`) is a data structure that allows a mixed behavior of `stack` and `queue`. Its basic operations include `is_empty`, `add_front`, `add_rear`, `remove_front`, and `remove_rear`. Although a `double-ended queue` can be used as a `stack` or a `queue`, it is recommended to use it only as a `double-ended queue`.

A `LinkedList` is a collection of items, each of which is stored in a relative position in the list. The implementation of a linked list itself maintains logical order and does not need to be stored in physical order. Modifying the head of a linked list is a special case.

`Vec` is a data container that comes with Rust, and the default implementation uses dynamic arrays. In this chapter, however, we use a linked list for this purpose.

Chapter 5

Recursion

5.1 Objectives

- Understanding simple recursive solutions
- Learning how to write programs using recursion
- Understanding and applying the three laws of recursion
- Understanding recursion as a form of iteration
- Formulating problems into recursive solutions
- Understanding how computers implement recursion

5.2 What is Recursion?

Recursion is a problem-solving method that involves breaking down a complex problem into smaller sub-problems until a simple base case is reached. By solving the base case and combining the results, the solution to the original problem can be obtained. Recursion is particularly useful for solving difficult problems elegantly.

For instance, consider the task of computing the sum of an integer array like [2, 1, 7, 4, 5]. The most straightforward way to do it is by using an accumulator to add each value iteratively.

```
1 // iterative sum
2
3 fn nums_sum(nums: Vec<i32>) -> i32 {
4     // use sum as accumulator
5     let mut sum = 0;
6
7     for num in nums {
8         sum += num;
9     }
10
11     sum
12 }
```

However, if a programming language lacks for or while loops, the above sum code won't work. In such cases, one can still compute the sum by using recursion to solve smaller problems. To solve the sum problem recursively, you can break down the array into smaller additions, which can be constructed using the basic problem of addition. By recursively solving these smaller problems, you can eventually solve the original sum problem without loops.

The process of constructing small additions can be done using elementary school-level knowledge of constructing fully parenthesized expressions, of which there are of course two forms as shown below.

$$\begin{aligned}
 2 + 1 + 7 + 4 + 5 &= 2 + 1 + 7 + 4 + 5 \\
 sum &= (((2 + 1) + 7) + 4) + 5 \\
 sum &= (((3 + 7) + 4) + 5) \\
 sum &= (10 + 4) + 5 \\
 sum &= (14 + 5) \\
 sum &= 19 \\
 &= 2 + 1 + 7 + 4 + 5 \\
 sum &= (2 + (1 + (7 + (4 + 5)))) \\
 sum &= (2 + (1 + (7 + 9))) \\
 sum &= (2 + (1 + 16)) \\
 sum &= (2 + 17) \\
 sum &= 19
 \end{aligned} \tag{5.1}$$

Overall, recursion provides an elegant and effective solution to seemingly difficult problems, allowing for efficient computation and solving programming challenges even in languages with limited capabilities.

The given expression can be correctly parenthesized using either form of parentheses on the right-hand side. By following the rule of parentheses precedence and internal priority, we can treat the parenthesized expression as a sequence of small additions. The pattern of both parts and the entire expression is entirely recursive, and we can simulate it without using While or For loops.

If we observe the calculation starting with "sum" and read it from bottom to top, we can see that the first term is 19, followed by (2 + 17), and then (2 + (1 + 16)). The total sum is the sum of the first term and the remaining terms on the right-hand side. We can further decompose it into the sum of its first term and the remaining terms on the right-hand side. Therefore, we can express this mathematically as follows:

$$Sum(nums) = First(nums) + Sum(restR(nums)) \tag{5.2}$$

This is the calculation method for the second form of parenthesized expressions. Similarly, we also have a calculation method for the first form of parenthesized expressions.

$$Sum(nums) = Last(nums) + Sum(restL(nums)) \tag{5.3}$$

The equation defines various functions such as First(nums), restR(nums), Last(nums), and restL(nums) that return the required elements from the array.

We can implement two recursive methods to calculate expressions in Rust. The `nums_sum1` function sums the remaining terms after `nums[0]`, while the `nums_sum2` function sums the last term and all the terms before it. Both implementations have the same time and space complexities and do not differ from each other.

```

1 // nums_sum12.rs
2
3 // Form : Sum(nums) = First(nums) + Sum(restR(nums))
4 fn nums_sum1(nums: &[i32]) -> i32 {
5     if 1 == nums.len() {
6         nums[0]
7     } else {
8         let first = nums[0];
9         first + nums_sum1(&nums[1..])
10    }

```

```

11 }
12
13 // Form : Sum(nums) = Last(nums) + Sum(restL(nums))
14 fn nums_sum2(nums: &[i32]) -> i32 {
15     if 1 == nums.len() {
16         nums[0]
17     } else {
18         let last = nums[nums.len() - 1];
19         nums_sum2(&nums[..nums.len() - 1]) + last
20     }
21 }
22
23 fn main() {
24     let nums = [2,1,7,4,5];
25     let sum1 = nums_sum1(&nums);
26     let sum2 = nums_sum2(&nums);
27     println!("sum1 = {sum1}, sum2 = {sum2}");
28     // sum1 = 19, sum2 = 19
29
30     let nums = [-1,7,1,2,5,4,10,100];
31     let sum1 = nums_sum1(&nums);
32     let sum2 = nums_sum2(&nums);
33     println!("sum1 = {sum1}, sum2 = {sum2}");
34     // sum1 = 128, sum2 = 128
35 }

```

The crucial part of the code is the if and else statements and their respective forms. The condition of `1 == nums.len()` is crucial as it marks the turning point of the function. At this point, the numerical value is directly returned without any further mathematical calculation. In the else statement, the function calls itself, achieving the effect of solving parentheses layer by layer and calculating their values. This is the essence of recursion, where a function calls itself until it reaches a base case.

5.2.1 The Three Laws of Recursion

By analyzing the above code, we can see that all recursive algorithms must follow these three basic laws:

- 1 Recursive algorithms must have a base case
- 2 Recursive algorithms must move towards the base case
- 3 Recursive algorithms must call themselves in a recursive manner

The first law is the base case, which is when the algorithm stops, in this case, `1 == nums.len()`. The second law involves the decomposition of the problem, where we return the number in the else clause and then calculate it again using the set of elements except for the returned number. This reduces the number of elements in the original array `nums` that need to be calculated. It is evident that the else clause moves towards the base case (`nums.len() == 1`), which occurs when `nums.len() > 1`. The third law involves calling itself, which is implemented in the else clause. It is important to note that calling itself is not a loop, and there are no while or for statements here. In summary, our recursive sum algorithm satisfies these three laws.

The recursive function call process can be visualized using a series of boxes representing the function call graph. Each recursion solves a small problem until the problem reaches the base case, where it can no longer be further decomposed. Finally, these intermediate calculated values are used to solve the larger problem.

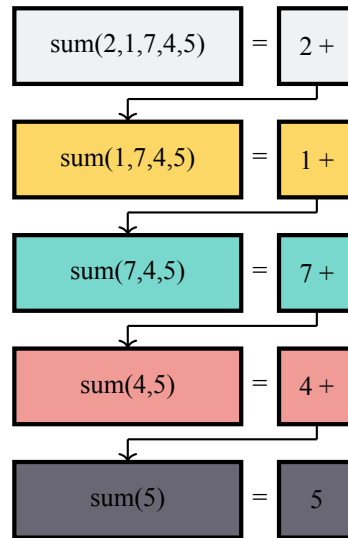


Figure 5.1: caculation by iteration

5.2.2 Converting an Integer to a String in Any Base

In the previous section, we discussed implementing algorithms for converting integers into binary and hexadecimal strings using stacks. However, recursive methods can also be used for base conversion. To design a recursive algorithm for base conversion, we must adhere to the three laws of recursion.

- 1 The recursive algorithm must have a base case
- 2 The recursive algorithm must move towards the base case
- 3 The recursive algorithm must call itself

The algorithm for base conversion can be derived as follows:

- 1 Simplify the original number into a series of individual digits
- 2 Use a lookup method to convert each digit into a character
- 3 Concatenate the individual characters to form the final result

To move towards the base case, we use division to change the number state. We divide the number by the base and obtain a remainder and a quotient. If the quotient is less than the base, we stop the operation and return the result. For instance, consider the decimal number 996 and converting it into a base-10 string. We divide 996 by 10, and the remainder is 6, and the quotient is 99. Since the remainder is less than 10, we can obtain the character '6'. The quotient 99 is less than 996, which moves us closer to the base case. We then recursively call the function with 99 as the input. We divide 99 by 10, and the remainder is 9, and the quotient is 9. Since the remainder is less than 10, we can obtain the character '9'. Finally, we get the decimal string '996'. Once we can solve base-10 conversion, we can easily handle base-2 to base-16 conversion.

Here's an algorithm for converting integers to any base (base-2 to base-16) string. The variable BASESTR holds the character representation for different digits, with digits greater than 10 represented by characters A-F.

```

1 // num2str_rec.rs
2
3 //Character table for various digits
4 const BASESTR: [&str; 16] = ["0","1","2","3","4","5","6","7",
5                               "8","9","A","B","C","D","E","F"];
  
```



```

6 fn num2str_rec(num: i32, base: i32) -> String {
7     if num < base {
8         BASESTR[num as usize].to_string()
9     } else {
10        // Append remainder to the end of the string
11        num2str_rec(num/base, base) +
12        BASESTR[(num % base) as usize]
13    }
14 }
15
16 fn main() {
17     let num = 100;
18     let sb = num2str_rec(num,2); // sb = str_binary
19     let so = num2str_rec(num,8); // so = str_octal
20     let sh = num2str_rec(num,16); // sh = str_hexdecimal
21     println!("{num} = b{sb}, o{so}, x{sh}");
22     // 100 = b1100100, o144, x64
23
24     let num = 1000;
25     let so = num2str_rec(num,8);
26     let sh = num2str_rec(num,16);
27     println!("{num} = o{so}, x{sh}");
28     // 1000 = o1750, x3E8
29 }

```

In the previous section, we utilized a stack to convert numbers to any base. Now, we still achieve similar functionality using recursion, indicating a relationship between stacks and recursion. Recursion can be viewed as a stack, but the stack is implicitly called by the compiler. The code only uses recursion, but the compiler employs a stack to save data. If we were to implement the following recursive code using a stack, the structure would be very similar.

```

1 // num2str_stk.rs
2
3 fn num2str_stk(mut num: i32, base: i32) -> String {
4     let digits: [&str; 16] = ["0","1","2","3","4","5","6","7",
5                                "8","9","A","B","C","D","E","F"];
6
7     let mut rem_stack = Stack::new();
8     while num > 0 {
9         if num < base {
10            // Push directly to stack if not exceeding the base
11            rem_stack.push(num);
12        } else {
13            // Push remainder to stack if exceeding the base
14            rem_stack.push(num % base);
15        }
16        num /= base;
17    }
18
19    // Pop remainder from stack and form a string
20    let mut numstr = "".to_string();
21    while !rem_stack.is_empty() {
22        numstr += digits[rem_stack.pop().unwrap() as usize];

```

```

23     }
24
25     numstr
26 }
27
28 fn main() {
29     let num = 100;
30     let sb = num2str_stk(100, 2);
31     let so = num2str_stk(100, 8);
32     let sh = num2str_stk(100, 16);
33     println!("{num} = b{sb}, o{so}, x{sh}");
34     // 100 = b1100100, o144, x64
35 }

```

5.2.3 Tower of Hanoi

The Tower of Hanoi puzzle was invented by French mathematician Edouard Lucas in 1883. The puzzle involves moving 64 gold discs of decreasing size from one of three poles to another, one disc at a time, with the constraint that a larger disc cannot be placed on a smaller one. The inspiration for this puzzle came from a legend about a puzzle given to young priests by the abbot of an Indian Hindu temple. In reality, moving all 64 discs would take 5.85 trillion years, longer than the known age of the universe.

Figure (5.2) illustrates the process of moving disks from the first peg to the third peg, which resembles a stack. If you haven't played this before, you can try it now. You don't need disks, a pile of bricks, books, or paper will do, and you don't need 64, 10 is enough. See if it really takes that long.

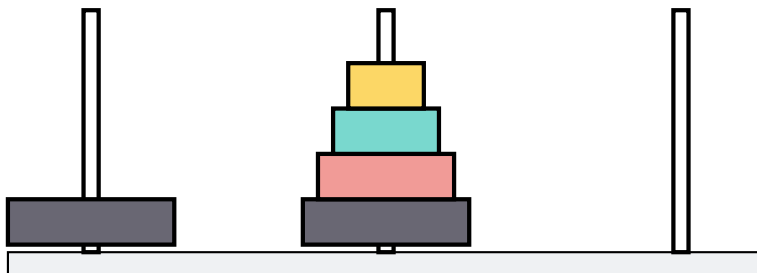


Figure 5.2: Hanoi

To solve this problem recursively, we need to determine the base case. Suppose there is a tower of Hanoi with three pegs: left, middle, and right, and five disks on the left peg. We can assume that we know how to move one disk to the right. This is the base case. If we know how to move $n-1$ disks to the middle peg, we can easily move the bottom disk to the right peg and then move the $n-1$ disks from the middle peg to the right peg. If we don't know how to move $n-1$ disks to the middle peg, we can assume that we know how to move $n-2$ disks to the middle peg, then move the $n-1$ th disk to the right peg, and then move the $n-2$ disks from the middle peg to the top of the $n-1$ th disk on the right peg. This process is an abstraction of the disk movement process.

In summary, we can organize the above operation process into the following algorithm to solve the Tower of Hanoi puzzle recursively.

- 1 Move height-1 disks to the middle peg using the target peg as a helper.
- 2 Move the last disk to the target peg.
- 3 Move height-1 disks from the middle peg to the target peg using the starting peg as a helper.

To solve the Towers of Hanoi problem, the recursive three laws must be followed to ensure that larger discs remain at the bottom of the stack. The base case is when there is only one disc, which can be moved to its final destination. The algorithm then reduces the height of the Towers of Hanoi through steps 1 and 3, moving towards the base case. The solution can be implemented in a few lines of Rust code using recursion, as shown below.

```

1 // hanoi.rs
2
3 // p: pole
4 fn hanoi(height:u32, src_p:&str, des_p:&str, mid_p:&str) {
5     if height >= 1 {
6         hanoi(height - 1, src_p, mid_p, des_p);
7         println!("move disk[{height}] from
8                 {src_p} to {des_p}");
9         hanoi(height - 1, mid_p, des_p, src_p);
10    }
11 }
12
13 fn main() {
14     hanoi(1, "A", "B", "C");
15     hanoi(2, "A", "B", "C");
16     hanoi(3, "A", "B", "C");
17     hanoi(4, "A", "B", "C");
18     hanoi(5, "A", "B", "C");
19     hanoi(6, "A", "B", "C");
20 }
```

To simulate the problem, one can use three pens and paper to move the discs according to the output of the hanoi function for heights 1, 2, 3, and 4.

5.3 Tail Recursion

The aforementioned recursive computation is called regular recursion, which involves saving values during the computation process. This can lead to stack overflow if there are too many recursive calls and memory becomes limited. To avoid this, tail recursion can be used. Tail recursion processes the current result of the operation as a parameter for the next recursive function call, which reduces the memory consumption on the stack and makes the code more concise.

Tail recursion optimization is particularly useful when dealing with subproblems that need to be summed. It reduces the problem size by calculating the sum of the parameters beforehand, which optimizes the algorithm. However, tail-recursive code may be more difficult to understand since the results of subproblems are directly used as parameters for the next recursive call. The regular recursive version of the code for summing the numbers can be converted into a more concise tail-recursive form.

```

1 // nums_sum34.rs
2
3 fn nums_sum3(sum: i32, nums: &[i32]) -> i32 {
4     if 1 == nums.len() {
5         sum + nums[0]
6     } else {
7         nums_sum3(sum + nums[0], &nums[1..])
8     }
9 }
```

```

9  }
10
11 fn nums_sum4(sum: i32, nums: &[i32]) -> i32 {
12     if 1 == nums.len() {
13         sum + nums[0]
14     } else {
15         nums_sum4(sum + nums[nums.len() - 1],
16                   &nums[..nums.len() - 1])
17     }
18 }
19
20 fn main() {
21     let nums = [2,1,7,4,5];
22     let sum1 = nums_sum3(0, &nums);
23     let sum2 = nums_sum4(0, &nums);
24     println!("sum1 is {sum1}, sum2 is {sum2}");
25     // sum1 is 19, sum2 is 19
26 }

```

Ultimately, the implementation of recursive programs depends on the programmer's ability to write clear and efficient code. If tail recursion does not cause stack overflow and is easy to understand, it can be a useful tool in algorithm optimization.

5.3.1 Recursion VS Iteration

To compute the array [2, 1, 7, 4, 5], we employed three methods: loop iteration, recursion, and tail recursion. These methods demonstrate that recursion and iteration can accomplish the same goal. So, what is the relationship between recursion and iteration?

- Recursion refers to a process of repeating a transaction in a self-similar way. In mathematics and computer science, it involves using the function itself in its definition. When recursion is unfolded, it creates a tree-like structure. This can be thought of as a process of repetitive recursion and backtracking. Once the recursion reaches the bottom, backtracking begins, which is equivalent to a depth-first traversal of the tree.

- On the other hand, iteration involves repeating a feedback process, where the outcome of each iteration becomes the starting point for the next iteration. Iteration is a loop structure, starting from the initial state, where each iteration traverses the loop and updates the state. Multiple iterations continue until the end state is reached.

While all iterations can be converted into recursion, converting recursion into iteration is not always possible because changing a tree into a loop is not always feasible. Therefore, it is essential to choose the appropriate method for a given problem, taking into account its complexity, readability, and efficiency.

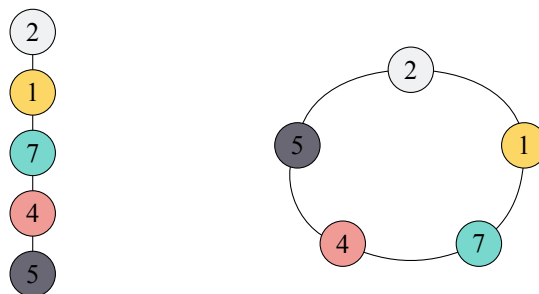


Figure 5.3: Recursion VS Iteration

5.4 Dynamic Programming

In computer science, optimization problems that involve finding the maximum or minimum value of a function are common. These problems are crucial for achieving carbon neutrality, saving energy, and improving the passenger experience. This section aims to demonstrate different solutions to optimization problems and highlight dynamic programming as an effective approach for solving such problems.

A typical example of an optimization problem is the "minimum number of bills/coins to make change" problem, commonly seen in subway ticketing and vending machines. The objective is to minimize the number of bills/coins returned for a single transaction. For example, returning two bills (a five and a one) for a six dollars transaction, instead of six one-dollar bills.

The problem is how to plan the return of different denominations of bills/coins from a total task, such as making change for six dollars. The most intuitive approach is the greedy method, which starts with the largest denomination of bills/coins and uses as many of them as possible, then moves on to the next smaller denomination and repeats the process until the change is complete.

However, the greedy method may not always yield the optimal solution. For instance, if a country has a complex currency system that includes not only the usual 1, 5, 10, and 25 dollar bills/coins, but also a 22 dollar bill/coin, the greedy method cannot find the best solution for making change for 66 dollars. Even with the addition of the 22 dollars bill/coin, the greedy method still finds a solution, but it requires five bills (two 25 dollar bills, one 10 dollar bill, one 5 dollar bill, and one 1 dollar bill), while the optimal answer is three 22 dollar bills.

To solve the change-making problem, recursion is an effective method. The first step is to identify the basic problem, which is to make change for one bill/coin with an unspecified denomination but equal to the total amount needed. This is because making change with only one bill/coin represents the minimum number of bills/coins required except when the amount is zero. Assuming US dollars bills with denominations of 1, 5, 10, 20, and 50 dollar are used, the number of bills needed to make change for the original amount can be calculated by adding one to the number of bills needed for the remaining amount after subtracting the denomination from the total amount. Therefore, the formula for calculating the number of bills needed is straightforward.

$$\text{numBills}(\text{amount}) = \begin{cases} 1 + \text{numBills}(\text{amount} - 1) \\ 1 + \text{numBills}(\text{amount} - 5) \\ 1 + \text{numBills}(\text{amount} - 10) \\ 1 + \text{numBills}(\text{amount} - 20) \\ 1 + \text{numBills}(\text{amount} - 50) \end{cases} \quad (5.4)$$

To implement this approach, the `numBills` function is used to calculate the number of bills required for a given amount of change. The function considers five cases, where the change can be made up of different denominations of bills. The algorithm checks if the change amount is equal to a denomination of bill on line 8, which serves as a base case. If not, the function recursively calls itself with a reduced change amount and the next smallest denomination of bill. The count of bills used is incremented before each recursive call to keep track of the total number of bills used in the final solution

```

1 // rec_mc1.rs
2
3 fn rec_mc1(cashes: &[u32], amount: u32) -> u32 {
4     // Calculates the minimum number of coins required for
5     // change when using only 1-yuan bills.
6     let mut min_cashes = amount;
7
8     if cashes.contains(&amount) {
9         return 1;
10    } else {

```

```

11      // Extracts valid denominations (denominations must
12      // be smaller than the change amount).
13      for c in cashes.iter()
14          .filter(|&c| c <= amount)
15          .collect::<Vec<u32>>() {
16          // Subtracts c from amount to indicate that a bill
17          // of denomination c has been used,
18          // and increments num_cashes by 1.
19          let num_cashes = 1 + rec_mc1(&cashes, amount - c);
20
21          // If num_cashes is smaller than min_cashes,
22          // update min_cashes.
23          if num_cashes < min_cashes {
24              min_cashes = num_cashes;
25          }
26      }
27  }
28
29  min_cashes
30 }
31
32 fn main() {
33     // cashes stores denominations of bills.
34     let cashes = [1,5,10,20,50];
35     let amount = 31u32;
36     let cashes_num = rec_mc1(&cashes, amount);
37     println!("need refund {cashes_num} cashes");
38     // need refund 3 cashes
39 }

```

To avoid lengthy wait times for the program to return a result, it is important to optimize the code by reducing redundant calculations. In the given code, changing the amount to 90 results in a long wait time because the program needs to make a large number of recursive calls to find the combination of three bills: [50, 20, 20].

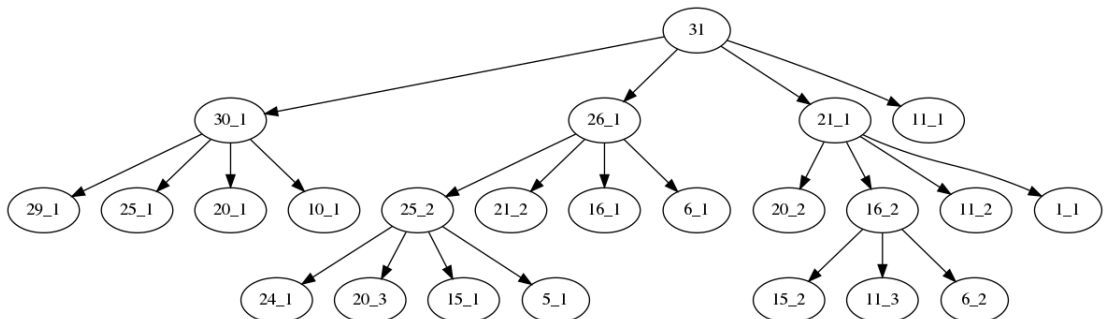


Figure 5.4: solve the task by recursion

To address this issue, one solution is to store previously calculated results to avoid redundant calculations. One approach is to store the current minimum number of bills in a list and check this list before calculating a new minimum value. If the result already exists, use the stored value instead of recalculating it. This technique is an example of trading space for time in algorithm design.

```

1 // rc_mc2.rs
2
3 fn rec_mc2(cashes: &[u32], amount: u32,
4           min_cashes: &mut [u32]) -> u32
5 {
6     // Calculates the minimum number of coins required for
7     // change when using only 1-yuan bills.
8     let mut min_cashe_num = amount;
9
10    if cashes.contains(&amount) {
11        // Collects denominations that match the
12        // current change value.
13        min_cashes[amount as usize] = 1;
14        return 1;
15    } else if min_cashes[amount as usize] > 0 {
16        // If the change amount already has the minimum number
17        // of coins required for change, return directly.
18        return min_cashes[amount as usize];
19    } else {
20        for c in cashes.iter()
21            .filter(|&&c| c <= amount)
22            .collect::<Vec<&u32>>() {
23
24            let cashe_num = 1 + rec_mc2(cashes,
25                                     amount - c,
26                                     min_cashes);
27
28            // Updates the minimum number of coins
29            // required for change.
30            if cashe_num < min_cashe_num {
31                min_cashe_num = cashe_num;
32                min_cashes[amount as usize] = min_cashe_num;
33            }
34        }
35    }
36
37    min_cashe_num
38 }
39
40 fn main() {
41     let amount = 90u32;
42     let cashes: [u32; 5] = [1,5,10,20,50];
43     let mut min_cashes: [u32; 91] = [0; 91];
44     let cashe_num = rec_mc2(&cashes, amount, &mut min_cashes);
45     println!("need refund {cashe_num} cashes");
46     // need refund 3 cashes
47 }

```

The `rec_mc2` is less time-consuming because it uses the variable `min_cashes` to store intermediate values. It is important to note that although this section discusses dynamic programming, both programs presented are recursive rather than dynamic programming. The second program only saves intermediate values during recursion, using a memory or cache technique to reduce the time required for calculations.

5.4.1 What is Dynamic Programming?

Dynamic programming(DP) is a mathematical technique for solving optimization problems in decision-making processes. The change-making problem described above is an optimization problem that seeks to minimize the number of bills/coins used, and this is the optimization objective.

The key to dynamic programming is identifying whether a problem can be solved by this technique and expressing the problem as a state that can be transitioned between. The state is critical to the calculation and is used to derive the next step in the problem-solving process.

Unlike the previous greedy algorithm, which goes from large to small denominations, dynamic programming assumes that the current best result has been obtained and deduces the next step based on this result. The technique decomposes a large problem into smaller sub-problems and deduces from small to large problems. The intermediate values of the deducing process are cached, and this process is known as state transition.

For the change-making problem, three parameters are required for dynamic programming: a list of available bills, the amount to be changed, and a list that contains the minimum number of bills needed for each amount. The algorithm saves the calculated values to the list, starting from the smallest amount and gradually building up to the desired change amount. The implementation of the algorithm uses iteration.

By using dynamic programming, redundant calculations can be avoided, and a significant improvement in efficiency can be achieved. The program caches previously calculated results, which reduces the workload and processing time.

```

1 // dp_rec_mc.rs
2
3 fn dp_rec_mc(cashes: &[u32], amount: u32,
4             min_cashes: &mut [u32]) -> u32 {
5     // Collect the minimum number of coins required to
6     // make change from 1 to amount, and then reconstructs
7     for denm in 1..=amount {
8         let mut min_cashe_num = denm;
9         for c in cashes.iter()
10             .filter(|&c| c <= denm)
11             .collect::<Vec<&u32>>() {
12             let index = (denm - c) as usize;
13             let cashe_num = 1 + min_cashes[index];
14             if cashe_num < min_cashe_num {
15                 min_cashe_num = cashe_num;
16             }
17         }
18         min_cashes[denm as usize] = min_cashe_num;
19     }
20     // directly return the result
21     min_cashes[amount as usize]
22 }
23
24 fn main() {
25     let amount = 90u32;
26     let cashes = [1,5,10,20,50];
27     let mut min_cashes: [u32; 91] = [0; 91];
28     let cash_num = dp_rec_mc(&cashes,amount,&mut min_cashes);
29     println!("Refund for ${amount} need {cash_num} cashes");
30     // Refund for $90 need 3 cashes
31 }

```


The iterative dynamic programming code is much more concise than the previous recursive versions and reduces the use of the stack. However, it is important to note that just because a problem can be solved using recursion does not mean it is the best solution.

While the dynamic programming algorithm finds the minimum number of bills required, it does not indicate which denominations of bills are used. To obtain this information, a table, `cashes_used`, can be added to record the denominations and quantities of bills used. The algorithm can be extended by adding the denomination of the last bill used for each amount to the `cashes_used` table and then continuously finding the last bill used for the previous amount until the end.

```

1 // dp_rc_mc_show.rs
2
3 // The algorithm uses cashes_used to collect
4 // the denominations of coins used,
5 fn dp_rec_mc_show(cashes: &[u32], amount: u32,
6                   min_cashes: &mut [u32],
7                   cashes_used: &mut [u32]) -> u32 {
8     for denm in 1..=amount {
9         let mut min_cashe_num = denm ;
10        // With a minimum denomination of 1 yuan
11        let mut used_cashe = 1;
12        for c in cashes.iter()
13            .filter(|&c| *c <= denm)
14            .collect::<Vec<&u32>>() {
15            let index = (denm - c) as usize;
16            let cashe_num = 1 + min_cashes[index];
17            if cashe_num < min_cashe_num {
18                min_cashe_num = cashe_num;
19                used_cashe = *c;
20            }
21        }
22        // update the minimum number of coins
23        // required for each amount
24        min_cashes[denm as usize] = min_cashe_num;
25        cashes_used[denm as usize] = used_cashe;
26    }
27
28    min_cashes[amount as usize]
29 }
30
31 // prints the denominations of coins used
32 fn print_cashes(cashes_used: &[u32], mut amount: u32) {
33     while amount > 0 {
34         let curr = cashes_used[amount as usize];
35         println!("{}", curr);
36         amount -= curr;
37     }
38 }
39
40 fn main() {
41     let amount = 81u32; let cashes = [1,5,10,20,50];
42     let mut min_cashes: [u32; 82] = [0; 82];
43     let mut cashes_used: [u32; 82] = [0; 82];

```

```

44     let cs_num = dp_rec_mc_show(&cashs, amount,
45                                &mut min_cashes,
46                                &mut cashs_used);
47     println!("Refund for ${amount} need {cs_num} cashs:");
48     print_cashes(&cashs_used, amount);
49 }

```

For instance, when making change for 90 dollars, only three bills ([50, 20, 20]) are needed.

```

Refund for $90 requires 3 cashs:
$20
$20
$50

```

5.4.2 Dynamic Programming VS Recursion

Recursion and dynamic programming are two different techniques used to solve problems. Recursion breaks down a large problem into smaller ones and solves them by calling itself, while dynamic programming solves large problems using solutions to smaller problems. Recursion can be memory-intensive and lead to stack overflow, while dynamic programming requires determining transition rules and initial conditions, but has simple code and a steep learning curve. Although both techniques can solve the same problems, such as the Fibonacci sequence problem.

```

1  // fibnacci_dp_rec.rs
2
3  fn fibnacci_dp(n: u32) -> u32 {
4      // Using only two positions to store values can save memory
5      let mut dp = [1, 1];
6      for i in 2..=n {
7          let idx1 = (i % 2) as usize;
8          let idx2 = ((i - 1) % 2) as usize;
9          let idx3 = ((i - 2) % 2) as usize;
10         dp[idx1] = dp[idx2] + dp[idx3];
11     }
12
13     dp[(((n-1) % 2) as usize)]
14 }
15
16 fn fibnacci_rec(n: u32) -> u32 {
17     if n == 1 || n == 2 {
18         return 1;
19     } else {
20         fibnacci_rec(n-1) + fibnacci_rec(n-2)
21     }
22 }
23
24 fn main() {
25     println!("fib(10): {}", fibnacci_dp(10));
26     println!("fib(10): {}", fibnacci_rec(10));
27     // fib(10): 55
28     // fib(10): 55
29 }

```

It is important to note that a problem that can be solved with dynamic programming may not necessarily be solvable with recursion as they have different requirements.

5.5 Summary

In this chapter, we covered both recursive and iterative algorithms. Recursive algorithms must satisfy three laws, and while recursion can sometimes replace iteration, it is not always the optimal solution. Recursion can be a natural way to express a problem, but tail recursion is an optimization technique that can reduce stack usage. Dynamic programming is a useful approach for solving optimization problems by breaking down large problems into smaller ones and gradually constructing larger solutions. While recursion solves problems by breaking them down, dynamic programming works by building up from small to large problems.

Chapter 6

Searching

6.1 Objectives

- Be able to implement sequential search and binary search algorithms.
- Understand the concept of using hash tables as a search technique.
- Use Vec to implement a HashMap data structure.

6.2 What is Searching?

In this chapter, we will use the data structures we have previously implemented (stacks, queues, and linked lists) to solve practical search and sort problems. Search and sort are fundamental tasks in computer science, and many software applications and algorithms are developed around them. You are likely familiar with the search function in various software applications you have used, such as finding specific characters in a Word document or using the search bar in Google browser to find information. Search and lookup mean the same thing, and this book does not distinguish between the two.

Searching is the process of finding a specific item within a set of items. Search algorithms typically return either true or false to indicate whether the item exists in the set, and sometimes they also return the position of the item. In Rust, we can use the `contains()` function to query whether an item is in a set. This function checks whether the data contains a certain value and performs a search for us.

```
1 fn main() {
2     let data = vec![1,2,3,4,5];
3     if data.contains(&3) {
4         println!("Yes");
5     } else {
6         println!("No");
7     }
8 }
```

While `contains()` is a simple search algorithm, there are many different search methods available, including sequential search, binary search, and hash search. We will explore how these different algorithms work and what their complexities are.

6.3 The Sequential Search

When data items are stored in collections such as Vec, arrays, or slices, they have a linear relationship since each item is stored at a position relative to the other items. This means that the index values of the

data items in slices are ordered, and can be accessed in order, making these data structures linear as well. Similarly, the stacks, queues, and linked lists that we have previously studied are also linear. Based on this same linear logic inherent in the physical world, a natural search technique is linear search, which is also known as sequential search.

Linear search works by starting at the first item in the slice and moving from one item to another in order until the target item is found or the entire slice is traversed. If the item being searched for is not found after traversing the entire slice, it means that the item does not exist. The following figure illustrates how this search works.

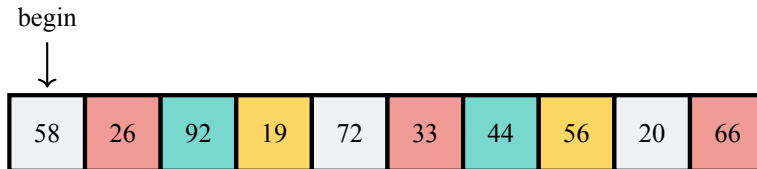


Figure 6.1: Sequential search

6.3.1 Implementing an Sequential Search in Rust

The following Rust code implements sequential search, and it is straightforward to understand. The program takes a slice of integers and a target integer as input. It iterates through each element in the slice and compares it with the target value. If the value is found, the function returns a boolean value: `true`. Otherwise, it returns `false`.

```
1 // sequential_search.rs
2
3 fn sequential_search(nums: &[i32], num: i32) -> bool {
4     let mut pos = 0;
5     let mut found = false;
6
7     // Continue looping if pos is within the index range and
8     // the item has not been found.
9     while pos < nums.len() && !found {
10         if num == nums[pos] {
11             found = true;
12         } else {
13             pos += 1;
14         }
15     }
16
17     found
18 }
```

Here is an example of sequential search execution.

```
1 fn main() {
2     let num = 8;
3     let nums = [9,3,7,4,1,6,2,8,5];
4     let found = sequential_search(&nums, num);
5     println!("nums contains {num}: {found}");
6     // nums contains 8: true
7 }
```

Of course, sequential search can also return the specific position of the search item, or return None if it is not found.

```

1 // sequential_search_pos.rs
2
3 fn sequential_search_pos(nums:&[i32],num:i32) -> Option<usize>{
4     let mut pos: usize = 0;
5     let mut found = false;
6     while pos < nums.len() && !found {
7         if num == nums[pos] {
8             found = true;
9         } else {
10             pos += 1;
11         }
12     }
13
14     if found { Some(pos) } else { None }
15 }
16
17 fn main() {
18     let num = 8;
19     let nums = [9,3,7,4,1,6,2,8,5];
20     match sequential_search_pos(&nums, num) {
21         Some(pos) => println!("{num}'s index: {pos}"),
22         None => println!("nums does not contain {num}"),
23     }
24     // 8's index: 7
25 }

```

6.3.2 Analysis of Sequential Search

To analyze the complexity of the sequential search algorithm, we need to set a basic computational unit. Since comparison operation is the primary operation in searching, counting the number of comparisons is the most important. Data items in a set are randomly placed and unordered, and the probability of a data item being at any position in the set is the same according to probability theory.

If the target item is not in the set, the only way to know this result is to compare the target with all the data items in the set. If there are n items, then the sequential search requires n comparisons, and the complexity is $O(n)$. What if the target item is in the set? Is the complexity still $O(n)$?

When searching for a target item in a set, the analysis of the algorithm's complexity is not always straightforward. There are three possibilities: the best case scenario is when the target item is at the beginning of the set, and only one comparison is needed, resulting in a complexity of $O(1)$. In the worst case scenario, the target item is at the end of the set, and it takes n comparisons to find it, resulting in a complexity of $O(n)$. Finally, the target item may be distributed throughout the set, and there are $n-2$ possible complexities, ranging from $O(2)$ to $O(n-1)$. In this scenario, the number of possible comparisons ranges from 1 to n times, and the average complexity equals the sum of all possible complexities divided by the total number of comparisons.

$$\sum_{i=1}^n O(i)/n = O(n/2) = O(n) \quad (6.1)$$

When the set becomes large, the complexity of sequential search on a random sequence is $O(n)$, as $1/2$ can be ignored. However, if the data items are ordered in ascending order and the probability of the

target item existing in any of the n positions is still the same, the search performance can be improved. If the target item does not exist, the search can be accelerated through techniques like binary search. For example, if searching for the target item 50, the comparison is performed in order until 56. At this point, it can be determined that there is no target value of 50 behind it because the items behind 56 are larger than 56, and the algorithm stops searching.

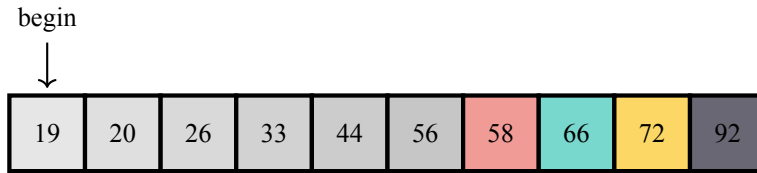


Figure 6.2: Sequential search on ordered data

The sequential search algorithm used on sorted data sets is shown below. It optimizes the algorithm by setting the "stop" variable to control the search and stop immediately when it goes beyond the range for saving time.

```

1 // ordered_sequential_search.rs
2
3 fn ordered_sequential_search(nums:&[i32], num:i32) -> bool {
4     let mut pos = 0;
5     let mut found = false;
6     // return when reads on ordered data
7     let mut stop = false;
8
9     while pos < nums.len() && !found && !stop {
10         if num == nums[pos] {
11             found = true;
12         } else if num < nums[pos] {
13             // ordered data, return
14             stop = true;
15         } else {
16             pos += 1;
17         }
18     }
19
20     found
21 }
22
23 fn main() {
24     let nums = [1,3,8,10,15,32,44,48,50,55,60,62,64];
25     let num = 44;
26     let found = ordered_sequential_search(&nums, num);
27     println!("nums contains {num}: {found}");
28     // nums contains 44: true
29
30     let num = 49;
31     let found = ordered_sequential_search(&nums, num);
32     println!("nums contains {num}: {found}");
33     // nums contains 49: false
34 }

```

In the case of sorted data, if the target item is not in the set and is less than the first item, only one comparison is needed to determine that the item is not in the set. In the worst case, the algorithm would need to compare the target item with all n items in the set, resulting in n comparisons. The average number of comparisons is still $n/2$, and the complexity remains at $O(n)$. However, the $O(n)$ complexity is better than the search on unordered data because most searches follow the average case. In fact, the average case complexity for sorted data sets can be twice as fast as unordered ones. Thus, sorting has always been a crucial topic in computer science. The complexity of sequential search for unordered and ordered data sets is summarized in the table below.

Table 6.1: Complexity of Sequential Search

Case	Min	Avg	Max	Search Class
target exist	$O(1)$	$O(\frac{n}{2})$	$O(n)$	unordered search
target not exist	$O(n)$	$O(n)$	$O(n)$	unordered search
target exist	$O(1)$	$O(\frac{n}{2})$	$O(n)$	ordered search
target not exist	$O(1)$	$O(\frac{n}{2})$	$O(n)$	ordered search

6.4 The Binary Search

An ordered dataset is advantageous for search algorithms because sequential search on sorted data sets requires comparing only $n-1$ remaining items if the first item is not the one being searched for. However, although this method can be terminated when a value outside of the range is encountered, it is still relatively slow. Is there a faster search algorithm for sorted datasets? Absolutely, it's called binary search. In the following discussion, we will analyze and implement binary search in Rust.

6.4.1 Implementing a Binary Search

As its name implies, binary search is a very important search algorithm that works by dividing the dataset into two parts for searching using low, mid, and high to control the range of the search. It starts from the middle item, not in order. If the middle item is the one being searched for, the search is completed. If it is not, we can use the ordered property of the sorted set to eliminate half of the remaining items. If the item being searched for is greater than the middle item, we can eliminate the middle item and the half of the elements smaller than the middle item, which means we don't need to compare the first item to the middle item. This is because, if the target item is greater than the middle value, it is definitely not in the first half, whether it is in the set or not. Conversely, if the target item is less than the middle item, the second half of the data can be skipped. After removing half of the data, look at the middle item in the remaining half of the data, repeat the comparison and omission process, and finally obtain the result. This search is relatively fast and very intuitive, so it is called binary search.

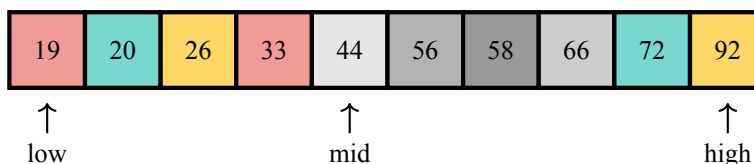


Figure 6.3: Binary search

Binary search is a search algorithm that is used on sorted data sets. It is faster than sequential search since it divides the data set into two parts using low, mid, and high to control the range of the search. If the middle item is the one being searched for, the search is completed. If not, the ordered property of the

sorted set is used to eliminate half of the remaining items. By repeating the comparison and omission process, the target item is found relatively quickly.

To implement binary search, we set low and high to the far left and far right, respectively. For instance, if we want to find 60, we compare it with the middle value of 44. If it is greater than 44, we move low to 56 and mid to 66. We then compare it with 66 and find that it is less than 66, so we move high to 58 and mid to 56. This process continues until we reach the target item.

```

1 // binary_search.rs
2
3 fn binary_search1(nums: &[i32], num: i32) -> bool {
4     let mut low = 0;
5     let mut high = nums.len() - 1;
6     let mut found = false;
7
8     // note: <= not <
9     while low <= high && !found {
10         let mid: usize = (low + high) >> 1;
11
12         //low + high may cause overflow, use subtraction
13         //let mid: usize = low + ((high - low) >> 1);
14
15         if num == nums[mid] {
16             found = true;
17         } else if num < nums[mid] {
18             // num < mid, drop right half of data
19             high = mid - 1;
20         } else {
21             // num >= mid, drop left half of data
22             low = mid + 1;
23         }
24     }
25
26     found
27 }
28
29 fn main() {
30     let nums = [1,3,8,10,15,32,44,48,50,55,60,62,64];
31
32     let target = 3;
33     let found = binary_search1(&nums, target);
34     println!("nums contains {target}: {found}");
35     // nums contains 3: true
36
37     let target = 63;
38     let found = binary_search1(&nums, target);
39     println!("nums contains {target}: {found}");
40     // nums contains 63: false
41 }

```

Binary search satisfies the three laws of recursion and can be implemented using recursion. However, recursive implementation involves slicing the dataset and discarding the mid item, which can cause stack overflow risk. Therefore, it is generally recommended to use an iterative approach to implement binary search.

```

1 // binary_search.rs
2
3 fn binary_search2(nums: &[i32], num: i32) -> bool {
4     // base case1: target does not exist
5     if 0 == nums.len() { return false; }
6
7     let mid: usize = nums.len() >> 1;
8
9     // base case2: target exists
10    if num == nums[mid] {
11        return true;
12    } else if num < nums[mid] {
13        // minimize problem size
14        return binary_search2(&nums[..mid], num);
15    } else {
16        return binary_search2(&nums[mid+1..], num);
17    }
18 }
19
20 fn main() {
21     let nums = [1,3,8,10,15,32,44,48,50,55,60,62,64];
22
23     let target = 3;
24     let found = binary_search2(&nums, target);
25     println!("nums contains {target}: {found}");
26     // nums contains 3: true
27
28     let target = 63;
29     let found = binary_search2(&nums, target);
30     println!("nums contains {target}: {found}");
31     // nums contains 63: false
32 }

```

In summary, binary search is a fast and intuitive search algorithm that works best on sorted data sets. It can be implemented using recursion but an iterative approach is generally preferred to avoid stack overflow risk.

6.4.2 Analysis of Binary Search

The binary search algorithm has a best-case scenario when the middle item is the target, and its time complexity is $O(1)$. With each comparison, binary search eliminates half of the remaining items, allowing us to determine the worst-case complexity by finding the maximum number of comparisons. After the first comparison, there are $n/2$ remaining items, then $n/4$, and so on, until the remaining items are reduced to $n/2^i = 1$, at which point the binary search ends.

$$\begin{aligned} \frac{n}{2^i} &= 1 \\ i &= \log_2(n) \end{aligned} \tag{6.2}$$

As a result, the binary search algorithm performs at most $\log_2(n)$ comparisons, giving it a time complexity of $O(\log_2(n))$, which is superior to algorithms with a time complexity of $O(n)$. However, it's important to note that in the recursive implementation of binary search, the default stack usage consumes memory, resulting in a space complexity that's inferior to the iterative version.

While binary search may seem efficient, it's not worth sorting and using binary search when n is very small. In such cases, sequential search may be more efficient. Additionally, sorting large datasets for binary search can be time-consuming and memory-intensive, making sequential search a more efficient option. However, binary search is well-suited for datasets that are neither too large nor too small, making it an ideal choice for many practical scenarios.

6.4.3 The Interpolation Search

Interpolation search is a variant of binary search that's specifically designed for sorted data. When the data is evenly distributed, interpolation search can quickly approach the search area and improve efficiency.

Unlike binary search, interpolation search doesn't use the median directly to bound the search range. Instead, it finds the upper and lower bounds using an interpolation algorithm. This algorithm uses two points (x_0, y_0) and (x_1, y_1) to perform linear interpolation and determine the value of y corresponding to any point x in the range $[x_0, x_1]$ or the value of x corresponding to any y .

$$\begin{aligned}\frac{y - y_0}{x - x_0} &= \frac{y_1 - y_0}{x_1 - x_0} \\ x &= \frac{(y - y_0)(x_1 - x_0)}{y_1 - y_0} + x_0\end{aligned}\tag{6.3}$$

To illustrate this, let's consider an example of searching for the element 27 in the sorted set of 14 elements $[1, 9, 10, 15, 16, 17, 19, 23, 27, 28, 29, 30, 32, 35]$. We use the index as the x -axis and the element value as the y -axis. It's found that $x_0 = 0, x_1 = 13$, and $y_0 = 1, y_1 = 35$. Therefore, the x value corresponding to $y = 27$ can be calculated.

$$\begin{aligned}x &= \frac{(27 - 1)(13 - 0)}{35 - 1} + 0 \\ x &= 9\end{aligned}\tag{6.4}$$

Starting with `nums[9]` which has a value of 28, which is greater than 27, we use it as the upper bound. Since the index of 28 is 9, we continue the interpolation algorithm to search for elements in the range $[0, 8]$.

$$\begin{aligned}x &= \frac{(27 - 1)(8 - 0)}{27 - 1} + 0 \\ x &= 8\end{aligned}\tag{6.5}$$

Finally, we arrive at `nums[9]`, which has a value of 27, and the algorithm ends.

```
1 // interpolation_search.rs
2
3 fn interpolation_search(nums: &[i32], target: i32) -> bool {
4     if nums.is_empty() {
5         return false;
6     }
7
8     let mut low = 0usize;
9     let mut high = nums.len() - 1;
10    loop {
11        let low_val = nums[low];
12        let high_val = nums[high];
13
14        // in this case, target isn't in the nums
15        if high <= low || target < low_val
```

```

16         || target > high_val {
17         return false;
18     }
19
20     // calculate the position for insertion
21     let offset = (target - low_val) * (high - low) as i32
22                 / (high_val - low_val);
23     let interpolant = low + offset as usize;
24
25     // update high and low
26     if nums[interpolant] > target {
27         high = interpolant - 1;
28     } else if nums[interpolant] < target {
29         low = interpolant + 1;
30     } else {
31         // found: nums[interpolant] = target
32         return true;
33     }
34 }
35 }

```

Here is an use example of this search.

```

1 fn main() {
2     let nums = [1,9,10,15,16,17,19,23,27,28,29,30,32,35];
3     let target = 27;
4     let found = interpolation_search(&nums, target);
5     println!("nums contains {target}: {found}");
6     // nums contains 27: true
7
8     let nums = [0,1,2,10,16,19,31,35,36,38,40,42,43,55];
9     let found = interpolation_search(&nums, target);
10    println!("nums contains {target}: {found}");
11    // nums contains 27: false
12 }

```

Upon careful analysis of the code, we can see that the only difference between interpolation search and binary search is the calculation method of the interpolant. In all other aspects, the two algorithms are almost identical.

When the data is evenly distributed, the time complexity of interpolation search is $O(\log \log(n))$. The proof of this result is complex and can be found in this paper^[11]. However, the worst-case and average-case complexity of interpolation search are both $O(n)$.

6.4.4 The Exponential Search

Exponential search is a variant of binary search that utilizes the exponential function to estimate the midpoint, making it suitable for sorted data without bounds. The algorithm continuously compares the value at positions of 2^0 , 2^1 , 2^2 , 2^k to the target value, then determines the search range, and finally performs binary search within that range. The time complexity of exponential search is $O(\log(i))$, where i is the position of the target value.

For instance, to find the value 22 in the sorted set of 15 elements [2,3,4,6,7,8,10,13,15,19,20,22,23,24,28], exponential search first checks whether the value at position $2^0 = 1$ is greater than 22. It then continues

to check the values at positions 2^1 , 2^2 , and 2^3 , which are 4, 7, and 15, all of which are less than 22. Checking position $2^4 = 16$ is beyond the range, so the upper bound is the last index, 14.

```

1 // exponential_search.rs
2
3 fn exponential_search(nums: &[i32], target: i32) -> bool {
4     let size = nums.len();
5     if size == 0 { return false; }
6
7     // find the upper bound
8     let mut high = 1usize;
9     while high < size && nums[high] < target {
10         high <=< 1;
11     }
12     // use the half of the upper bound as the lower bound
13     let low = high >> 1;
14
15     // use binary_search method implemented previously
16     binary_search(&nums[low..size.min(high+1)], target)
17 }
18
19 fn main() {
20     let nums = [1,9,10,15,16,17,19,23,27,28,29,30,32,35];
21     let target = 27;
22     let found = exponential_search(&nums, target);
23     println!("nums contains {target}: {found}");
24     // nums contains 27: true
25 }

```

Note that in the implementation of the exponential search algorithm, the lower bound at line 13 is half of the high but can also be set to 0. However, setting it to 0 may reduce efficiency.

The complexity of exponential search is divided into two parts: finding the upper bound for dividing the search range, and performing binary search. The complexity of finding the upper bound is related to the target value i , and its complexity is $O(\log(i))$. The complexity of binary search is $O(\log(n))$, where n is the length of the search range. The length of the search range is $high - low = 2^{\log(i)} - 2^{\log(i)-1} = 2^{\log(i)-1}$, and its complexity is $O(\log(2^{\log(i)-1})) = O(\log(i))$. Therefore, the total complexity is $O(\log(i) + \log(i)) = O(\log(i))$.

6.5 The Hash Search

The search algorithms we have discussed so far use the positional information of items in a collection to perform searches. Binary search can find data items in logarithmic time by sorting the collection. However, if an algorithm has prior knowledge of the stored addresses of different items, it can retrieve them directly without comparing them one by one. This method is called Hash Search, a search algorithm with a complexity of $O(1)$, making it the fastest search algorithm available.

To implement Hash Search, the item's address must exist first. A data structure that provides an efficient way to store and retrieve items' addresses is needed. This data structure is known as a hash table, which stores data items in such a way that they are easy to locate, with each item's position referred to as a slot or address. These slots are named starting from 0, or any other value. Once the first slot is chosen, all subsequent slots are incremented accordingly. Initially, the hash table contains no items, and each slot is empty. A hash table can be implemented using Vec, with each element initialized to None. The figure below shows a hash table with size $m = 11$, meaning that there are m slots named 0 to 10.

0	1	2	3	4	5	6	7	8	9	10
None	None	None	None	None	None	None	None	None	None	None

Figure 6.4: Hash table

The mapping between data items and their corresponding slots in the hash table is accomplished through a hash function. The hash function takes any item in the collection and returns a specific slot name, a process known as hashing. For example, if there are integer items [24, 61, 84, 41, 56, 31], and a hash table with a capacity of 11 slots, each item's location in the hash table can be obtained by inputting it into the hash function. A simple hash function involves using modulo, since any number modulo 11 will have a remainder within 11, ensuring that there is always a slot available to store the data.

$$\text{hash}(\text{item}) = \text{item} \% 11 \quad (6.6)$$

Here is the result after modulo hash calculation.

0	1	2	3	4	5	6	7	8	9	10
None	56	24	None	None	None	61	84	41	31	None

Figure 6.5: Modulo hash

Calculated the hash value, the item can be inserted into the corresponding slot in the hash table, as depicted in the figure above. Currently, 6 of the 11 slots in the table are occupied, resulting in a load factor of $\lambda = 6/11$. The load factor is a useful metric to evaluate the hash table, particularly when it needs to store a large number of items. A high load factor indicates that there is limited space for additional items, necessitating resizing the table. In languages like Rust and Go, resizing occurs automatically when the load factor exceeds a particular threshold, preparing for future data insertion.

Although the data stored in the hash table is unsorted, the hash function enables the calculation of the slot of a data item regardless of its disorderliness. For instance, to check if the number 56 exists in the table, its hash value can be computed ($\text{hash}(56) = 1$), and slot 1 can be examined to locate 56, indicating its presence in the table. The search operation has a complexity of $O(1)$, making hash search very efficient. However, conflicts may occur and must be resolved, or else the hash table cannot be utilized. For example, if 97 is added, and its hash value ($\text{hash}(97) = 9$) corresponds to an occupied slot (9) that contains 31 instead of 97, a collision occurs that necessitates resolution.

6.5.1 Hash Functions

In the previous section, the hash function computed the remainder of the item directly, ensuring that the remainder falls within a specific range. Any algorithm that can produce a value within a particular range based on the item can be used as a hash function. By improving the hash function, we can minimize the probability of collisions and create an effective hash table.

One such approach to improving the hash function is the "sum of groups" method, which divides items into equal-sized blocks (the last block may be different), adds them up, and then computes the remainder. For instance, if the data item is the phone number "316-545-0134," it can be split into two-digit numbers, padded with zeroes if required, resulting in [31, 65, 45, 01, 34]. The sum is 176, and taking the remainder of 11 yields a hash value (slot) of 0. The phone number may also be split into three-digit numbers or reversed before computing the remainder. Many hash functions are available, as any value that can be computed and then have its remainder taken can be used in various ways.

Another hash algorithm is the "mid-square" method. The item is squared first, and then the middle portion of the square is extracted as the value for which the remainder is computed. For example, if the number is 36, its square is 1296, and taking the middle portion, 29, and then computing the remainder of 11 yields $\text{hash}(29) = 7$, indicating that 36 should be stored in slot 7.

When storing a string, the remainder can also be calculated based on the ASCII values of its characters. The string "rust" has four characters with ASCII values of [114, 117, 115, 116], and their sum is 462. Computing the remainder results in $\text{hash}(462) = 0$. Other strings, such as "Java" with ASCII values [74, 97, 118, 97], may also be tried. Its sum is 386, and taking the remainder gives $\text{hash}(386) = 1$, indicating that "Java" should be stored in slot 1, as shown in the figure below.

0	1	2	3	4	5	6	7	8	9	10
rust	Java	None	C#	None	go	None	None	html	C++	css

The following is an implementation of hash function for ASCII value.

```

1 // hash.rs
2
3 fn hash1(astr: &str, size: usize) -> usize {
4     let mut sum = 0;
5     for c in astr.chars() {
6         sum += c as usize;
7     }
8
9     sum % size
10 }
11
12 fn main() {
13     let s1 = "rust"; let s2 = "Rust";
14     let size = 11;
15
16     let p1 = hash1(s1, size);
17     let p2 = hash1(s2, size);
18     println!("{s1} in slot {p1}, {s2} in slot {p2}");
19     // rust in slot 0, Rust in slot 1
20 }
```

To mitigate collisions that may arise when using the above hash function, one can modify it slightly. For example, one can use the position of different characters in a string as weights and multiply the ASCII values by their positional weights.

$$\begin{aligned}
 \text{hash}(\text{rust}) &= (0 * 114 + 1 * 117 + 2 * 115 + 3 * 116) \% 11 \\
 &= 695 \% 11 \\
 &= 2
 \end{aligned}
 \tag{6.7}$$

Starting the index from 1 is preferred to ensure that the first character contributes to the total. Below is an example calculation and code snippet.

$$\begin{aligned}
 \text{hash}(\text{rust}) &= (1 * 114 + 2 * 117 + 3 * 115 + 4 * 116) \% 11 \\
 &= 1157 \% 11 \\
 &= 2
 \end{aligned}
 \tag{6.8}$$

```

1 // hash.rs
2 fn hash2(astr: &str, size: usize) -> usize {
3     let mut sum = 0;
4     for (i, c) in astr.chars().enumerate() {
5         sum += (i + 1) * (c as usize);
6     }
7     sum % size
8 }
9
10 fn main() {
11     let (s1, s2, size) = ("rust", "Rust", 11);
12     let p1 = hash2(s1, size);
13     let p2 = hash2(s2, size);
14     // rust in slot 2, Rust in slot 3
15 }

```

Efficiency is crucial when designing a hash function to ensure that it does not become the bottleneck of the system. If the hash function is too complex, it could break the $O(1)$ complexity.

6.5.2 Collision Resolution

To address the issue of hash collisions, collision resolution methods are necessary. For instance, if a hash function with position as weight is used and the index starts from 0, the strings "rust" and "Rust" will collide. When two items are hashed to the same slot, a method must be used to insert the conflicting item into another slot of the hash table. This process is called collision resolution.

Although perfect hash tables do not exist due to limited memory and complex real-world situations, one intuitive method for resolving collisions is to search the hash table and attempt to find the next empty slot to save the conflicting item. The simplest method is called open addressing, specifically linear probing. Linear probing attempts to linearly probe the next empty slot in the hash table, starting from the original collision point until the first empty slot is found. Note that you can start searching again from the beginning after reaching the end. The following figure shows a hash table with 11 slots, where we try to insert 35 and 47.

0	1	2	3	4	5	6	7	8	9	10
None	56	24	None	None	None	61	84	41	31	None

When inserting 35, its position should be slot 2, but we find that slot 2 already contains 24. Therefore, we start searching for an empty slot from slot 2 and find that slot 3 is empty, so we insert 35 there.

0	1	2	3	4	5	6	7	8	9	10
None	56	24	35	None	None	61	84	41	31	None

When we insert 47, we find that slot 3 has the value 35, so we search for the next empty slot and find that slot 4 is empty, so we insert 47 there.

0	1	2	3	4	5	6	7	8	9	10
None	56	24	35	47	None	61	84	41	31	None

To search for items in a hash table built using open addressing, we must use the same method as used during its construction. For instance, if we are searching for item 56 with a hash of 1, we locate it in slot 56 and return true. However, if we are looking for item 35, which has a hash of 2, we may find another item like 24 instead of 35 due to collisions. To avoid returning false, we need to conduct a sequential search until we locate 35, an empty slot, or we loop back to 24 before returning the result.

However, using a sequential search may lead to data clustering, where items cluster in the table due to multiple collisions occurring in the same hash slot. Linear probing fills subsequent slots with conflicting items, forcing the originally intended values to be inserted elsewhere. Sequential searches are time-consuming, with a complexity greater than $O(1)$. To address data clustering, open addressing technology can be extended to skip several slots instead of sequentially searching for the next open slot. By checking every three slots when a conflict occurs, for example, the conflicting items are more evenly distributed, thus dispersing the conflict. This method has a significant effect and can alleviate data clustering, as shown in the figure below.

0	1	2	3	4	5	6	7	8	9	10
None	56	24	47	None	35	61	84	41	31	None

When adding item 35, a conflict may occur, prompting the search to skip every three slots from that point onwards, thus dispersing the conflict. Consequently, item 47 can be inserted without any conflict. This method has proven effective in alleviating data clustering, as demonstrated in the following figure.

Rehashing is the process of finding an alternative slot in a hash table after a collision occurs. This process involves calculating a new hash value using a specified skip size, which must ensure that all slots in the table can eventually be accessed. To guarantee this, it is advisable to use a prime number as the table size, as exemplified by the use of 11 in the accompanying example.

$$rehash(pos) = (pos + n) \% size \quad (6.9)$$

Another approach to resolving conflicts is the chain method. This method involves setting up a linked list to store data items for each conflicting position, as shown in Figure (6.6). When searching, conflicts are resolved by sequentially searching the chain, which has a complexity of $O(n)$. If the data in the conflict chain is sorted, a binary search can be utilized to achieve a complexity of $O(\log 2(n))$. If the chain becomes too long, it can be transformed into a red-black tree to enhance its structural stability. In many programming languages, the chain method is the default implementation for resolving conflicts in hash table data structures.

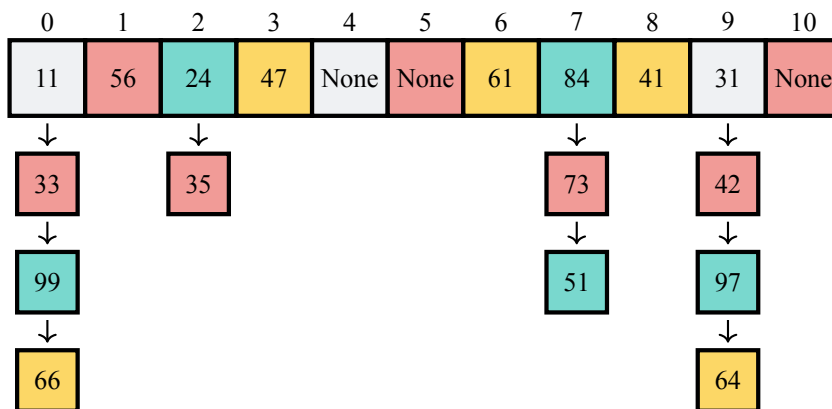


Figure 6.6: Collision Resolution by Chain Method

6.5.3 Implementing a HashMap in Rust

The most commonly used collection type in Rust is the HashMap. It is an associative data type that stores key-value pairs. The key is utilized to locate the position of the value, and due to the position uncertainty, this type of search is akin to searching on a map. Thus, this data structure is called HashMap.

HashMap is an unordered collection of key-value pairs where the keys are unique, and there is a one-to-one relationship between the keys and the values. Here is the abstract data type definition of HashMap:

- `new()`: creates a new HashMap, takes no arguments and returns an empty HashMap collection.
- `insert(k,v)`: adds a new key-value pair to the HashMap, taking parameters `k` and `v`. If the key exists, the old value is replaced with the new value. No return value is provided.
- `remove(k)`: removes value corresponding to `k` from the HashMap, takes parameter `k`, and returns the corresponding value `v`.
- `get(k)`: returns the value `v` stored in the HashMap for the given key `k` (which may be empty `None`), takes parameter `k`.
- `contains(k)`: returns `true` if the key `k` exists in the HashMap, otherwise `false`, takes parameter `k`.
- `len()`: returns the number of key-value pairs stored in the HashMap, takes no arguments.

The table below displays the results of operations assuming that 'h' is an already created empty HashMap (represented as `{}` here).

Table 6.2: HashMap operations

Operation	Value	Return
<code>h.is_empty()</code>	<code>{}</code>	<code>true</code>
<code>h.insert("a", 1)</code>	<code>{a:1}</code>	
<code>h.insert("b", 2)</code>	<code>{a:1, b:2}</code>	
<code>h.insert("c", 3)</code>	<code>{a:1, b:2, c:3}</code>	
<code>h.get("b")</code>	<code>{a:1, b:2, c:3}</code>	<code>Some(2)</code>
<code>h.get("d")</code>	<code>{a:1, b:2, c:3}</code>	<code>None</code>
<code>h.len()</code>	<code>{a:1, b:2, c:3}</code>	<code>3</code>
<code>h.contains("c")</code>	<code>{a:1, b:2, c:3}</code>	<code>true</code>
<code>h.contains("e")</code>	<code>{a:1, b:2, c:3}</code>	<code>false</code>
<code>h.remove("a")</code>	<code>{b:2, c:3}</code>	<code>Some(1)</code>
<code>h.remove("a")</code>	<code>{b:2, c:3}</code>	<code>None</code>
<code>h.insert("a", 1)</code>	<code>{b:2, c:3, a:1}</code>	
<code>h.contains("a")</code>	<code>{b:2, c:3, a:1}</code>	<code>true</code>
<code>h.len()</code>	<code>{b:2, b:3, a:1}</code>	<code>3</code>
<code>h.get("a")</code>	<code>{b:2, c:3, a:1}</code>	<code>Some(1)</code>
<code>h.remove("c")</code>	<code>{b:2, a:1}</code>	<code>Some(3)</code>

In actual implementation, a HashMap in Rust is made up of two separate Vecs - one for storing the keys (called slot) and the other for storing the values (called data). The slot Vec saves the keys with the index starting from 1, while the default value in the slot Vec is 0. The HashMap is encapsulated by a struct, and a cap parameter is added to control the capacity.

```

1 // hashmap.rs
2
3 #[derive(Debug, Clone, PartialEq)]
4 struct HashMap <T> {
5     cap: usize, // capacity
6     slot: Vec<usize>, // store data address (index)
7     data: Vec<T>, // store elements
8 }

```

The rehash function in Rust can be implemented using a linear search method that adds 1, which is simple and easy to implement. The initial size of the HashMap is set to 11, but it can also be set to other prime numbers, such as 13, 17, 19, 23, 29, etc. Below is the complete implementation code for a HashMap in Rust.

```

1 // hashmap.rs
2
3 impl<T: Clone + PartialEq + Default> HashMap<T> {
4     fn new(cap: usize) -> Self {
5         // Initialize slot and data
6         let mut slot = Vec::with_capacity(cap);
7         let mut data = Vec::with_capacity(cap);
8         for _i in 0..cap{
9             slot.push(0);
10            data.push(Default::default());
11        }
12
13        HashMap { cap, slot, data }
14    }
15
16    fn len(&self) -> usize {
17        let mut len = 0;
18        for &d in self.slot.iter() {
19            // If slot is not empty, then increase len by 1
20            if 0 != d {
21                len += 1;
22            }
23        }
24        len
25    }
26
27    fn is_empty(&self) -> bool {
28        let mut empty = true;
29        for &d in self.slot.iter() {
30            if 0 != d {
31                empty = false;
32                break;
33            }
34        }
35        empty
36    }
37
38    fn clear(&mut self) {
39        let mut slot = Vec::with_capacity(self.cap);
40        let mut data = Vec::with_capacity(self.cap);
41        for _i in 0..self.cap{
42            slot.push(0);
43            data.push(Default::default());
44        }
45
46        self.slot = slot;
47        self.data = data;

```

```

48     }
49
50     fn hash(&self, key: usize) -> usize {
51         key % self.cap
52     }
53
54     fn rehash(&self, pos: usize) -> usize {
55         (pos + 1) % self.cap
56     }
57
58     fn insert(&mut self, key: usize, value: T) {
59         if 0 == key { panic!("Error: key must > 0"); }
60
61         let pos = self.hash(key);
62         if 0 == self.slot[pos] {
63             // If the slot is empty, insert directly
64             self.slot[pos] = key;
65             self.data[pos] = value;
66         } else {
67             // If the slot is not empty, then
68             // find next available position
69             let mut next = self.rehash(pos);
70             while 0 != self.slot[next]
71                 && key != self.slot[next] {
72                 next = self.rehash(next);
73
74                 // If the slot is full, exit
75                 if next == pos {
76                     println!("Error: slot is full!");
77                     return;
78                 }
79             }
80
81             // Insert the data in the found slot
82             if 0 == self.slot[next] {
83                 self.slot[next] = key;
84                 self.data[next] = value;
85             } else {
86                 self.data[next] = value;
87             }
88         }
89     }
90
91     fn remove(&mut self, key: usize) -> Option<T> {
92         if 0 == key { panic!("Error: key must > 0"); }
93
94         let pos = self.hash(key);
95         if 0 == self.slot[pos] {
96             // If the slot is empty, return None
97             None
98         } else if key == self.slot[pos] {
99             // If found the same key,

```

```

100         // update the slot and data
101         self.slot[pos] = 0;
102         let data = Some(self.data[pos].clone());
103         self.data[pos] = Default::default();
104         data
105     } else {
106         let mut data: Option<T> = None;
107         let mut stop = false;
108         let mut found = false;
109         let mut curr = pos;
110
111         while 0 != self.slot[curr] && !found && !stop {
112             if key == self.slot[curr] {
113                 // If the value is found, delete the data
114                 found = true;
115                 self.slot[curr] = 0;
116                 data = Some(self.data[curr].clone());
117                 self.data[curr] = Default::default();
118             } else {
119                 // If rehashing returns to the initial
120                 // position, it means that it has searched
121                 // a full circle and still not found
122                 curr = self.rehash(curr);
123                 if curr == pos {
124                     stop = true;
125                 }
126             }
127         }
128
129         data
130     }
131 }
132
133 fn get_pos(&self, key: usize) -> usize {
134     if 0 == key { panic!("Error: key must > 0"); }
135
136     // Calculate the data position
137     let pos = self.hash(key);
138     let mut stop = false;
139     let mut found = false;
140     let mut curr = pos;
141
142     // Loop through to find the data
143     while 0 != self.slot[curr] && !found && !stop {
144         if key == self.slot[curr] {
145             found = true;
146         } else {
147             // If rehashing returns to the initial position
148             // it means that it has searched a full circle
149             // and still not found
150             curr = self.rehash(curr);
151             if curr == pos {

```

```

152             stop = true;
153         }
154     }
155 }
156
157     curr
158 }
159
160 // Get a reference and a mutable reference to val
161 fn get(&self, key: usize) -> Option<&T> {
162     let curr = self.get_pos(key);
163     self.data.get(curr)
164 }
165
166 fn get_mut(&mut self, key: usize) -> Option<&mut T> {
167     let curr = self.get_pos(key);
168     self.data.get_mut(curr)
169 }
170
171 fn contains(&self, key: usize) -> bool {
172     if 0 == key {
173         panic!("Error: key must > 0");
174     }
175     self.slot.contains(&key)
176 }
177
178 // Implement iteration and mutable iteration for hashmap
179 fn iter(&self) -> Iter<T> {
180     let mut iterator = Iter { stack: Vec::new() };
181     for item in self.data.iter() {
182         iterator.stack.push(item);
183     }
184     iterator
185 }
186
187 fn iter_mut(&mut self) -> IterMut<T> {
188     let mut iterator = IterMut { stack: Vec::new() };
189     for item in self.data.iter_mut() {
190         iterator.stack.push(item);
191     }
192     iterator
193 }
194 }
195
196 // Implementation of iteration
197 struct Iter<'a, T: 'a> { stack: Vec<&'a T>, }
198 impl<'a, T> Iterator for Iter<'a, T> {
199     type Item = &'a T;
200     fn next(&mut self) -> Option<Self::Item> {
201         self.stack.pop()
202     }
203 }

```

```

204
205 struct IterMut<'a, T: 'a> { stack: Vec<&'a mut T>, }
206 impl<'a, T> Iterator for IterMut<'a, T> {
207     type Item = &'a mut T;
208     fn next(&mut self) -> Option<Self::Item> {
209         self.stack.pop()
210     }
211 }
212
213 fn main() {
214     basic();
215     iter();
216
217     fn basic() {
218         let mut hmap = HashMap::new(11);
219         hmap.insert(2,"dog");
220         hmap.insert(3,"tiger");
221         hmap.insert(10,"cat");
222
223         println!("empty: {}, size: {:?}",
224                 hmap.is_empty(), hmap.len());
225         println!("contains key 2: {}", hmap.contains(2));
226
227         println!("key 3: {:?}", hmap.get(3));
228         let val_ptr = hmap.get_mut(3).unwrap();
229         *val_ptr = "fish";
230         println!("key 3: {:?}", hmap.get(3));
231         println!("remove key 3: {:?}", hmap.remove(3));
232         println!("remove key 3: {:?}", hmap.remove(3));
233
234         hmap.clear();
235         println!("empty: {}, size: {:?}",
236                 hmap.is_empty(), hmap.len());
237     }
238
239     fn iter() {
240         let mut hmap = HashMap::new(11);
241         hmap.insert(2,"dog");
242         hmap.insert(3,"tiger");
243         hmap.insert(10,"cat");
244
245         for item in hmap.iter() {
246             println!("val: {item}");
247         }
248
249         for item in hmap.iter_mut() {
250             *item = "fish";
251         }
252
253         for item in hmap.iter() {
254             println!("val: {item}");
255         }

```

```

256     }
257 }

```

Here is the code execution output.

```

empty: false, size: 3
contains key 2: true
key 3: Some("tiger")
key 3: Some("fish")
remove key 3: Some("fish")
remove key 3: None
empty: true, size: 0
val: cat
val:
val:
val:
val:
val:
val:
val: tiger
val: dog
val:
val:
val: fish
val: fish
val: fish
val: fish
val: fish
val: fish
val: fish
val: fish
val: fish
val: fish
val: fish
val: fish

```

6.5.4 Analysis of HashMap

A hash table provides a lookup complexity of $O(1)$ in the best case. However, due to collisions, the number of comparisons during the lookup process can vary. The severity of collisions affects the performance of the hash table, and a good evaluation metric is the load factor λ . A small load factor means the chance of collisions is low, and items are more likely to be in their corresponding slot. A large load factor means the table is filling up quickly, and there are more collisions, leading to a more complex conflict resolution process.

The result of each search is either success or failure. For searches performed using linear probing open addressing, the average number of comparisons for success is about $\frac{1 + \frac{1}{1-\lambda}}{2}$, and for failure, it is about $\frac{1 + (\frac{1}{1-\lambda})^2}{2}$. Even when using chaining, the average number of comparisons for success is $1 + \lambda/2$, and for failure, it is λ . Overall, the lookup complexity of a hash table is around $O(\lambda)$.

When using the chaining method in a hash table, an increase in collisions results in more items being added to the same chain. As a result, searching for a specific item in the chain takes longer, making it the main contributor to the search time.

6.6 Summary

This chapter covers several search algorithms, namely sequential search, binary search, interpolation search, exponential search and hash search. Sequential search is a straightforward algorithm that has a complexity of $O(n)$. Binary search, on the other hand, is a fast algorithm that cuts the data set in half each time, but it requires the data to be sorted, with a complexity of $O(\log_2(n))$. Other search algorithms, such as interpolation search and exponential search, build on binary search and are suitable for different types of data distribution. Hash search is a highly efficient $O(1)$ search algorithm that uses a HashMap. However, it is crucial to consider that hash tables are prone to collisions and need appropriate measures, such as open addressing and chaining, to resolve them. Sorting is a helpful technique that speeds up search algorithms, and in the next chapter, we will delve into sorting algorithms.

Chapter 7

Sorting

7.1 Objectives

- Learn sorting algorithms.
- Be able to implement the ten basic sorting algorithms in Rust.

7.2 What is Sorting?

Sorting involves arranging elements in a specific order in a collection. For example, playing cards are shuffled and arranged by suit or order to play effectively, and words can be sorted alphabetically or by length. Similarly, Chinese cities can be sorted by population, ethnicity, or region. Orderliness is a crucial skill for humans as it enables smooth progress in various areas.

Sorting is an important field in computer science, and many pioneers have contributed to the development of sorting algorithms. Various algorithms, including sequential search and binary search, benefit from sorting datasets. Sorting algorithm efficiency is dependent on the number of items being processed, and complex sorting methods are too expensive for small collections while simple algorithms are not suitable for large datasets. In this chapter, we will explore different sorting algorithms and compare their performance.

Comparison is the core operation in sorting, as it determines sequence, order, and orderliness. Comparison involves looking at which element is larger or smaller based on a criterion, which can be simple (numerical size) or abstract (health index). If the comparison shows that the order is incorrect, data positions need to be exchanged. However, exchanging data positions is an expensive operation in computer science, and the total number of exchanges is crucial for evaluating the efficiency of algorithms. In this chapter, we will analyze various sorting algorithms using the Big O analysis method, which is the most intuitive indicator for evaluating their efficiency.

Apart from efficiency, sorting also has the issue of stability. For instance, in a list like [1,4,9,8,5,5,2,3,7,6], there are two 5s, and different sorts may swap the order of the two 5s. Although they are adjacent in the end, the original order relationship has been destroyed. For numeric data, this may not matter, but for data structures containing keys like personal information, rash sorting may change the order, which is not desirable.

```
person {          person {
    amount: 5,      amount: 5,
    name: Shieber,  name: Kew,
    phone: 133xx,   phone: 133xx,
    age: 20,        age: 24,
}                  }
```

To ensure that the order of the elements in a sorted collection remains stable, it is important to consider stability when evaluating sorting algorithms. Stability refers to whether the relative order of equal elements in the original collection is maintained in the sorted collection. For example, if the amount: 5 appears twice in a collection and sorting changes the order of the two occurrences, this can cause issues for algorithms that rely on the stability of the sequence, such as deduction operations.

There are numerous sorting algorithms available for collections, but there are ten basic types of sorting algorithms that serve as the foundation for most of them. These include bubble sort, quick sort, selection sort, heap sort, insertion sort, shell sort, merge sort, counting sort, bucket sort, and radix sort. Many improved algorithms have been derived from these ten basic algorithms. In this article, we will explain several of these improved algorithms, including new bubble sort, cocktail sort, comb sort, binary insertion sort, flash sort, and Tim sort.

7.3 The Bubble Sort

Bubble sort is a simple sorting algorithm that requires multiple iterations through the collection, comparing adjacent items and swapping those that are unordered. Each iteration places the largest value in its correct position. The algorithm gets its name from the similarity of the sorting process to bubbles rising to the top of a pot of boiling water.

In the first pass, the algorithm compares adjacent items and swaps them if they are in the wrong order. If there are n items in the list, there will be $n-1$ comparisons in the first pass. In the second pass, the largest value in the data set is already in the correct position, and the remaining $n-1$ items need to be sorted, which means there will be $n-2$ comparisons. The algorithm requires $n-1$ iterations to place the next largest value in its proper position. After completing $n-1$ rounds of iteration and comparison, the smallest item will definitely be in the correct position, and no further processing is needed.

92	84	66	56	44	31	72	19	24
84	92	66	56	44	31	72	19	24
84	66	92	56	44	31	72	19	24
84	66	56	92	44	31	72	19	24
84	66	56	44	92	31	72	19	24
84	66	56	44	31	92	72	19	24
84	66	56	44	31	72	92	19	24
84	66	56	44	31	72	19	92	24
84	66	56	44	31	72	19	24	92

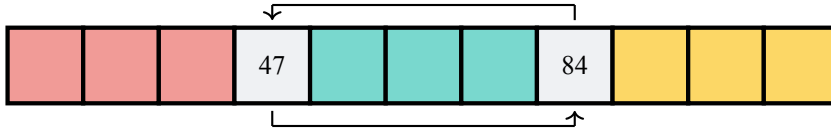
Figure 7.1: Bubble sort

The diagonal line in the figure above represents the maximum value, and it keeps moving towards the right, just like bubbles rising to the top. Bubble sort involves frequent swap operations, which are com-

monly used auxiliary operations in comparisons. In Rust, the `Vec` data structure defaults to implementing the `swap()` function, but you can also implement the following swap operation.

```
1 // swap
2 let temp = data[i];
3 data[i] = data[j];
4 data[j] = temp;
```

Some programming languages provide a convenient way to swap values without using temporary variables, such as: `data[i], data[j] = data[j], data[i]`. Although this feature still uses variables internally, it operates on two variables simultaneously, as shown in the figure below.



In this chapter, we only deal with sets of numbers to simplify the algorithm design. Therefore, we can use a `Vec` to implement bubble sort.

```
1 // bubble_sort.rs
2
3 fn bubble_sort1(nums: &mut [i32]) {
4     if nums.len() < 2 {
5         return;
6     }
7
8     for i in 1..nums.len() {
9         for j in 0..nums.len()-i {
10             if nums[j] > nums[j+1] {
11                 nums.swap(j, j+1);
12             }
13         }
14     }
15 }
```

Here is an example of bubble sort.

```
1 fn main() {
2     let mut nums = [54,26,93,17,77,31,44,55,20];
3     bubble_sort1(&mut nums);
4     println!("sorted nums: {:?}", nums);
5     // sorted nums: [17, 20, 26, 31, 44, 54, 55, 77, 93]
6 }
```

It is worth noting that to avoid a double for-loop, we can also use a while-loop to implement bubble sort.

```
1 // bubble_sort.rs
2
3 fn bubble_sort2(nums: &mut [i32]) {
4     let mut len = nums.len() - 1;
5
6     while len > 0 {
7         for i in 0..len {
```

```

8             if nums[i] > nums[i+1] {
9                 nums.swap(i, i+1);
10            }
11        }
12
13        len -= 1;
14    }
15 }
16
17 fn main() {
18     let mut nums = [54,26,93,17,77,31,44,55,20];
19     bubble_sort2(&mut nums);
20     println!("sorted nums: {:?}", nums);
21     // sorted nums: [17, 20, 26, 31, 44, 54, 55, 77, 93]
22 }

```

Bubble sort involves comparing adjacent items and swapping those that are unordered, which repeats until the largest item is in its correct position. The algorithm performs $n - 1$ rounds of traversal to sort n numbers, regardless of the initial arrangement. During the first round, $n - 1$ comparisons are required, followed by $n - 2$ comparisons in the second round, and so on, until only one comparison is needed. The total number of comparisons can be calculated using the formula:

$$1 + 2 + \dots + n - 1 = \frac{n^2}{2} + \frac{n}{2} \quad (7.1)$$

This means that the time complexity of bubble sort is $O(\frac{n^2}{2} + \frac{n}{2}) = O(n^2)$.

Although both bubble sort algorithms presented above achieve sorting, even a sorted set requires continuous comparisons and swapping of data items. To optimize the algorithm, we can add a variable to control whether comparisons should continue and exit directly when encountering a sorted set.

```

1 // bubble_sort.rs
2
3 fn bubble_sort3(nums: &mut [i32]) {
4     // compare controls whether to continue comparing
5     let mut compare = true;
6     let mut len = nums.len() - 1;
7
8     while len > 0 && compare {
9         compare = false;
10        for i in 0..len {
11            if nums[i] > nums[i+1] {
12                // Data is unordered and need to compare
13                nums.swap(i, i+1);
14                compare = true;
15            }
16        }
17
18        len -= 1;
19    }
20 }
21
22 fn main() {
23     let mut nums = [54,26,93,17,77,31,44,55,20];

```

```

24     bubble_sort3(&mut nums);
25     println!("sorted nums: {:?}", nums);
26     // sorted nums: [17, 20, 26, 31, 44, 54, 55, 77, 93]
27 }

```

Bubble sort is a sorting algorithm that compares adjacent elements in an array, starting from the first number and exchanging positions based on their relative size. Elements are only swapped from left to right. However, can we perform bubble sort from right to left? Yes, we can, and this bidirectional sorting method is called cocktail sort. Cocktail sort is a variant of bubble sort that sorts in descending order when sorting from right to left. While it slightly optimizes bubble sort, its time complexity is still $O(n^2)$, but it approaches $O(n)$ if the sequence is already sorted.

```

1  // cocktail_sort.rs
2
3  fn cocktail_sort(nums: &mut [i32]) {
4      if nums.len() <= 1 { return; }
5
6      // bubble controls the sort process
7      let mut bubble = true;
8      let len = nums.len();
9      for i in 0..(len >> 1) {
10         if bubble {
11             bubble = false;
12             // bubble from left to right
13             for j in i..(len - i - 1) {
14                 if nums[j] > nums[j+1] {
15                     nums.swap(j, j+1);
16                     bubble = true
17                 }
18             }
19             // bubble from right to left
20             for j in (i+1..(len - i - 1)).rev() {
21                 if nums[j] < nums[j-1] {
22                     nums.swap(j-1, j);
23                     bubble = true
24                 }
25             }
26         } else {
27             break;
28         }
29     }
30 }
31
32 fn main() {
33     let mut nums = [1,3,2,8,3,6,4,9,5,10,6,7];
34     cocktail_sort(&mut nums);
35     println!("sorted nums {:?}", nums);
36     // sorted nums [1, 2, 3, 3, 4, 5, 6, 6, 7, 8, 9, 10]
37 }

```

In contrast to bubble sort, comb sort can compare items with a distance greater than 1. Comb sort starts by setting the gap to the length of the array and decreasing it by a fixed ratio in each iteration of the loop, typically by multiplying it by 0.8, which is the most effective ratio determined by the original

author through experimentation. When the gap is 1, comb sort degenerates into bubble sort. Comb sort aims to move inverted numbers forward as much as possible and ensure that the numbers in the current gap are sorted, similar to combing hair with a comb, where the gap is similar to the gap between the teeth of the comb. Comb sort has a time complexity of $O(n \log n)$, with a space complexity of $O(1)$, and it is an unstable sorting algorithm.

```

1 // comb_sort.rs
2
3 fn comb_sort(nums: &mut [i32]) {
4     if nums.len() <= 1 { return; }
5     let mut i;
6     let mut gap: usize = nums.len();
7
8     // ordered basicly
9     while gap > 0 {
10         gap = (gap as f32 * 0.8) as usize;
11         i = gap;
12         while i < nums.len() {
13             if nums[i-gap] > nums[i] {
14                 nums.swap(i-gap, i);
15             }
16             i += 1;
17         }
18     }
19
20     // rearrange the element properly.
21     // exchange controls the process
22     let mut exchange = true;
23     while exchange {
24         exchange = false;
25         i = 0;
26         while i < nums.len() - 1 {
27             if nums[i] > nums[i+1] {
28                 nums.swap(i, i+1);
29                 exchange = true;
30             }
31             i += 1;
32         }
33     }
34 }

```

```

1 fn main() {
2     let mut nums = [1,2,8,3,4,9,5,6,7];
3     comb_sort(&mut nums);
4     println!("sorted nums {:?}", nums);
5     // sorted nums [1, 2, 3, 4, 5, 6, 7, 8, 9]
6 }

```

Bubble sort suffers from the problem that the boundary index, such as $i, j, i+1, j+1$, must be arranged properly and cannot be wrong. In 2021, a new sorting algorithm^[12] was published that does not require handling boundary index values. It is intuitive and resembles bubble sort at first glance, but it is actually similar to insertion sort. Although it looks like a descending sort, it is actually an ascending sort.

```

1 // CantBelieveItCanSort.rs
2
3 fn cbic_sort1(nums: &mut [i32]) {
4     for i in 0..nums.len() {
5         for j in 0..nums.len() {
6             if nums[i] < nums[j] { nums.swap(i, j); }
7         }
8     }
9 }
10
11 fn main() {
12     let mut nums = [54,32,99,18,75,31,43,56,21,22];
13     cbic_sort1(&mut nums);
14     println!("sorted nums {:?}", nums);
15     // sorted nums [18, 21, 22, 31, 32, 43, 54, 56, 75, 99]
16 }

```

Of course, it can also be implemented as a descending sort by changing the less than symbol to a greater than symbol.

```

1 // CantBelieveItCanSort.rs
2
3 fn cbic_sort2(nums: &mut [i32]) {
4     for i in 0..nums.len() {
5         for j in 0..nums.len() {
6             if nums[i] > nums[j] {
7                 nums.swap(i, j);
8             }
9         }
10     }
11 }

```

```

1 fn main() {
2     let mut nums = [54,32,99,18,75,31,43,56,21,22];
3     cbic_sort2(&mut nums);
4     println!("sorted nums {:?}", nums);
5     // sorted nums [99, 75, 56, 54, 43, 32, 31, 22, 21, 18]
6 }

```

This algorithm uses only two for loops, and the index values do not need to be handled separately. While it resembles the definition of bubble sort, it is not actually a bubble sort algorithm.

7.4 The Quick Sort

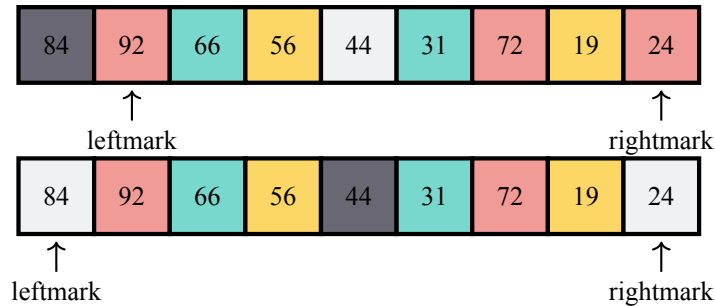
Quicksort and bubble sort do have some similarities, but it is not accurate to say that quicksort is an upgraded version of bubble sort. Quicksort is a completely different algorithm that uses the divide-and-conquer strategy to speed up the sorting process.

The quicksort algorithm involves two main steps: selecting a pivot value and partitioning the set. The pivot value can be any value in the collection, but it is typically chosen to be a value that is close to the middle of the final sorted collection for the fastest sorting speed. There are many different methods for selecting a pivot value, but this article only explains the principle and does not consider algorithm optimization.

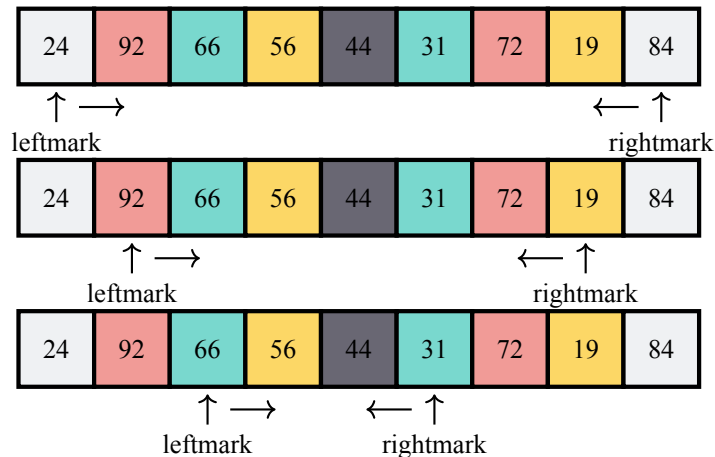
In the following figure, the pivot value of 84 is not necessarily at the middle of the sorted collection, and it would be more efficient to choose a value closer to the middle, such as 56, as the pivot value. However, choose a correct pivot value is not the focus of this article.



To implement quicksort, two markers need to be set for comparison after selecting the pivot value (the dark gray value). The left and right markers should be located at the far left and far right extremes of the collection, except for the pivot value.



The goal of partitioning is to move the items that are misaligned with respect to the pivot value. By comparing the values at the left and right markers with the pivot value and swapping the smaller value to the left marker and the larger value to the right marker, a basically sorted collection can be achieved quickly through repeated swaps.

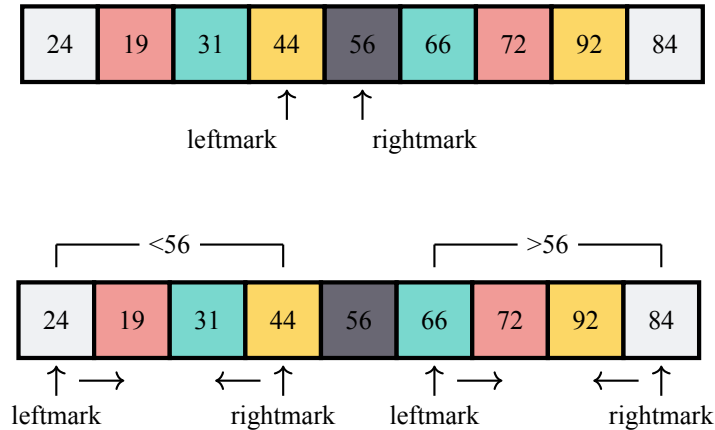


To begin partitioning, right shift the left index until a value greater than or equal to the pivot value is found. Then left shift the decreasing right index until a value less than or equal to the pivot value is found. If the value of the left index is greater than the right index, swap the values. In this case, 84 and 24 satisfy this condition, so the values are directly swapped. Repeat this process until the left and right indices cross each other.

After crossing, compare the values of the left and right indices. If the right is less than the left, swap the right index value with the pivot value. Otherwise, swap the left index value with the pivot value. The right index value serves as the splitting point, dividing the set into two intervals.

Recursively call quicksort on the left and right intervals until the sorting is completed. It is important to note that the pivot value does not necessarily have to be the value in the middle of the collection but

should be a value that is in the middle or close to the middle of the final sorted collection for the fastest sorting speed.



Once quicksort is executed on both the left and right sides, the sorting process is completed. If the set's length is less than or equal to one, it's already sorted, and the program can exit directly. To implement quicksort, we have a dedicated partition function that selects the first item (or selects a random value in the data set) as the pivot value.

```

1 // quick_sort.rs
2
3 fn quick_sort1(nums: &mut [i32], low: usize, high: usize) {
4     if low < high {
5         let split = partition(nums, low, high);
6
7         // avoid out of range (split <= 1) and syntax error
8         if split > 1 {
9             quick_sort1(nums, low, split - 1);
10        }
11
12        quick_sort1(nums, split + 1, high);
13    }
14 }
15
16 fn partition(nums:&mut[i32], low:usize,high:usize) -> usize {
17     // left marker and right marker
18     let mut lm = low;
19     let mut rm = high;
20
21     loop {
22         // left mark move to right gradually
23         while lm <= rm && nums[lm] <= nums[low] {
24             lm += 1;
25         }
26
27         // right mark move to left gradually
28         while lm <= rm && nums[rm] >= nums[low] {
29             rm -= 1;
30         }

```

```

31
32     // once lm > rm, return and exchange data between
33     // position lm and rm
34     if lm > rm {
35         break;
36     } else {
37         nums.swap(lm, rm);
38     }
39 }
40
41     nums.swap(low, rm);
42
43     rm
44 }
45
46 fn main() {
47     let mut nums = [54,26,93,17,77,31,44,55,20];
48     let high = nums.len() - 1;
49     quick_sort1(&mut nums, 0, high);
50     println!("sorted nums: {:?}", nums);
51     // sorted nums: [17, 20, 26, 31, 44, 54, 55, 77, 93]
52 }

```

However, it's also possible to implement quicksort directly through the recursive method without using the partition function.

```

1 // quick_sort.rs
2
3 fn quick_sort2(nums: &mut [i32], low: usize, high: usize) {
4     if low >= high {
5         return;
6     }
7
8     // left marker and right marker
9     let mut lm = low;
10    let mut rm = high;
11    while lm < rm {
12        // right marker move to left gradually
13        while lm < rm && nums[low] <= nums[rm] {
14            rm -= 1;
15        }
16
17        // left marker move to right gradually
18        while lm < rm && nums[lm] <= nums[low] {
19            lm += 1;
20        }
21
22        // exchange data between position lm and rm
23        nums.swap(lm, rm);
24    }
25
26    // exchange data between position low and lm
27    nums.swap(low, lm);

```

```

28
29     if lm > 1 {
30         quick_sort2(nums, low, lm - 1);
31     }
32
33     quick_sort2(nums, rm + 1, high);
34 }
35
36 fn main() {
37     let mut nums = [54,26,93,17,77,31,44,55,20];
38     let high = nums.len() - 1;
39     quick_sort2(&mut nums, 0, high);
40     println!("sorted nums: {:?}", nums);
41     // sorted nums: [17, 20, 26, 31, 44, 54, 55, 77, 93]
42
43     let mut nums = [1000,-1,2,-20,89,64,0,99,73];
44     let high = nums.len() - 1;
45     quick_sort2(&mut nums, 0, high);
46     println!("sorted nums: {:?}", nums);
47     // sorted nums: [-20, -1, 0, 2, 64, 73, 89, 99, 1000]
48 }

```

To partition a set of length n in quicksort, there will be $\log(n)$ partitions if the partition is always in the middle. However, finding the splitting point requires checking the pivot value against each of the n items, resulting in a complexity of $n\log(n)$. In the worst case, when the splitting point is far from the middle, sorting of items 1 and $n - 1$ will be repeated n times, leading to a complexity of $O(n^2)$.

Quicksort relies on recursion, but excessive depth can decrease its performance. To overcome this limitation, introsort switches to heap sort after the recursion depth exceeds $\log(n)$. For small numbers ($n < 20$), introsort switches to insertion sort. This mixed sorting algorithm can achieve the high performance of quicksort on regular datasets and maintain a performance of $O(n\log(n))$ in the worst case. Introsort is a built-in sorting algorithm in C++.

Quicksort divides the array to be sorted into two areas for sorting. However, if there are a large number of duplicate elements, quicksort will repeatedly compare them, resulting in wasted performance. To address this issue, the array can be divided into three areas for sorting (three-area quicksort). Duplicate elements are placed in the third area, and only the other two areas are sorted. The repeated data is chosen as the pivot value, and the less-than pivot value is put into the left area, the greater-than pivot value into the right area, and the equal pivot value into the middle area. Then, three-area quicksort is recursively called on the left and right areas.

7.5 The Insertion Sort

Insertion sort achieves sorting by iteratively inserting data items into a sorted subsequence. Although its performance is still $O(n^2)$, its approach is distinct from other sorting algorithms. It always maintains a sorted subsequence at the lower end of the dataset and inserts new items into the subsequence, expanding it until the entire collection is sorted.

The algorithm starts by assuming that the initial subsequence has only one item located at position 0. During the next traversal, items 1 to $n-1$ are compared with the first item. If an item is smaller than the first item, it is inserted before the first item. If it is larger than the first item, the subsequence is expanded by one. This process is repeated for the remaining unsorted items until the entire collection is sorted. In the following figure, the sorted region is shown in red, and the unsorted region in blue. Each row represents a sorting operation, and the bottom row shows the sorted result.

84	92	66	56	44	31	72	19	24	84 ordered
84	92	66	56	44	31	72	19	24	still ordered
66	84	92	56	44	31	72	19	24	insert 66
56	66	84	92	44	31	72	19	24	insert 56
44	56	66	84	92	31	72	19	24	insert 44
31	44	56	66	84	92	72	19	24	insert 31
31	44	56	66	72	84	92	19	24	insert 72
19	31	44	56	66	72	84	92	24	insert 19
19	24	31	44	56	66	72	84	92	insert 24

Figure 7.2: Insertion sort

```

1 // insertion_sort.rs
2 fn insertion_sort(nums: &mut [i32]) {
3     if nums.len() < 2 { return; }
4     for i in 1..nums.len() {
5         let mut pos = i;
6         let curr = nums[i];
7         while pos > 0 && curr < nums[pos-1] {
8             // move element to right
9             nums[pos] = nums[pos-1];
10            pos -= 1;
11        }
12
13        // insert element: curr
14        nums[pos] = curr;
15    }
16 }
17 fn main() {
18     let mut nums = [54,32,99,18,75,31,43,56,21];
19     insertion_sort(&mut nums);
20     println!("sorted nums: {:?}", nums);
21     // sorted nums: [18, 21, 31, 32, 43, 54, 56, 75, 99]
22 }

```

The insertion sort requires comparing each new element with the already sorted elements one by one. However, the binary search algorithm discussed in Chapter 6 can efficiently locate the position of an element in a sorted subsequence. Therefore, binary search can be utilized for acceleration.

```
1 // binary_insertion_sort.rs
2
3 fn binary_insertion_sort(nums: &mut [i32]) {
4     let mut temp;
5     let mut left;
6     let mut mid;
7     let mut right;
8
9     for i in 1..nums.len() {
10         left = 0;
11
12         // Sorted array boundaries
13         right = i - 1;
14
15         // Data to be sorted
16         temp = nums[i];
17
18         // Binary search finds the position of temp
19         while left <= right {
20             mid = (left + right) >> 1;
21             if temp < nums[mid] {
22                 // To prevent right = 0 - 1 case
23                 if 0 == mid {
24                     break;
25                 }
26                 right = mid - 1;
27             } else {
28                 left = mid + 1;
29             }
30         }
31
32         // Move data back
33         for j in (left..i-1).rev() {
34             nums.swap(j, j+1);
35         }
36
37         // Insert temp into the empty space
38         if left != i {
39             nums[left] = temp;
40         }
41     }
42 }
43
44 fn main() {
45     let mut nums = [1,3,2,8,6,4,9,7,5,10];
46     binary_insertion_sort(&mut nums);
47     println!("sorted nums: {:?}", nums);
48     // sorted nums: [1, 2, 3, 4, 5, 6, 7, 8, 9, 10]
49 }
```

7.6 The Shell Sort

Shell sort, also known as diminishing increment sort, is a sorting algorithm that divides the original set into smaller subsets and applies insertion sort to each subset. The way of selecting subsets is the key to Shell sort. Rather than evenly splitting the set into contiguous sublists, Shell sort selects items every few items to add to the subset, with the distance between them called the gap.

84	92	66	56	44	31	72	19	24
84	92	66	56	44	31	72	19	24
84	92	66	56	44	31	72	19	24

Figure 7.3: Shell sort

To understand Shell sort, consider the figure above. The set in the figure has nine items. If a gap of three is used, there will be a total of three subsets, each with three items of the same color. These separated elements can be considered as connected together, so insertion sort can be used to sort elements of the same color.

After sorting the subsets, the overall set is still unordered, as shown in the figure below. Although the overall set is unordered, it is not completely unordered. Subsets of the same color are ordered. By sorting the entire set using insertion sort, the set can be completely sorted quickly. The number of insertion sort moves is small at this point because adjacent items are in their own subsets' ordered positions, and these adjacent items are almost ordered. Therefore, only a few insertion moves are needed to complete the sorting.

56	19	24	72	44	31	84	92	66
----	----	----	----	----	----	----	----	----

Figure 7.4: Shell sort

In Shell sort, the increment is the key feature, and different increments can be used to determine the number of subsets. The gap value is adjusted continuously in the implementation of Shell sort to achieve sorting.

```

1 // shell_sort.rs
2
3 fn shell_sort(nums: &mut [i32]) {
4     // Internal function for insertion sort
5     // with elements exchanged distance is gap
6     fn ist_sort(nums: &mut [i32], start: usize, gap: usize) {
7         let mut i = start + gap;
8
9         while i < nums.len() {
10             let mut pos = i;
11             let curr = nums[pos];
12
13             while pos >= gap && curr < nums[pos - gap] {
14                 nums[pos] = nums[pos - gap];
15                 pos -= gap;

```

```

16         }
17
18         nums[pos] = curr;
19         i += gap;
20     }
21 }
22
23 // minimize the gap in every loop untill 1
24 let mut gap = nums.len() / 2;
25 while gap > 0 {
26     for start in 0..gap {
27         ist_sort(nums, start, gap);
28     }
29
30     gap /= 2;
31 }
32 }

```

The following are use cases.

```

1 fn main() {
2     let mut nums = [54,32,99,18,75,31,43,56,21,22];
3     shell_sort(&mut nums);
4     println!("sorted nums: {:?}", nums);
5     // sorted nums: [18, 21, 22, 31, 32, 43, 54, 56, 75, 99]
6
7     let mut nums = [1000,-1,2,-20,89,64,0,99,73];
8     shell_sort(&mut nums);
9     println!("sorted nums: {:?}", nums);
10    // sorted nums: [-20, -1, 0, 2, 64, 73, 89, 99, 1000]
11 }

```

Shell sort divides the original set into smaller subsets and applies insertion sort to each subset. The subsets are selected by choosing an item every few items with the distance between them called the gap. The set is then considered as connected together, and insertion sort is used to sort elements of the same color. The final insertion operation in Shell sort is much less than in insertion sort because the collection has been pre-sorted by earlier increments. This makes the overall sorting very efficient.

Although the complexity analysis of Shell sort is slightly more complicated, it is roughly between $O(n)$ and $O(n^2)$. By changing the gap value according to the formula $2^k - 1$ (1, 3, 7, 15, 31), the complexity of Shell sort is approximately $O(n^{\frac{3}{2}})$, which is very fast. Additionally, binary search can be used to further improve Shell sort, similar to insertion sort. However, the index processing in binary search needs to add the gap value. Readers are encouraged to think about implementing this improved algorithm.

7.7 The Merge Sort

The previously learned divide and conquer strategy can be used to improve the performance of sorting algorithms, specifically merge sort. Both merge sort and quicksort use recursion to continuously divide the list in half for sorting. If the set has only one item, it is considered sorted. If there are multiple items, the set is split, and recursive calls to merge sort are made for both intervals. Once sorted, a merge operation combines the two intervals into a single sorted set. This process is illustrated in the figures shown below.

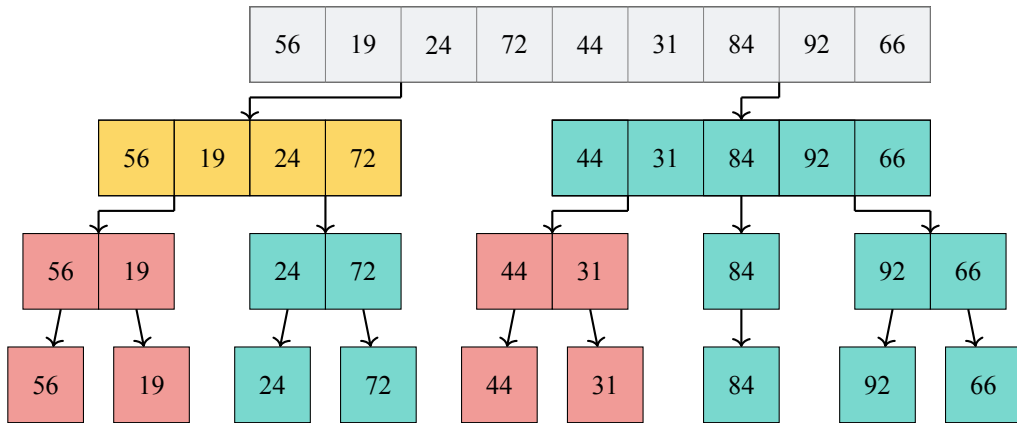


Figure 7.5: Split procedures of merge sort

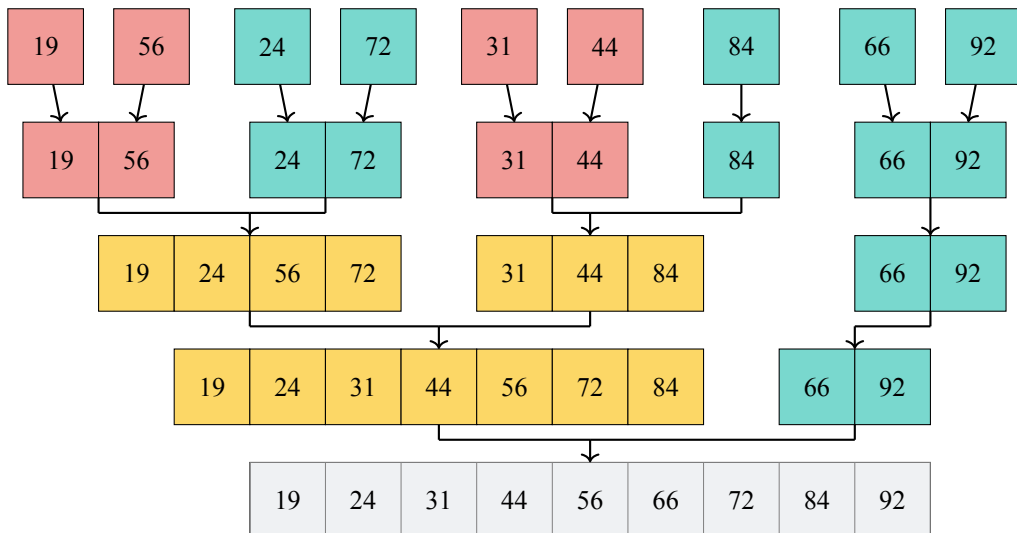


Figure 7.6: Merge procedures of merge sort

Merge sort breaks down the set into basic cases with one or two elements, allowing for easy direct comparison. Then, the merge process begins by first sorting the smallest subsequence of the basic case. Two-by-two merging is performed until the set is completely sorted. Even if the set is not evenly divided, the performance is not affected, as the difference is at most one element. The merge operation is straightforward, as each subsequence is already sorted, and only one comparison is necessary at a time. The resulting sequence is always ordered. Below is a Rust implementation of merge sort, consisting of two sorts and one merge operation, which is straightforward.

```

1 // merge_sort.rs
2
3 fn merge_sort(nums: &mut [i32]) {
4     if nums.len() > 1 {
5         let mid = nums.len() >> 1;
6         merge_sort(&mut nums[..mid]); // sort the first half
7         merge_sort(&mut nums[mid..]); // sort the last half
8         merge(nums, mid); // merge all

```

```

9      }
10   }
11
12   fn merge(nums: &mut [i32], mid: usize) {
13       let mut i = 0; // mark element in first half of data
14       let mut k = mid; // mark element in last half of data
15       let mut temp = Vec::new();
16
17       for _j in 0..nums.len() {
18           if k == nums.len() || i == mid {
19               break;
20           }
21
22           // put into a temp collection
23           if nums[i] < nums[k] {
24               temp.push(nums[i]);
25               i += 1;
26           } else {
27               temp.push(nums[k]);
28               k += 1;
29           }
30       }
31
32       // to make sure all data been solved
33       if i < mid && k == nums.len() {
34           for j in i..mid {
35               temp.push(nums[j]);
36           }
37       } else if i == mid && k < nums.len() {
38           for j in k..nums.len() {
39               temp.push(nums[j]);
40           }
41       }
42
43       // put temp data back to nums, finish sort
44       for j in 0..nums.len() {
45           nums[j] = temp[j];
46       }
47   }
48
49   fn main() {
50       let mut nums = [54,32,99,22,18,75,31,43,56,21];
51       merge_sort(&mut nums);
52       println!("sorted nums: {:?}", nums);
53       // sorted nums: [18, 21, 22, 31, 32, 43, 54, 56, 75, 99]
54   }

```

To analyze the time complexity of merge sort, we can divide the sorting process into two parts: sorting and merging. The sorting complexity is $O(\log_2(n))$, as we learned in the binary search section. The merging step involves placing each item in the set on the sorted list, with a maximum of n times, resulting in a complexity of $O(n)$. As recursive and merging operations are combined, the performance of merge sort is $O(n\log_2(n))$.

The space complexity of merge sort is relatively high at $O(n)$. To reduce space usage, we can consider optimizing merge sort using insertion sort. One approach is to use insertion sort directly when the length is less than a certain threshold (such as 64) and to use merge sort when the length is greater than the threshold. This algorithm is called insertion merge sort and improves the merge sort algorithm to some extent. However, implementation details are left to the reader.

7.8 The Selection Sort

Selection sort is a more efficient version of bubble sort, as it performs only one data exchange per round by seeking only the index of the maximum value during traversal. After completing the traversal, it swaps that maximum item into its correct position. Like bubble sort, after the first traversal, the maximum item is in the last position, and after the second traversal, the second-largest value is in the second-to-last position. To sort n items, $n-1$ passes are required.

Although selection sort and bubble sort have the same number of comparisons, selection sort is faster because of its reduced number of data exchanges. The time complexity of both algorithms is $O(n^2)$. The figure below shows the entire sorting process of selection sort, in which the maximum unsorted item is selected during each traversal and placed in the correct position. Unlike bubble sort, selection sort does not exchange pairs during the traversal.

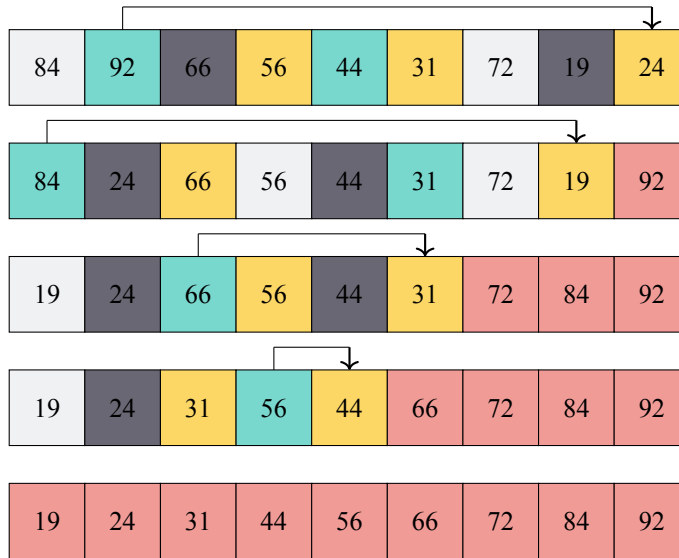


Figure 7.7: Selection sort

```

1 // selection_sort.rs
2
3 fn selection_sort(nums: &mut Vec<i32>) {
4     let mut left = nums.len() - 1; // left marker
5     while left > 0 {
6         let mut pos_max = 0;
7         for i in 1..=left {
8             if nums[i] > nums[pos_max] {
9                 pos_max = i; // current loop's max value
10            }
11        }
12        nums.swap(left, pos_max);

```

```

13         left -= 1; // unsorted elements reduced by 1
14     }
15 }
16
17 fn main() {
18     let mut nums = vec![54,32,99,18,75,31,43,56,21,22];
19     selection_sort(&mut nums);
20     println!("sorted nums: {:?}", nums);
21     // sorted nums: [18, 21, 22, 31, 32, 43, 54, 56, 75, 99]
22 }

```

Although bubble sort and selection sort have the same time complexity, there are still opportunities for optimization. One such optimization is to perform bidirectional sorting, which can be achieved by adopting the cocktail sort approach. Similarly, selection sort can also be modified to perform bidirectional sorting, which changes the coefficient of complexity but not the overall time complexity. Readers are encouraged to consider the implementation of this approach.

7.9 The Heap Sort

In addition to linear data structures like stacks and queues, non-linear data structures also exist in computers, one of which is the heap. A heap is a non-linear complete binary tree with left and right child nodes. The height of a tree with n nodes is $\log(n)$. Though trees are not studied until Chapter 8, comprehending them here can help understand the properties of heaps. There are two types of heaps: min-heaps and max-heaps. A min-heap has every node with a value less than or equal to its left and right child nodes, whereas a max-heap has every node with a value greater than or equal to its left and right child nodes. Figure (7.8) illustrates an example of a min-heap.

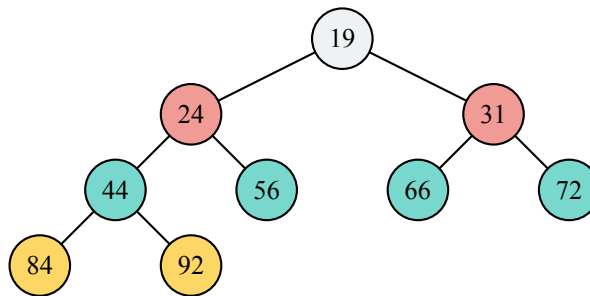


Figure 7.8: Min heap

Heap sort is a sorting algorithm designed using the heap data structure. It repeatedly selects the top element and moves it to the end, then rebuilds the heap to implement the sorting. It is a selection sort with worst-case, best-case, and average time complexity of $O(n\log_2(n))$ and is an unstable sort. As shown in the figure above, the heap is similar to a linked list with multiple connections. The nodes in the heap are numbered by layer, and if this logical structure is mapped to an array, it looks like the figure below, where the first position, index 0, is occupied by 0.

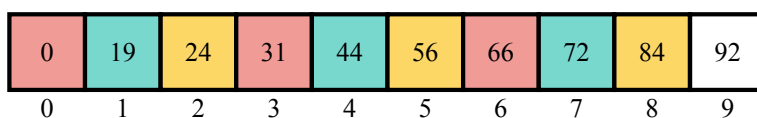


Figure 7.9: Array as a min heap

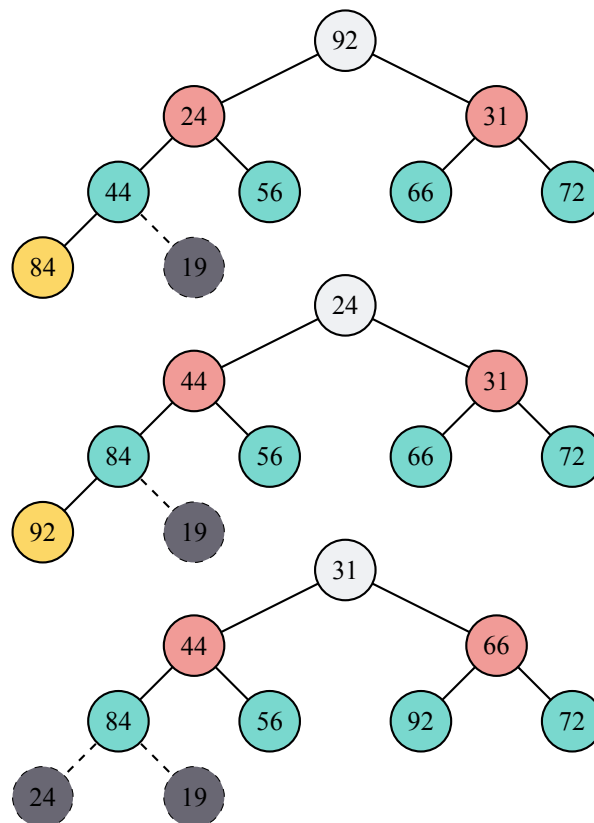
It is possible to represent heaps not only with trees, but also with arrays or Vecs (shown in figure above). In fact, using arrays or Vecs to represent heaps is more in line with the literal meaning of the word "heap," which refers to a collection of things gathered together. Note that our index starts from 1, which allows us to represent the indices of the left and right child nodes as $2i$ and $2i+1$, making it easier to calculate.

To satisfy the requirements of a binary tree's node relationships, a heap represented by an array should meet the following criteria:

- Max heap: $\text{arr}[i] \geq \text{arr}[2i]$ and $\text{arr}[i] \geq \text{arr}[2i+1]$
- Min heap: $\text{arr}[i] \leq \text{arr}[2i]$ and $\text{arr}[i] \leq \text{arr}[2i+1]$

Heap sort involves constructing an unsorted sequence into a min heap. The minimum value of the entire sequence is then the root node of the heap. Swap it with the last element of the sequence, and the last element becomes the minimum value. This minimum value is no longer considered in the heap, and the remaining $n-1$ elements are reconstructed into a heap, producing a new minimum value. Swap this minimum value to the end of the new heap, and you will have two sorted values. Repeat this process until the entire sequence is sorted. A min heap produces a descending order sorting, while a max heap produces an ascending order sorting.

To better illustrate the heap sort process, a figure is provided below. The dark gray color represents the minimum element, which has been replaced by 92 and is no longer part of the heap. When 92 is at the top of the heap, it is no longer a min heap, so a new min heap is constructed to make the minimum value 24 at the top of the heap. Then, 24 is swapped with the last element in the heap, and the second-to-last element becomes dark gray. Note that the dotted line indicates that this element no longer belongs to the heap. Continuing to swap, the dark gray subsequence gradually fills the entire heap in reverse order from the last level, achieving a reverse sorting from largest to smallest. To achieve sorting from smallest to largest, a max heap should be constructed, and the corresponding logic in the min heap should be modified.



Here is an implementation of heap sort.

```

1 // heap_sort.rs
2
3 // calculate index of parent node
4 macro_rules! parent {
5     ($child:ident) => {
6         $child >> 1
7     };
8 }
9
10 // calculate index of left child node
11 macro_rules! left_child {
12     ($parent:ident) => {
13         $parent << 1
14     };
15 }
16
17 // calculate index of right child node
18 macro_rules! right_child {
19     ($parent:ident) => {
20         ($parent << 1) + 1
21     };
22 }
23
24 fn heap_sort(nums: &mut [i32]) {
25     if nums.len() < 2 { return; }
26
27     let len = nums.len() - 1;
28     let last_parent = parent!(len);
29     for i in (1..=last_parent).rev() { // build a min-heap
30         move_down(nums, i);           // begins from 1
31     }
32
33     for end in (1..nums.len()).rev() { // rebuild min-heap
34         nums.swap(1, end);
35         move_down(&mut nums[..end], 1);
36     }
37 }
38
39 // move down greater element
40 fn move_down(nums: &mut [i32], mut parent: usize) {
41     let last = nums.len() - 1;
42     loop {
43         let left = left_child!(parent);
44         let right = right_child!(parent);
45         if left > last { break; }
46
47         // right <= last: determin if right child node exists
48         let child = if right <= last
49             && nums[left] < nums[right] {
50             right

```

```

51         } else {
52             left
53         };
54
55         // swap data if child node is greater than parent node
56         if nums[child] > nums[parent] {
57             nums.swap(parent, child);
58         }
59
60         // update parent and child relationship
61         parent = child;
62     }
63 }
64
65 fn main() {
66     let mut nums = [0,54,32,99,18,75,31,43,56,21,22];
67     heap_sort(&mut nums);
68     println!("sorted nums: {:?}", nums);
69     // sorted nums: [0, 18, 21, 22, 31, 32, 43, 54, 56, 75, 99]
70 }

```

The heap is a binary tree with a time complexity of $O(n\log(n))$. The time is mainly consumed in two parts: building the heap and adjusting the heap n times. Building the heap processes n elements with a complexity of $O(n)$. The longest path for each adjustment is from the root to the leaf, which is the height of the heap, $\log n$. Therefore, the time complexity of n adjustments is $O(n\log(n))$, and the final total time complexity is $O(n\log(n))$. Macros are used here to obtain node indexes, and functions can also be used to implement them. Heap sort and selection sort are similar in that they both find the maximum or minimum value in a set.

7.10 The Bucket Sort

The previously discussed different sorting techniques, including comparison-based algorithms. However, there are also sorting algorithms that do not involve comparisons, such as bucket sort, counting sort, and radix sort. Non-comparison sorting works by determining the number of elements that come before each element in the sorted set. For example, given a set of numbers "nums", calculating the number of elements before "nums[i]" uniquely determines its position in the sorted set. Non-comparison sorting can be completed in a single pass with a time complexity of $O(n)$, as it only requires counting the number of elements before each element.

Although non-comparison sorting has a low time complexity, it requires additional space to determine the position, making it suitable for specific data sizes and distributions (especially numbers). Non-comparison sorting requires indexable information within the data to determine the position. In contrast, comparison sorting is suitable for data of different sizes and distributions, making it applicable in various sorting situations.

Bucket sorting is one type of non-comparison sorting, where buckets are similar to slots in a hash table, but can hold multiple elements. The basic idea of bucket sorting are:

- Step 1: Divide the elements into different buckets, setting the number of buckets to k after finding $\max V$ and $\min V$, and dividing the interval $[\min V, \max V]$ into k equal intervals. Elements in the sequence are hashed to each bucket using a hash function (such as taking the remainder).
- Step 2: Sort the elements in each bucket with any available sorting algorithm.
- Step 3: Merge the ordered elements in each bucket into a large ordered set.

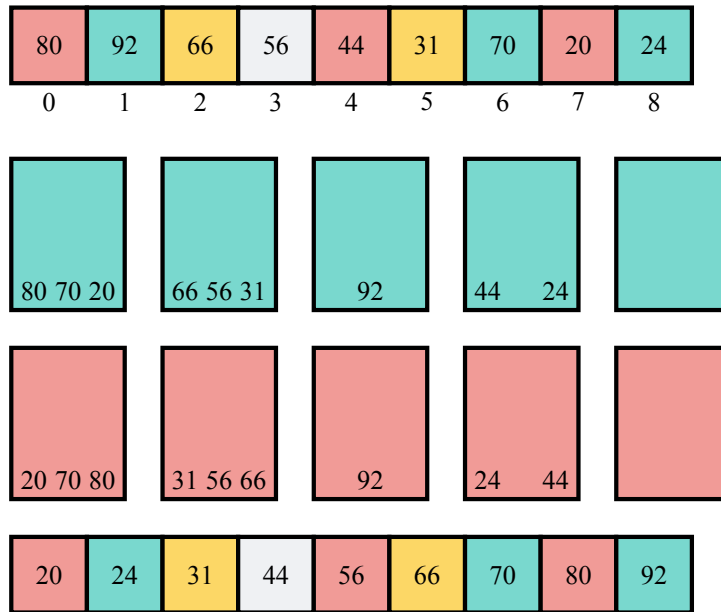
In Rust, a bucket can be defined as a structure that contains a hash function named `hasher` and a dataset named `values`.

```

1 // bucket_sort.rs
2
3 struct Bucket<H, T> {
4     hasher: H,      // hasher: a function, recieved when called
5     values: Vec<T>, // values: a container for data
6 }

```

To help readers understand bucket sort, the figure below illustrates the process of hashing data into buckets, sorting within each bucket, and merging the sorted data to obtain the final ordered set.



Here is an implementation of bucket sort.

```

1 // bucket_sort.rs
2
3 use std::fmt::Debug;
4
5 impl<H, T> Bucket<H, T> {
6     fn new(hasher: H, value: T) -> Bucket<H, T> {
7         Bucket {
8             hasher: hasher,
9             values: vec![value]
10        }
11    }
12 }
13
14 // Bucket sort, Debug feature is used for printing T
15 fn bucket_sort<H, T, F>(nums: &mut [T], hasher: F)
16     where H: Ord,
17           T: Ord + Clone + Debug,
18           F: Fn(&T) -> H,
19 {
20     let mut buckets: Vec<Bucket<H, T>> = Vec::new();

```



```

21     for val in nums.iter() {
22         let hasher = hasher(&val);
23
24         // Binary search and sort bucket data
25         match buckets.binary_search_by(|bct|
26             bct.hasher.cmp(&hasher)) {
27             Ok(idx) => buckets[idx].values.push(val.clone()),
28             Err(idx) => buckets.insert(idx,
29                 Bucket::new(hasher, val.clone())),
30         }
31     }
32
33     // Split the buckets and merge them into a single Vec
34     let ret = buckets.into_iter().flat_map(|mut bucket| {
35         bucket.values.sort();
36         bucket.values
37     }).collect::<Vec<T>>();
38
39     nums.clone_from_slice(&ret);
40 }
41
42 fn main() {
43     let mut nums = [0,54,32,99,18,75,31,43,4,56,21,22,1,100];
44     bucket_sort(&mut nums, |t| t / 5);
45     println!("{}", nums);
46     // [0, 1, 4, 18, 21, 22, 31, 32, 43, 54, 56, 75, 99, 100]
47 }

```

Implementing bucket sort is more complex than comparison-based sorting algorithms because it involves creating a bucket structure and implementing a sorting algorithm that operates on this structure. In this implementation, the data is distributed into buckets using division by 5. If another value is used, the number of buckets needs to be adjusted accordingly.

Bucket sort has a time complexity of $O(n + n \log n - n \log k)$, where n is the number of elements to be sorted and k is the number of buckets used. Assuming the data is uniformly distributed, the average number of elements in each bucket is n/k . If quicksort is used to sort each bucket, the time complexity of each sorting operation is $O(n/k \log(n/k))$. Therefore, the total time complexity can be expressed as $O(n) + kO(n/k \log(n/k)) = O(n + n \log(n/k))$. When k is close to n , bucket sorting can be regarded as having a time complexity of $O(n)$.

One drawback of bucket sort is that it may create too many unnecessary buckets, resulting in wasted space. To address this issue, FlashSort is an optimized bucket sorting algorithm that estimates the number of buckets needed based on the number of elements to be sorted. If the number of buckets is greater than the number of elements, the number of buckets can be calculated by $m = f * n$, where f is a decimal such as 0.2. The rule for placing elements into buckets is defined by equation shown below, where $K(A_i)$ denotes the bucket index of element A_i .

$$K(A_i) = 1 + \text{int}((m - 1) \frac{A_i - A_{\min}}{A_{\max} - A_{\min}}) \quad (7.2)$$

Assuming each bucket has an average of n/m elements, using insertion sort for each bucket results in a total time complexity of $O(n^2/m)$. Experimental results show that FlashSort performs optimally at $O(n)$ when $m = 0.42n$. Moreover, when $m = 0.1n$, FlashSort is faster than quicksort as long as $n > 80$, and when $n = 10000$, it is twice as fast. Additionally, when $m = 0.2n$, it is 15% faster than when $m = 0.1n$. Even when m is as low as $0.05n$, FlashSort is still significantly faster than quicksort when $n > 200$.

7.11 The Counting Sort

Another type of non-comparison sorting is counting sort, which is a special case of bucket sort that only handles data of the same type, and therefore requires more space. The basic idea of counting sort is as follows:

- First, initialize a counter set of length $\text{maxV} - \text{minV} + 1$, where maxV is the maximum value in the set to be sorted, and minV is the minimum value. Set all values in the counter set to 0.
- Second, scan the set to be sorted and use the current value minus minV as the index in the counter set, and increment the counter at this index.
- Third, scan the counter set and write the values back to the original set in order to complete the sorting.

For example, given `nums=[0,7,1,7,3,1,5,8,4,4,5]`, first traverse `nums` to obtain the minimum and maximum values, $\text{maxV}=8$, $\text{minV}=0$, and then initialize a counter set of length $8 - 0 + 1$.

$$\text{counter} = [0, 0, 0, 0, 0, 0, 0, 0, 0]$$

Then scan `nums` and calculate the index by subtracting minV from the current value. For example, when scanning 0, the index is $0 - 0 = 0$, so the value at index 0 of counter is incremented by 1. At this point, counter is `[1,0,0,0,0,0,0,0,0]`. Continue scanning until the final value of counter is obtained.

$$[1, 2, 0, 1, 2, 2, 0, 2, 1]$$

When traversing the counter set, if a value at a certain index is not 0, write the corresponding index value into `nums`, and decrement the value in counter. For example, if the value at the first position 0 of counter is 1, indicating that there is one 0 in `nums`, write it into `nums`. Continuing, the value at index 1 is 2, indicating that there are two 1s in `nums`, write them into `nums`. Finally, `nums` is sorted as:

$$[0, 1, 1, 3, 4, 4, 5, 5, 7, 8]$$

The sorting process is completed by scanning the counter set, and the sorting process does not involve comparison, exchange, or other operations, making it very fast. Here is an implementation of counting sort:

```

1 // counting_sort.rs
2
3 fn counting_sort(nums: &mut [usize]) {
4     if nums.len() <= 1 {
5         return;
6     }
7
8     // bucket number is the maximum value in nums plus 1
9     let max_bkt_num = 1 + nums.iter().max().unwrap();
10
11     // save the number of each value in nums
12     let mut counter = vec![0; max_bkt_num];
13     for &v in nums.iter() {
14         counter[v] += 1;
15     }
16
17     // write data back to original nums slice
18     let mut j = 0;
19     for i in 0..max_bkt_num {
20         while counter[i] > 0 {
21             nums[j] = i;

```

```

22         counter[i] -= 1;
23         j += 1;
24     }
25 }
26 }
27
28 fn main() {
29     let mut nums = [54,32,99,18,75,31,43,56,21,22];
30     counting_sort(&mut nums);
31     println!("sorted nums: {:?}", nums);
32     // sorted nums: [18, 21, 22, 31, 32, 43, 54, 56, 75, 99]
33 }

```

7.12 The Radix Sort

The third type of non-comparison sorting is radix sort, which uses the radix rule of positive numbers to sort. The sorting process involves collecting and distributing data, and the specific steps are as follows:

- Step 1: Find the maximum value in the given set of integers, and determine the number of digits in it. Then, pad zeros to the left of the numbers to unify the number of digits.
- Step 2: Starting from the lowest digit, perform stable sorting, collect the numbers according to their digits, and then sort the numbers according to the next higher digit until the sorting is complete.

For instance, consider an integer sequence [1, 134, 532, 45, 36, 346, 999, 102]. The maximum value in the set is 999, which has three digits. Therefore, three rounds of sorting are required for the ones, tens, and hundreds digits. Firstly, pad zeros to the left of the numbers to make all numbers have three digits, as shown in the second column of the figure. Then, sort the numbers according to their ones digit, tens digit, and hundreds digit in order. After three rounds of sorting, the data is sorted. The red digit in the figure represents the digit being sorted in the current round, and it is apparent that the numbers are sorted based on their corresponding digit at each iteration.

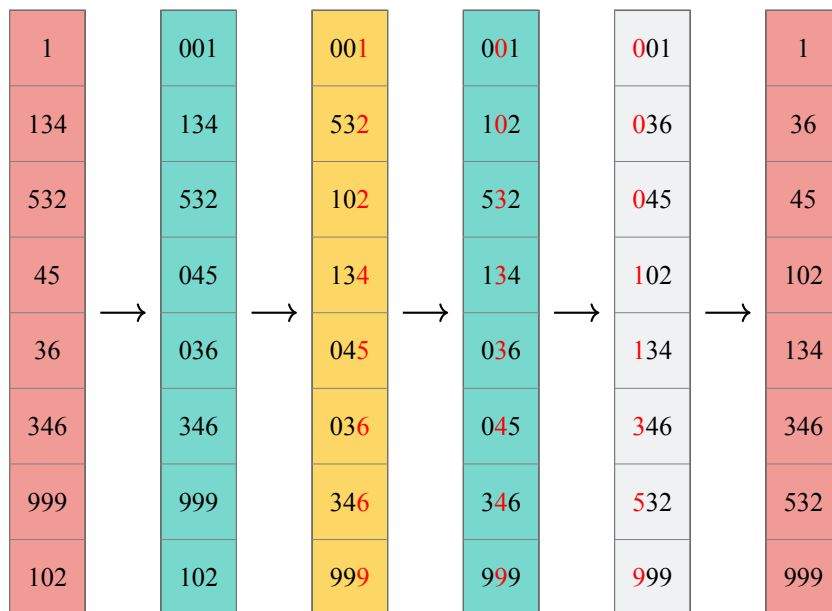


Figure 7.10: Radix sort

```

1 // radix_sort.rs
2
3 fn radix_sort(nums: &mut [usize]) {
4     if nums.len() <= 1 { return; }
5
6     // Find the largest number, which has the most digits.
7     let max_num = match nums.iter().max() {
8         Some(&x) => x,
9         None => return,
10    };
11
12    // Find the power of 2 that is greater than or equals to
13    // the length of nums as the bucket size. For example,
14    // the closest and greater power of 2 to 10 is 2^4 = 16,
15    // the closest and greater power of 2 to 17 is 2^5 = 32.
16    let radix = nums.len().next_power_of_two();
17
18    // The variable 'digit' represents the count of numbers
19    // in a bucket that are less than a certain digit.
20    // The ones, tens, hundreds, and thousands digits
21    // correspond to positions 1, 2, 3, and 4, respectively.
22    // The counting starts from the ones digit, so it is 1.
23    let mut digit = 1;
24    while digit <= max_num {
25        // Calculate the position of the data in the bucket.
26        let index_of = |x| x / digit % radix;
27        let mut counter = vec![0; radix];
28        for &x in nums.iter() {
29            counter[index_of(x)] += 1;
30        }
31        for i in 1..radix {
32            counter[i] += counter[i-1];
33        }
34
35        // sorting
36        for &x in nums.to_owned().iter().rev() {
37            counter[index_of(x)] -= 1;
38            nums[counter[index_of(x)]] = x;
39        }
40
41        // process next bucket
42        digit *= radix;
43    }
44 }
45
46 fn main() {
47     let mut nums = [0,54,32,99,18,75,31,43,56,21,22,100];
48     radix_sort(&mut nums);
49     println!("sorted nums: {:?}", nums);
50     // sorted nums: [18, 21, 22, 31, 32, 43, 54, 56, 75, 99]
51 }

```

To sort by the same digit, stable sorting is used because it preserves the previous sorting result. For instance, sorting the tens digit preserves the sorting result of the ones digit, and sorting the hundreds digit preserves the sorting result of the tens digit. Binary can also be used to solve any non-negative integer sequence with the radix sorting algorithm. The time complexity is $O(64n)$ assuming the maximum integer in the sequence is 64 bits. Although comparison-based sorting algorithms have a time complexity of $O(n \log(n))$, they are still faster than radix sorting since the coefficient of 64 is too large to be practical. To use binary, $k=2$ is the smallest and the number of digits is the largest, so the time complexity becomes $O(nd)$ and the space complexity becomes $O(n + k)$, which is smaller. Conversely, when using the maximum value as the radix, $k=\max V$ is the largest and the number of digits is the smallest, resulting in a smaller time complexity of $O(nd)$ but an increased space complexity of $O(n + k)$, leading to radix sorting degenerating into counting sorting.

In summary, these three non-comparison sorting methods (counting sorting, bucket sorting, and radix sorting) are interconnected. Counting sorting is a specific case of bucket sorting, and radix sorting degenerates into counting sorting if it uses the minimum number of digits to sort. Bucket sorting is suitable for evenly distributed elements, counting sorting requires a small difference between $\max V$ and $\min V$, and radix sorting can only handle positive numbers with $\max V$ and $\min V$ being as close as possible. Therefore, these three sorting methods are only applicable for sorting small amounts of data, ideally less than 10000.

7.13 The Tim Sort

The TimSort hybrid sorting algorithm, proposed by Tim Peters in 2002, combines multiple sorting methods to address the limitations of individual algorithms. TimSort is efficient, stable, and adaptive to data distribution, making it superior to most sorting algorithms. It is currently the default sorting algorithm in many programming languages and platforms, including Java, Python, Rust etc.

TimSort is a hybrid stable sorting algorithm that uses merge and insertion sorts. When the number of elements to be sorted is less than 64, TimSort calls insertion sort directly. However, when the number of elements is greater than 64, TimSort combines insertion and merge sorts.

In practice, the data to be sorted often contains partially ordered blocks, where some parts of the data are already sorted in ascending or descending order, as shown in the figure below. TimSort leverages this characteristic of the data to sort it, and the existence of partially ordered blocks is the core of TimSort.

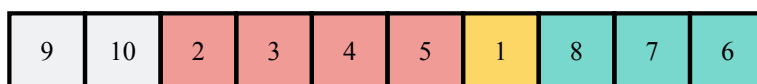


Figure 7.11: Partially ordered data

To sort the set of elements, TimSort divides them into runs or partitions. These runs can be seen as individual computational units. TimSort iterates through the elements and places them in different runs while merging the runs according to certain rules. This merging continues until only one run remains, which is the sorted result. To set the appropriate partition size, TimSort sets the minrun parameter. Each partition cannot have fewer elements than minrun. If a partition has fewer elements than minrun, the number of runs is expanded to minrun using insertion sort and then merged.

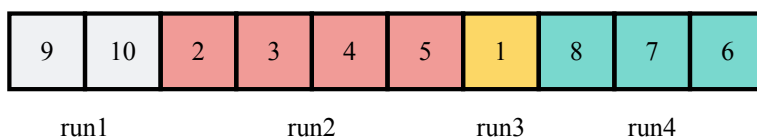


Figure 7.12: Data Runs

The approximate process of TimSort is as follows:

- Scan the set to be sorted to determine whether the number of elements is greater than 64.
- If it is less than or equal to 64, use the insertion sort algorithm to sort the data and return.
- If it is greater than 64, scan the set, calculate the minrun, and find various ordered blocks.
- Merge the blocks in pairs. If an individual block has fewer elements than minrun, use insertion sort to expand it.

- Repeat this process until only one block remains, which is the sorted set.

TimSort is a sorting algorithm that can be completed by addressing three important points: determining the minrun value, finding ordered blocks, and expanding and merging small blocks.

The first point is to determine the value of minrun. TimSort selects the first six binary digits of the length n of the collection to be sorted as the minrun. If the remaining digits are not 0, then minrun is incremented by 1. The range of minrun is from 32 to 64.

- For a collection of length $n = 189$, its binary representation is: 10111101. The first six digits, 101111, equal 47, and the remaining digits, 01, lead to a minrun value of 48.

- For a collection of length $n = 976$, its binary representation is: 1111010000. The first six digits, 111101, equal 61, and the remaining digits, 0000, lead to a minrun value of 61.

In fact, the maximum value of the first six binary digits is $111111 = 63$, and the minimum is $100000 = 32$. Therefore, the minrun value ranges from 32 to 64.

The second point is to find ordered blocks. TimSort finds ordered blocks by identifying increasing or decreasing sequences in the collection. This is achieved by an algorithm that only needs to judge the compare relationship of consecutive data to determine whether it is a block.

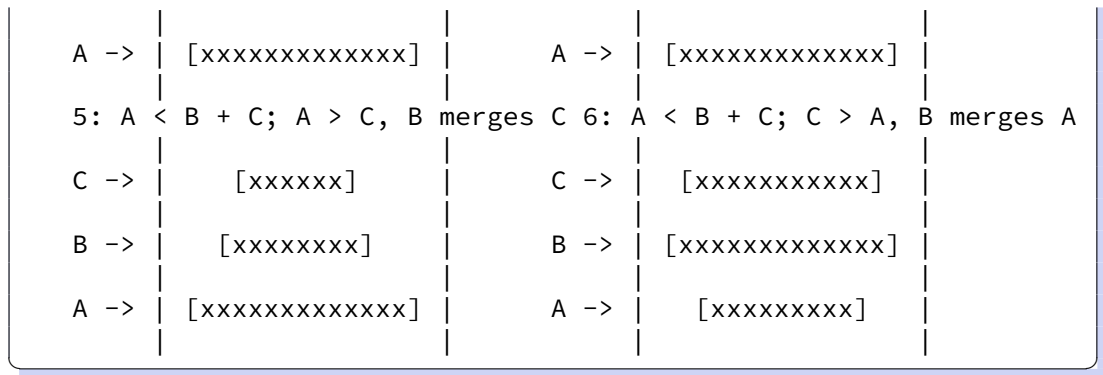
The third point is to expand and merge small blocks. Once the minrun value is calculated, TimSort adjusts any blocks that are in reverse order in place to be in the correct order. If the block has fewer elements than minrun, insertion sort is used to expand the block until it reaches minrun. The algorithm then checks the relationship between the current block and the previous two blocks. If the length of the three blocks does not satisfy $A > B + C$, $B > C$, then A and B or B and C are merged using insertion sort depending on the situation. If not, the algorithm continues to find the next block and repeat the block expansion and merging. Finally, the collection becomes several ordered blocks, and these merged blocks have lengths from longest to shortest. After the block division is completed, the small blocks are merged in pairs from the end of the collection to the beginning, and the collection becomes a large block, indicating that the sorting is completed.

The first two points are clear, but the third point may require further explanation. Reversing the order of a block is a simple process. To expand a block, new elements are inserted into the current block. However, merging blocks can be complicated as there are six possible scenarios.

When there are only two blocks, the algorithm only needs to consider the condition of $A > B$. When there are three blocks, the algorithm checks whether $A > B + C$ and $B > C$. Here are the illustrations of the six cases, where A is at the bottom of the stack, and B or C is at the top:

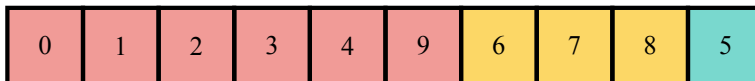
- $A > B$: Do not merge.
- $A > B + C$ and $B > C$: Do not merge.
- $A < B + C$ and $A > C$: Merge run B with C.
- $A < B$: Merge run A with B.
- $A > B + C$ and $B < C$: Merge run B with C.
- $A < B + C$ and $A < C$: Merge run A with B.

1: $A > B$, no merge		2: $A < B$, B merges A	
B ->	[xxxxxxxx]	B ->	[xxxxxxxxxxxxxx]
A ->	[xxxxxxxxxxxxxxxx]	A ->	[xxxxxxxx]
3: $A > B + C$; $B > C$, no merge		4: $A > B + C$; $B < C$, B merges C	
C ->	[xxx]	C ->	[xxxxxxx]
B ->	[xxxxxxxx]	B ->	[xxx]



The figure depicts the merging of three ordered blocks with a minimum run of 3. The characters A, B, and C represent the run blocks [xx], while the vertical lines represent a temporary stack used to merge these blocks. To determine whether and how to merge the blocks, the lengths of the three blocks are compared. The ideal merged state is shown in cases 1 and 3, while the other four merging scenarios are used to approximate these two cases.

The condition that $A > B + C$, $B > C$ is necessary to ensure efficient merging. For example, merging the block [6,7,8] directly with [0,1,2,3,4,9,5] would result in two remaining blocks with a large difference in length, making insertion sort inefficient. In contrast, when the blocks have similar lengths, merging them is very fast. Therefore, in this case, the blocks are not merged and are processed separately before being merged in reverse order at the end: [6,7,8] and [5], followed by [0,1,2,3,4,9] and [5,6,7,8].



To simplify the sorting process, a minimum run length is set, below which binary insertion sort is used. If the run length equals 64, minrun is also set to 64. Otherwise, a value between 32-64 is chosen for minrun such that $k = \frac{n}{minrun}$ is less than or equal to a power of 2. k represents the number of remaining blocks after scanning and processing, and their lengths must be sorted from largest to smallest. This enables merging from the tail, resulting in longer blocks and faster merging, akin to binary search. Hence, k is required to be less than or equal to a power of 2.

1	2	4	7	8	23	19	16	14	13	12	10	20	18	17	15	11	9	0	5	6	1	3	21	22
2	2	4	7	8	23	19	16	14	13	12	10	20	18	17	15	11	9	0	5	6	1	3	21	22
3	2	4	7	8	23	19	16	14	13	12	10	20	18	17	15	11	9	0	5	6	1	3	21	22
4	2	4	7	8	23	10	12	13	14	16	19	20	18	17	15	11	9	0	5	6	1	3	21	22
5	2	4	7	8	10	12	13	14	16	19	23	20	18	17	15	11	9	0	5	6	1	3	21	22
6	2	4	7	8	10	12	13	14	16	19	23	20	18	17	15	11	9	0	5	6	1	3	21	22
7	2	4	7	8	10	12	13	14	16	19	23	9	11	15	17	18	20	0	5	6	1	3	21	22
8	2	4	7	8	10	12	13	14	16	19	23	9	11	15	17	18	20	0	5	6	1	3	21	22
9	2	4	7	8	10	12	13	14	16	19	23	9	11	15	17	18	20	0	1	3	5	6	21	22
10	2	4	7	8	10	12	13	14	16	19	23	9	11	15	17	18	20	0	1	3	5	6	21	22
11	2	4	7	8	10	12	13	14	16	19	23	9	11	15	17	18	20	0	1	3	5	6	21	22
12	2	4	7	8	10	12	13	14	16	19	23	0	1	3	5	6	9	11	15	17	18	20	21	22
13	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23

Figure 7.13: TimSort

To illustrate the block expansion and merging mechanism, diagrams are presented above. For this example, $\text{minrun} = 5$, and each row represents an operation cycle. The leftmost column indicates the cycle number, and the square box contains the elements to be sorted.

The TimSort algorithm consists of several rounds of processing to sort a given set of data. In the first round, the algorithm obtains the data to be sorted from its parameters. In the second round, it searches for partitions and identifies a partition of length equal to the minimum run value, which is a predetermined set of values [2,4,7,8,23]. In the third round, it searches for the next partition and finds an inversion [19,16,14,13,12,10], which is then corrected to be in order as values [10,12,13,14,16,19] in the fourth round.

In the fifth round, the algorithm compares the length of the current partition to that of the previous partition and merges them using insertion sort if the current partition is longer. The result is [2,4,7,8,10,12,13,14,16,19,23], a new partition that is a combination of the two previous partitions. In the sixth round, it finds the next partition [20,18,17,15,11,9], detects an inversion, and corrects it to be in order in the seventh round. It then compares its length to the previous partition and continues searching for the next partition since it is shorter.

In the eighth round, the algorithm finds a new partition [0,5,6] that is shorter than the minimum run value and expands it using insertion sort. It then compares its length to the previous two partitions and determines that they satisfy certain conditions, which means that they do not need to be merged. This process continues until all the partitions are complete, which occurs after ten rounds.

In the eleventh round, the algorithm starts merging the smaller partitions from the end to the beginning ([0,1,3,5,6] and [21, 22]). In the twelfth round, it merges the two remaining partitions [9,11,15,17,18,20] and [0,1,3,5,6,21,22], which are of similar length. Finally, in the thirteenth round, the algorithm performs the final merge to obtain the sorted output.

To implement TimSort, the process can be broken down into several steps. Firstly, the original data list and the minimum number of elements required for merging (MIN_MERGE) need to be prepared. Next, the starting position of each run and its corresponding run can be obtained through the partition process. After that, since merge sorting involves temporary stacks and processing of two runs, these data related to sorting tasks can be implemented in a structure. Finally, the actual merging process can be performed from the small partitions at the end to the beginning, with longer partitions merged with shorter ones and the partitions of decreasing length for easier merging.

```

1 // tim_sort_without_gallop.rs
2
3 // The minimum length of a sequence involved in merging,
4 // shorter than which insertion sort is used.
5 const MIN_MERGE: usize = 64;
6
7 // Sorting state structure
8 struct SortState<'a> {
9     list: &'a mut [i32],
10     runs: Vec<Run>, // store all run(s)
11     pos: usize,
12 }
13
14 // Define the Run entity to save the starting index and
15 // interval length of the run in the list.
16 #[derive(Debug, Copy, Clone)]
17 struct Run {
18     pos: usize,
19     len: usize,
20 }
21
```



```

22 // merge_lo: a sorting state structure, used for merging
23 // sorting A and B.
24 struct MergeLo<'a> {
25     list_len: usize,      // The length of the data
26     first_pos: usize,     // first position of run1
27     first_len: usize,     // length of run1
28     second_pos: usize,    // first position of run2
29     dest_pos: usize,      // index of the sorted result.
30     list: &'a mut [i32], // partial interval of the data
31                          // to be sorted
32     temp: Vec<i32>,       // with a length set as the minimum
33                          // value between the lengths of
34                          // run1 and run2.
35 }
36
37 // merge_hi: a sorting state structure, used for merging
38 // sorting B and C.
39 struct MergeHi<'a> {
40     first_pos: isize,
41     second_pos: isize,
42     dest_pos: isize,
43     list: &'a mut [i32],
44     temp: Vec<i32>,       // used for memory alignment
45 }

```

For blocks with less than `MIN_MERGE` elements, binary insertion sort can be used to speed up the sorting process. This method has already been implemented in the previous section, so no further code will be provided.

When using TimSort, the first step is to calculate the minrun value and identify the sorted runs (including reverse runs). If necessary, the reverse runs should be converted to regular runs.

```

1 // tim_sort_without_gallop.rs
2
3 // Calculate minrun, actual range is [32, 64]
4 fn calc_minrun(len: usize) -> usize {
5     // If any bit in the low bits of len is 1, r is set to 1
6     let mut r = 0;
7     let mut new_len = len;
8     while new_len >= MIN_MERGE {
9         r |= new_len & 1;
10        new_len >>= 1;
11    }
12
13    new_len + r
14 }
15
16 // Calculate the starting index of the run and convert
17 // the reverse run to a forward run
18 fn count_run(list: &mut [i32]) -> usize {
19     let (ord, pos) = find_run(list);
20     if ord { // convert reverse run to a forward run
21         list.split_at_mut(pos).0.reverse();
22     }

```

```

23
24     pos
25 }
26
27 // Determine whether the relationship between list[i]
28 // and list[i+1] is ascending or descending and
29 // return the inflection point index
30 fn find_run(list: &[i32]) -> (bool, usize) {
31     let len = list.len();
32     if len < 2 {
33         return (false, len);
34     }
35
36     let mut pos = 1;
37     if list[1] < list[0] {
38         // descending: list[i+1] < list[i]
39         while pos < len - 1 && list[pos + 1] < list[pos] {
40             pos += 1;
41         }
42         (true, pos + 1)
43     } else {
44         // ascending: list[i+1] >= list[i]
45         while pos < len - 1 && list[pos + 1] >= list[pos] {
46             pos += 1;
47         }
48         (false, pos + 1)
49     }
50 }

```

Next, to sort the `SortState`, you will need to implement a constructor and a sort function. Finally, when the length of the partitions does not meet the requirements, partition merging needs to be performed using merge sort.

```

1 // tim_sort_without_gallop.rs
2
3 impl<'a> SortState<'a> {
4     fn new(list: &'a mut [i32]) -> Self {
5         SortState {
6             list: list,
7             runs: Vec::new(),
8             pos: 0,
9         }
10    }
11
12    fn sort(&mut self) {
13        let len = self.list.len();
14        // calculate the minrun
15        let minrun = calc_minrun(len);
16
17        while self.pos < len {
18            let pos = self.pos;
19            let mut run_len = count_run(self.list
20                                     .split_at_mut(pos)

```

```

21                                     .1);
22
23         // check if the remaining number of elements is
24         // less than minrun, if so,
25         // let run_minlen = len - pos
26         let run_minlen = if minrun > len - pos {
27             len - pos
28         } else {
29             minrun
30         };
31
32         // If the run is very short, extend its length
33         // to run_minlen, and the extended run needs to
34         // be sorted, so use binary insertion sort
35         if run_len < run_minlen {
36             run_len = run_minlen;
37             let left = self.list
38                 .split_at_mut(pos).1
39                 .split_at_mut(run_len).0;
40             binary_insertion_sort(left);
41         }
42
43         // Stack the runs, with each run having a
44         // different length
45         self.runs.push(Run {
46             pos: pos,
47             len: run_len,
48         });
49
50         // Find the next run position
51         self.pos += run_len;
52
53         // Merge runs that do not conform to the
54         // A > B + C and B > C rules
55         self.merge_collapse();
56     }
57
58
59     // Forcefully merge all remaining runs from the top
60     // of the stack until only one run remains,
61     // completing the tim_sort sort
62     self.merge_force_collapse();
63 }
64
65 // Merge runs to satisfy A > B + C and B > C
66 // If A <= B + C, merge B with the shorter of A and C
67 // If only A and B, and A <= B, merge A and B
68 fn merge_collapse(&mut self) {
69     let runs = &mut self.runs;
70     while runs.len() > 1 {
71         let n = runs.len() - 2;
72

```

```

73         // check the relationships between A, B, C, and D
74         // with D to prevent special cases of bugs
75         // A <= B + C || D <= A + B
76         if (n >= 1 && runs[n - 1].len
77             <= runs[n].len + runs[n + 1].len)
78             || (n >= 2 && runs[n - 2].len
79                 <= runs[n].len + runs[n - 1].len)
80         {
81             // Determine the length relationship between
82             // three consecutive runs A, B, and C,
83             // and merge n-1 corresponding to A,
84             // n corresponding to B, and n+1
85             // corresponding to C
86             let (pos1, pos2) = if runs[n-1].len
87                               < runs[n+1].len {
88                 (n - 1, n) // merge A and B
89             } else {
90                 (n, n + 1) // merge B and C
91             };
92
93             // Take out run1 and run2 to be merged
94             let (run1, run2) = (runs[pos1], runs[pos2]);
95             debug_assert_eq!(run1.pos+run1.len, run2.pos);
96
97             // Merge run to run1, i.e., update run1
98             // and delete run2,
99             // run1 index remains unchanged, and length
100            // becomes the len(run1) + len(run2)
101            runs.remove(pos2);
102            runs[pos1] = Run {
103                pos: run1.pos,
104                len: run1.len + run2.len,
105            };
106
107            // Take the merged run1 for merge sort
108            let new_list = self.list
109                .split_at_mut(run1.pos).1
110                .split_at_mut(run1.len + run2.len).0;
111            merge_sort(new_list, run1.len, run2.len);
112        } else {
113            break;
114        }
115    }
116 }
117
118 // Forcefully merge any remaining runs until
119 // only one run remains
120 fn merge_force_collapse(&mut self) {
121     let runs = &mut self.runs;
122     while runs.len() > 1 {
123         let n = runs.len() - 2;
124

```

```

125         // Check the length relationship between three
126         // consecutive runs A, B, and C,
127         // and merge n-1 corresponding to A,
128         // n corresponding to B, n+1 corresponding to C
129         let (pos1, pos2) = if n > 0
130             && runs[n - 1].len < runs[n + 1].len {
131             (n - 1, n)
132         } else {
133             (n, n + 1)
134         };
135
136         // Take out run1 and run2 to be merged
137         let (run1, run2) = (runs[pos1], runs[pos2]);
138         debug_assert_eq!(run1.len, run2.pos);
139
140         // Merge run to run1, i.e., update run1
141         // and delete run2,
142         // run1 index remains unchanged, and length
143         // becomes the sum of the lengths of run1 and run2
144         runs.remove(pos2);
145         runs[pos1] = Run {
146             pos: run1.pos,
147             len: run1.len + run2.len,
148         };
149
150         // Take the merged run1 for merge sort
151         let new_list = self.list
152             .split_at_mut(run1.pos).1
153             .split_at_mut(run1.len + run2.len).0;
154         merge_sort(new_list, run1.len, run2.len);
155     }
156 }
157 }

```

There are six cases of partitioning, which may require merging A and B or merging B and C. Since A, B, and C are contiguous in memory, merging A and B can be implemented separately from merging B and C, taking advantage of their relative positions.

```

1 // tim_sort_without_gallop.rs
2
3 // merge sort for A, B and C
4 fn merge_sort(
5     list: &mut [i32],
6     first_len: usize,
7     second_len: usize)
8 {
9     if 0 == first_len || 0 == second_len { return; }
10
11     if first_len > second_len {
12         // Merge B and C, and merge from the end
13         // of the list using temp
14         merge_hi(list, first_len, second_len);
15     } else {

```

```

16         // Merge B and A, and merge from the beginning
17         // of the list using temp
18         merge_lo(list, first_len);
19     }
20 }
21
22 // merge A and B to a single run
23 fn merge_lo(list: &mut [i32], first_len: usize) {
24     unsafe {
25         let mut state = MergeLo::new(list, first_len);
26         state.merge();
27     }
28 }
29
30 impl<'a> MergeLo<'a> {
31     unsafe fn new(list: &'a mut [i32], first_len: usize) ->
32         Self {
33         let mut ret_val = MergeLo {
34             list_len: list.len(),
35             first_pos: 0,
36             first_len: first_len,
37             second_pos: first_len, // run1 close to run2,
38                                   // the start index of run2
39                                   // is exactly equals the
40                                   // length of run1
41             dest_pos: 0,           // write the sorted data
42                                   // back to run1
43             list: list,
44             temp: Vec::with_capacity(first_len),
45         };
46
47         // copy run1 to temp
48         ret_val.temp.set_len(first_len);
49         for i in 0..first_len {
50             ret_val.temp[i] = ret_val.list[i];
51         }
52
53         ret_val
54     }
55
56     // merge sort
57     fn merge(&mut self) {
58         while self.second_pos > self.dest_pos
59             && self.second_pos < self.list_len {
60             debug_assert!((self.second_pos - self.first_len) +
61                 self.first_pos == self.dest_pos);
62
63             if self.temp[self.first_pos]
64                 > self.list[self.second_pos] {
65                 self.list[self.dest_pos]
66                     = self.list[self.second_pos];
67                 self.second_pos += 1;

```

```

67         } else {
68             self.list[self.dest_pos]
69             = self.temp[self.first_pos];
70             self.first_pos += 1;
71         }
72         self.dest_pos += 1;
73     }
74 }
75 }
76
77 // clear temp stack
78 impl<'a> Drop for MergeLo<'a> {
79     fn drop(&mut self) {
80         unsafe {
81             // put left date of temp into list(high digit)
82             if self.first_pos < self.first_len {
83                 for i in 0..(self.first_len - self.first_pos) {
84                     self.list[self.dest_pos + i]
85                     = self.temp[self.first_pos + i];
86                 }
87             }
88
89             // set the lenght of temp to 0
90             self.temp.set_len(0);
91         }
92     }
93 }
94
95 // merge B and C as a new run block
96 fn merge_hi(
97     list: &mut [i32],
98     first_len: usize,
99     second_len: usize)
100 {
101     unsafe {
102         let mut state = MergeHi::new(list,
103                                     first_len,
104                                     second_len);
105         state.merge();
106     }
107 }
108
109 impl<'a> MergeHi<'a> {
110     unsafe fn new(
111         list: &'a mut [i32],
112         first_len: usize,
113         second_len: usize) -> Self
114     {
115         let mut ret_val = MergeHi {
116             first_pos: first_len as isize - 1,
117             second_pos: second_len as isize - 1,
118             dest_pos: list.len() as isize - 1, // from the tail

```

```

119         list: list,
120         temp: Vec::with_capacity(second_len),
121     };
122
123     // copy run2 to temp
124     ret_val.temp.set_len(second_len);
125     for i in 0..second_len {
126         ret_val.temp[i] = ret_val.list[i + first_len];
127     }
128
129     ret_val
130 }
131
132 // merge sort
133 fn merge(&mut self) {
134     while self.first_pos < self.dest_pos
135         && self.first_pos >= 0 {
136         debug_assert!(self.first_pos + self.second_pos + 1
137             == self.dest_pos);
138         if self.temp[self.second_pos as usize]
139             >= self.list[self.first_pos as usize] {
140             self.list[self.dest_pos as usize]
141                 = self.temp[self.second_pos as usize];
142             self.second_pos -= 1;
143         } else {
144             self.list[self.dest_pos as usize]
145                 = self.list[self.first_pos as usize];
146             self.first_pos -= 1;
147         }
148         self.dest_pos -= 1;
149     }
150 }
151
152 }
153
154 // clear temp stack
155 impl<'a> Drop for MergeHi<'a> {
156     fn drop(&mut self) {
157         unsafe {
158             // put left date of temp into list(low digit)
159             if self.second_pos >= 0 {
160                 let size = self.second_pos + 1;
161                 let src = 0;
162                 let dest = self.dest_pos - size;
163                 for i in 0..size {
164                     self.list[(dest + i) as usize]
165                         = self.temp[(src + i) as usize];
166                 }
167             }
168
169             // set the length of temp stack to 0
170             self.temp.set_len(0);

```



```

171         }
172     }
173 }

```

Here is the main function of TimSort.

```

1 // timSort entry point
2
3 fn tim_sort(list: &mut [i32]) {
4     if list.len() < MIN_MERGE {
5         binary_insertion_sort(list);
6     } else {
7         let mut sort_state = SortState::new(list);
8         sort_state.sort();
9     }
10 }

```

Here is an example of TimSort.

```

1 fn main() {
2     let mut nums: Vec<i32> = vec![
3         2, 4, 7, 8, 23, 19, 16, 14, 13, 12, 10, 20,
4         18, 17, 15, 11, 9, -1, 5, 6, 1, 3, 21, 40,
5         22, 39, 38, 37, 36, 35, 34, 33, 24, 30, 31, 32,
6         25, 26, 27, 28, 29, 41, 42, 43, 44, 45, 46, 47,
7         48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59,
8         60, 80, 79, 78, 77, 76, 75, 74, 73, 72, 71, 70,
9         61, 62, 63, 64, 65, 66, 67, 68, 69, 95, 94, 93,
10        92, 91, 90, 85, 82, 83, 84, 81, 86, 87, 88, 89,
11    ];
12    tim_sort(&mut nums);
13    println!("sorted nums: {:?}", nums);
14 }

```

```

sorted nums: [-1, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11,
              12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23,
              24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35,
              36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47,
              48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59,
              60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71,
              72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83,
              84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95]

```

The implemented TimSort algorithm only works for `i32` type numbers, but it can be extended to support sorting algorithms for various types of numbers through the use of generics. Additionally, during merging, some data may already be sorted, but the implemented TimSort algorithm still goes through each comparison one by one. However, merging can be accelerated through a strategy called galloping. The TimSort implementation in this book's source code in `tim_sort_without_gallop.rs` is the non-accelerated version, but an accelerated version of Timsort has also been implemented in `tim_sort.rs`. Interested readers can refer to and compare the differences between the two algorithms.

Some readers may wonder how TimSort determines if the data to be sorted is block-sorted originally. In reality, the algorithm cannot determine this; it is a property of the partially ordered data that the author discovered. In physics, there is a concept of entropy^[13] that refers to the degree of disorder in a physical

system. The more disorderly a system is, the greater the entropy, and conversely, the more ordered it is, the smaller the entropy. In most cases, things have a certain degree of order. For example, humans are ordered animals in reverse entropy, and only become disordered with death. Another example is the principle of locality of reference^[14], where when accessing data on a hard drive, the CPU will read the data around it into memory because it is likely to be accessed next. These two phenomena are natural laws that are consistent with statistical principles, and Tim wrote TimSort based on these laws. This illustrates the importance of data structures and how different understandings can lead to very different algorithms.

7.14 Summary

In this chapter, we learned about ten different types of sorting algorithms. Bubble sort, selection sort, and insertion sort are $O(n^2)$ algorithms, while most other sorting algorithms have a complexity of $O(n \log_2(n))$. Selection sort is an improvement on bubble sort, shell sort improves on insertion sort, heap sort improves on selection sort, and quicksort and merge sort both use the divide-and-conquer approach.

All of these sorts are based on comparison, but there are also non-comparison sorts that rely solely on numerical patterns for sorting, such as bucket sort, counting sort, and radix sort. These sorts have a complexity of approximately $O(n)$ and are suitable for sorting small amounts of data. Counting sort is a special case of bucket sort, and radix sort is a multi-round bucket sort that can degrade into counting sort.

In addition to the basic sorting algorithms, we also learned about improved versions of some algorithms, particularly the TimSort algorithm, which is an efficient and stable hybrid sorting algorithm. Its improved version is already the default sorting algorithm for many languages and platforms.

The table below summarizes the various sorting algorithms, which readers can compare and understand on their own to deepen their understanding.

Table 7.1: Time and space complexity of various sorting algorithms

No	Algorithm	Time-worst	Time-best	Time-average	Space	Stability
1	Bubble sort	$O(n^2)$	$O(n)$	$O(n^2)$	$O(1)$	stable
2	Quick sort	$O(n^2)$	$O(n \log(n))$	$O(n \log(n))$	$O(n \log(n))$	unstable
3	Selection sort	$O(n^2)$	$O(n^2)$	$O(n^2)$	$O(1)$	unstable
4	Heap sort	$O(n \log(n))$	$O(n \log(n))$	$O(n \log(n))$	$O(1)$	unstable
5	Insertion sort	$O(n^2)$	$O(n)$	$O(n^2)$	$O(1)$	stable
6	Shell sort	$O(n^2)$	$O(n)$	$O(n^{1.3})$	$O(1)$	unstable
7	Merge sort	$O(n \log(n))$	$O(n \log(n))$	$O(n \log(n))$	$O(n)$	stable
8	Counting sort	$O(n + k)$	$O(n + k)$	$O(n + k)$	$O(n + k)$	stable
9	Bucket sort	$O(n^2)$	$O(n)$	$O(n + k)$	$O(n + k)$	stable
10	Radix sort	$O(nk)$	$O(nk)$	$O(nk)$	$O(n + k)$	stable
11	Tim sort	$O(n \log(n))$	$O(n)$	$O(n \log(n))$	$O(n)$	stable

Chapter 8

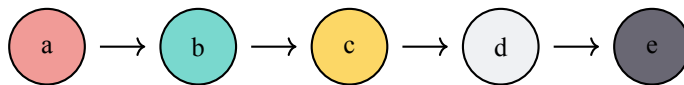
Trees

8.1 Objectives

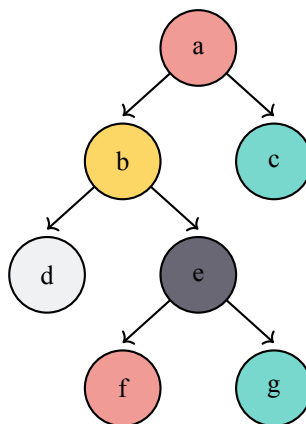
- Understanding trees and their usage.
- Implementing a priority queue using a binary heap.
- Understanding binary search trees and balanced binary trees.
- Implementing binary search trees and balanced binary trees.

8.2 What is Tree?

In previous chapters, we covered linear data structures such as stacks, queues, and linked lists, where each data item is connected to the next one.

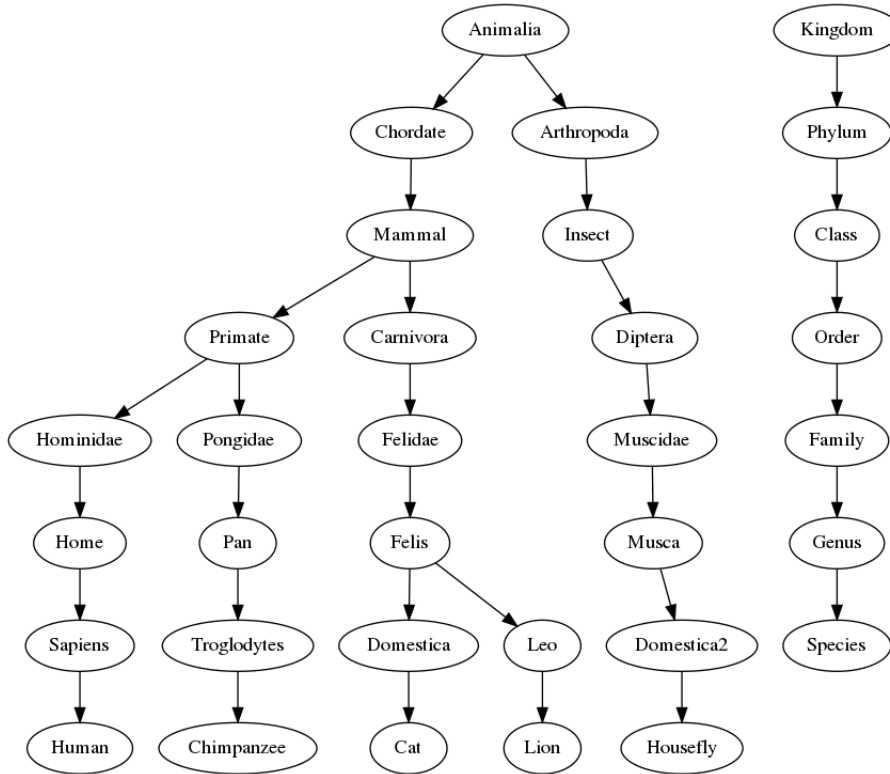


A tree is a new data structure that expands upon this linear structure by connecting multiple data items.



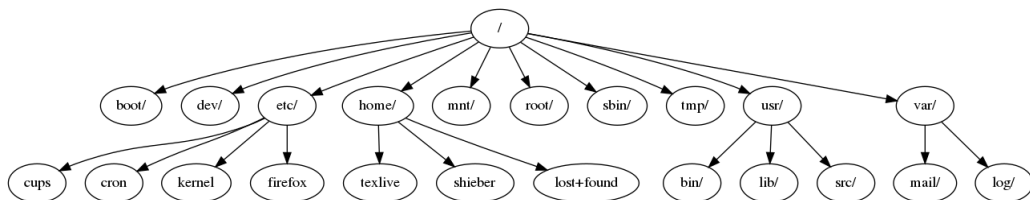
Like a natural tree, this data structure has a root, branches, and leaves, which are interconnected. Trees are used in various fields of computer science, such as operating systems, graphics, databases, and computer networks. Throughout the rest of the text, we will refer to this data structure as a "tree."

Before delving into trees, it's essential to understand some common tree examples, such as a biological classification tree. From this graph, we can see the specific location of entry 'People' (lower left), which is very helpful for studying relationships and properties.



This type of tree is hierarchical, with well-defined structures comprising seven levels: kingdom, phylum, class, order, family, genus, and species. The highest level represents the most abstract concept, while the lowest level represents the most specific. By starting from the root of the tree and following the arrows to the bottom, we can obtain a complete path that indicates the full name of a species. Organisms can find their place on this tree of life, displaying their relative relationships. For instance, the complete name of the "human" species is "animal kingdom-chordate phylum-mammal class-primate order-hominidae family-homo genus-homo sapiens species."

Each node's set of child nodes is independent of another node's set of child nodes, which makes the relationships between nodes clear. This property also means that modifying one node's child nodes will not affect other nodes. It is especially useful when trees are used as data storage containers, as it allows tools to modify data on specific nodes while keeping other data unchanged.



Lastly, each leaf node is unique, and there is only one unique path from the root to each leaf node. This property makes data storage efficient since the unique path can be used as a storage path. The file system in our computers is based on an improved tree structure. The file system tree and biological classification tree have many similarities, and the path from the root directory to any subdirectory uniquely identifies

that subdirectory and all files within it. If you're familiar with Unix-like operating systems, you should recognize paths like `/root`, `/home/user`, and `/etc` as nodes on a tree, where `/` is the root node.

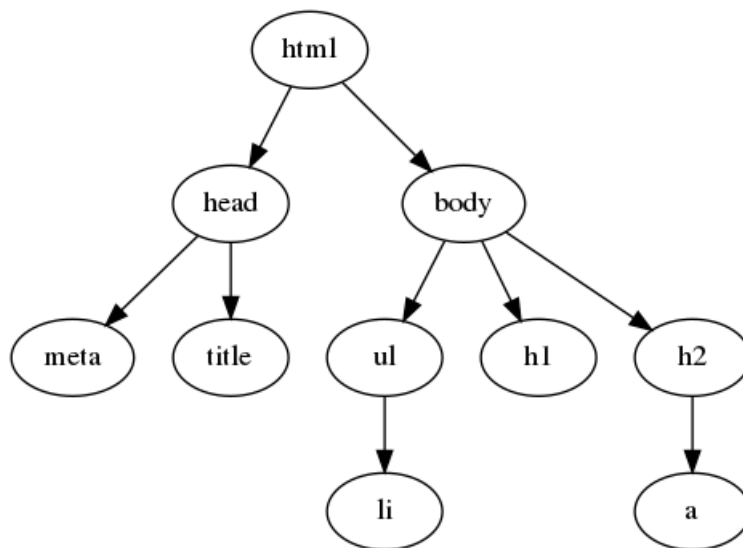
Trees can also be used to represent web page files, which are collections of resources that have a hierarchical structure. For example, the web page data of Google.cn's search interface shows that the `<` tag elements are also arranged hierarchically.

```

1 <html lang="zh"><head><meta charset="utf-8">
2   <head>
3     <meta charset="utf-8">
4     <title>Google</title>
5     <style>
6       html { background: #fff; margin: 0 1em; }
7       body { font: .8125em/1.5 arial, sans-serif; }
8     </style>
9   </head>
10  <body>
11  <div>
12    <h1><a href="http://www.google.com.hk/webhp?hl=zh-CN">
13      <strong id="target">google.com.hk</strong></a></h1>
14    <p>Please save our website</p>
15  </div>
16  <ul>
17    <li><a href="http://translate.google.cn/">translate</a></li>
18  </ul>
19 </body>
20 </html>

```

In web page files, the HTML root encapsulates all the other elements, as illustrated in the diagram below. This hierarchical structure allows web pages to be displayed in a consistent manner across different devices and browsers. By using trees to represent web page files, we can efficiently store and retrieve data and modify individual elements without affecting the entire page's layout.



8.2.1 Vocabularies and Definitions of Tree

After discussing some examples of trees, we will now define the various properties of a tree.

- **Node:** A node is the fundamental unit of a tree, consisting of a "key" name and an optional "payload" information. Payload information is not always necessary for tree algorithms, but it is typically important in applications that require additional information on nodes, such as file names, file content, or timestamps.

- **Root:** The root is the top-level node in a tree, with no incoming edges. All nodes in a tree can be reached by following edges from the root. This is analogous to the concepts of the / or C drive in an operating system.

- **Edge:** An edge connects two nodes and represents the relationship between them. Every node in a tree, except the root, has exactly one incoming edge and potentially several outgoing edges. An edge defines a path that can be used to locate a specific node.

- **Path:** A path is a sequence of edges that connect nodes. It is not a structure in itself, but rather an associative relationship that emerges from the logical arrangement of edges. For instance, /home/user/-files/sort.rs is a path that identifies the location of the file sort.rs.

- **Child Node:** A child node is a node that is one level below another node, originating from the same parent node. For example, sort.rs is a child node of files/. Child nodes are not unique, and there may be zero, one, or many child nodes, similar to the parent-child relationship in human society. Hence, child nodes are often named after kinship terms.

- **Parent Node:** A parent node is a node from which all lower-level nodes originate. For instance, files/ is the parent node of sort.rs. The parent node is unique, as a child can only have one parent.

- **Subtree:** A subtree is a set of nodes and edges that consists of a parent node and all its descendants. Since a tree is a recursive structure, taking any node from the tree creates a new subtree.

- **Leaf Node:** A leaf node is a node without child nodes, at the bottom level of the tree.

- **Internal Node:** An internal node has child nodes and a parent node.

- **Depth:** The depth of a node is the number of edges traversed from the root to that node. The depth of the root node is zero. In /home/user/files, the depth of files is 2. A depth of zero does not mean that there is no depth, but rather the depth is zero, indicating the first level.

- **Height:** The height of a tree is the maximum depth of any node in the tree.

To define a tree, we start with the foundational knowledge presented above:

- A tree has a root node, which is the topmost node in the hierarchy.

- Each node, except for the root node, is connected to a parent node and may have zero or more child nodes, forming a branching structure.

- The path from the root node to any other node in the tree is unique, meaning there is only one path to reach that node.

The following image shows an example of a tree structure with left and right child nodes labeled as "lc" and "rc," respectively. As trees are recursive structures, the structure of a subtree is the same as that of its parent node.

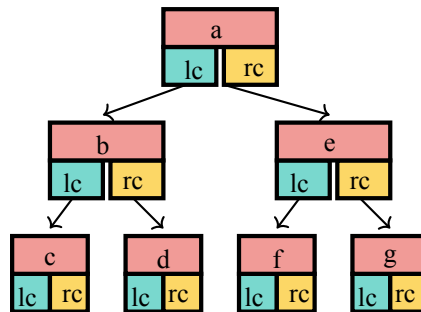


Figure 8.1: tree

8.2.2 Tree Representation

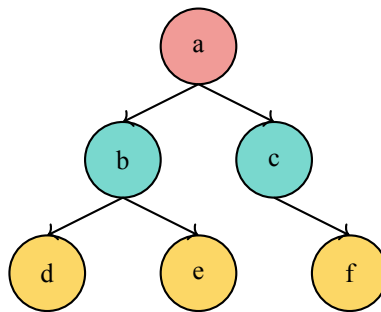
Trees are a powerful non-linear data structure, but representing them in a computer requires linear data structures. One common approach to representing a tree is using an array. In this approach, the elements of the tree are stored in an array, with the root at the first index and the child nodes following in a logical order.

```
tree = [ 'a',
        ['b',
         ['c', [], []],
         ['d', [], []],
        ],
        ['e',
         ['f', [], []],
         []
        ],
      ]
```

Array access methods can be used to access the tree elements, making it easy to obtain subtrees and elements. However, this approach becomes cumbersome and impractical for larger trees with many levels, as accessing deeper nodes requires complex nested access methods.

```
println!("root {:?}", tree[0]);
println!("left subtree {:?}", tree[1]);
println!("right subtree {:?}", tree[2]);
```

An alternative approach to representing trees is to use nodes. In this approach, each node in the tree is represented by an object or a struct, which stores the node's key and payload information, as well as links to its child nodes. The links between nodes form the edges of the tree. This approach is similar to the linked list data structure, where each node has a link to the next node. The node-based representation of a tree is more flexible than the array-based approach and can handle larger trees with ease. The figure below shows an example of a binary tree represented using nodes.



This structure is straightforward and avoids the complexity of nested arrays for accessing elements. The key now is how to define tree nodes. One feasible method is to use the struct to define a node.

```
1 use std::cmp::{max, Ordering::*};
2 use std::fmt::{Debug, Display};
3
4 // Binary tree child node link
5 type Link<T> = Option<Box<BinaryTree<T>>>;
6
7 // Binary tree definition
```

```

8 #[derive(Debug, Clone, PartialEq)]
9 struct BinaryTree<T> {
10     key: T,           // data storage
11     left: Link<T>,    // left child node address storage
12     right: Link<T>,   // right child node address storage
13 }

```

The struct has a key that stores data and left and right pointers that store the addresses of the left and right child nodes.

```

1 // binary_tree.rs
2
3 impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4     fn new(key: T) -> Self {
5         Self { key: key, left: None, right: None }
6     }
7
8     // New child node is added as the
9     // left child node of the root node
10    fn insert_left_tree(&mut self, key: T) {
11        if self.left.is_none() {
12            let node = BinaryTree::new(key);
13            self.left = Some(Box::new(node));
14        } else {
15            let mut node = BinaryTree::new(key);
16            node.left = self.left.take();
17            self.left = Some(Box::new(node));
18        }
19    }
20
21    // New child node is added as the
22    // right child node of the root node
23    fn insert_right_tree(&mut self, key: T) {
24        if self.right.is_none() {
25            let node = BinaryTree::new(key);
26            self.right = Some(Box::new(node));
27        } else {
28            let mut node = BinaryTree::new(key);
29            node.right = self.right.take();
30            self.right = Some(Box::new(node));
31        }
32    }
33 }

```

When inserting a new child node into a tree, there are two possible cases to consider. The first case is when the node has no child nodes, in which case the new node can be directly inserted as its child. The second case is when the node already has child nodes, in which case the new node's child nodes must first be attached to the appropriate positions in the tree before the new node itself can be added as a child of the original node.

The code defining the tree nodes may be confusing since it uses the term "BinaryTree" instead of "Node." However, when multiple nodes are linked together to form a tree, the entire subtree can be considered a single node, which simplifies the insertion process. It is important to note, however, that the internal structure of the node is still relevant, so it is written as "BinaryTree" rather than "Node."

Nonetheless, either term can be used to implement the tree.

Below is the code for calculating the number of different types of nodes and the depth of the tree.

```

1  // binary_tree.rs
2
3  impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4      // calculate total node number
5      fn size(&self) -> usize {
6          self.calc_size(0)
7      }
8
9      fn calc_size(&self, mut size: usize) -> usize {
10         size += 1;
11
12         if !self.left.is_none() {
13             size =self.left.as_ref().unwrap().calc_size(size);
14         }
15         if !self.right.is_none() {
16             size =self.right.as_ref().unwrap().calc_size(size);
17         }
18
19         size
20     }
21
22     // calculate the total leaf nodes number
23     fn leaf_size(&self) -> usize {
24         // both left and right nodes are none
25         // which means current node is a leaf node
26         if self.left.is_none() && self.right.is_none() {
27             return 1;
28         }
29
30         // calculate the total leaf nodes of
31         // both left and right subtrees
32         let left_leaf = match &self.left {
33             Some(left) => left.leaf_size(),
34             None => 0,
35         };
36         let right_leaf = match &self.right {
37             Some(right) => right.leaf_size(),
38             None => 0,
39         };
40
41         // leaf nodes = leaf nodes(left) + leaf nodes(right)
42         left_leaf + right_leaf
43     }
44
45     // calculate non-leaf nodes,
46     // it is actually very easy to calculate
47     fn none_leaf_size(&self) -> usize {
48         self.size() - self.leaf_size()
49     }

```

```

50
51 // calculate the depth of a tree
52 fn depth(&self) -> usize {
53     let mut left_depth = 1;
54     if let Some(left) = &self.left {
55         left_depth += left.depth();
56     }
57
58     let mut right_depth = 1;
59     if let Some(right) = &self.right {
60         right_depth += right.depth();
61     }
62
63     // return the max depth
64     max(left_depth, right_depth)
65 }
66 }

```

To manipulate the data of a binary tree node, you need to implement methods for accessing the left and right child nodes, setting and getting the root node value, and modifying node values. Additionally, methods for determining the existence of a node value and finding the maximum and minimum node values can also be helpful.

```

1 // binary_tree.rs
2
3 impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4     // get left subtree
5     fn get_left(&self) -> Link<T> {
6         self.left.clone()
7     }
8
9     fn get_right(&self) -> Link<T> {
10        self.right.clone()
11    }
12
13    // get and set key
14    fn get_key(&self) -> T {
15        self.key.clone()
16    }
17
18    fn set_key(&mut self, key: T) {
19        self.key = key;
20    }
21
22    // find min/max key in the tree
23    fn min(&self) -> Option<&T> {
24        match self.left {
25            None => Some(&self.key),
26            Some(ref node) => node.min(),
27        }
28    }
29
30    fn max(&self) -> Option<&T> {

```

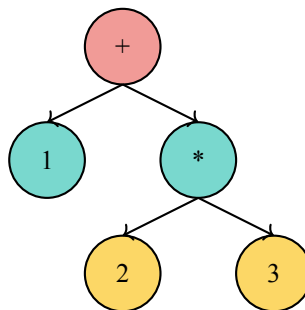
```

31         match self.right {
32             None => Some(&self.key),
33             Some(ref node) => node.max(),
34         }
35     }
36
37     // determine whether a key is in the tree
38     fn contains(&self, key: &T) -> bool {
39         match &self.key.cmp(key) {
40             Equal => true,
41             Greater => {
42                 match &self.left {
43                     Some(left) => left.contains(key),
44                     None => false,
45                 }
46             },
47             Less => {
48                 match &self.right {
49                     Some(right) => right.contains(key),
50                     None => false,
51                 }
52             },
53         }
54     }
55 }

```

8.2.3 Parse Tree

The following is an explanation of how a binary tree can be used to store data, using the example of storing the expression $(1 + (2 * 3))$. The expression is fully parenthesized, which indicates the order of operations. The tree structure that corresponds to this expression is shown in the accompanying diagram.



To store data in the tree, we follow certain rules based on the tree's structure. These rules are as follows:

- If the current symbol is `(`, a new node is added as the left child node, and we descend to that node.
- If the current symbol is one of `"+"`, `"-"`, `"*"`, or `"/"`, we set the root value to that symbol, add a new right child node, and descend to the right child node.
- If the current symbol is a number, we set the root value to that number and return to the parent node.
- If the current symbol is `)`, we return to the parent node of the current node.

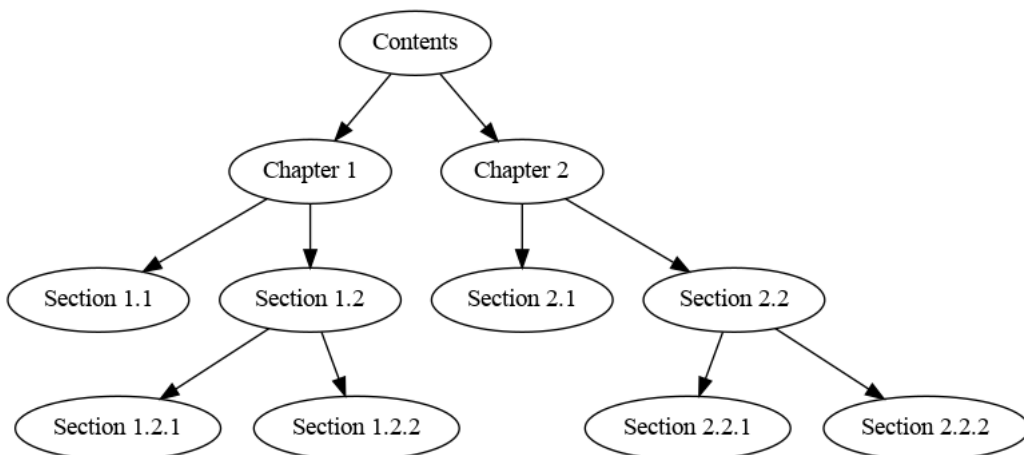
To convert the mathematical expression $(1 + (2 * 3))$ into the tree shown in the figure, we can use the data storage rules defined for the tree. The specific steps are as follows:

- (1) Create the root node.
- (2) Read the symbol $($, create a new left child node, and descend to the node.
- (3) Read the symbol 1, set the node value to 1, and return to the parent node.
- (4) Read the symbol $+$, set the node value to $+$, create a new right child node, and descend to the node.
- (5) Read the symbol $($, create a new left child node, and descend to the node.
- (6) Read the symbol 2, set the node value to 2, and return to the parent node.
- (7) Read the symbol $*$, set the node value to $*$, create a new right child node, and descend to the node.
- (8) Read the symbol 3, set the node value to 3, and return to the parent node.
- (9) Read the symbol $)$, return to the parent node.

Using a tree allows for the storage of the arithmetic expression while maintaining the structural information of the data. In fact, programming languages use trees to store all code during compilation and generate abstract syntax trees. By analyzing the functionality of each part of the syntax tree, intermediate code is generated, optimized, and then final code is generated. If you understand the principles of compilation, this should be familiar to you.

8.2.4 Tree Traversals

The purpose of data storage is to efficiently perform operations such as adding, deleting, searching, and modifying data. In a tree structure, locating a specific node is essential to perform these operations. However, unlike linear data structures that can be traversed using indexes, trees are nonlinear structures, so the search method is different. To access nodes in a tree, there are three commonly used methods that differ in the order in which nodes are visited. These methods are pre-order traversal, in-order traversal, and post-order traversal, and they are analogous to the traversal methods of linear data structures. To illustrate these methods, imagine this book as a tree with the table of contents as the root, each chapter as its child nodes, and each subsection as the child nodes of their respective chapters.



Pre-order traversal is the first method, and it starts from the root node, then moves to the left subtree, and finally the right subtree. To traverse a tree algorithmically using pre-order traversal, recursively call the pre-order traversal method on the left subtree from the root node. For example, in the book tree described above, the pre-order traversal algorithm visits the table of contents first, followed by chapter 1, section 1, section 2, and so on. Once chapter 1 is finished, the algorithm returns to the table of contents and recursively calls pre-order traversal on chapter 2, visiting chapter 2, section 1, section 2, and so on.

The pre-order traversal algorithm is simple and can be implemented as an internal method or an external function.

```

1 // binary_tree.rs
2
3 impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4     fn preorder(&self) {
5         println!("key: {:?}", &self.key);
6         match &self.left {
7             Some(node) => node.preorder(),
8             None => (),
9         }
10        match &self.right {
11            Some(node) => node.preorder(),
12            None => (),
13        }
14    }
15 }
16
17 // pre-order: implemented externally [by recursion]
18 fn preorder<T: Clone + Ord + ToString + Debug>(bt: Link<T>) {
19     if !bt.is_none() {
20         println!("key: {:?}", bt.as_ref().unwrap().get_key());
21         preorder(bt.as_ref().unwrap().get_left());
22         preorder(bt.as_ref().unwrap().get_right());
23     }
24 }

```

The post-order traversal starts from the left subtree, followed by the right subtree, and finally the root node.

```

1 // binary_tree.rs
2
3 impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4     fn postorder(&self) {
5         match &self.left {
6             Some(node) => node.postorder(),
7             None => (),
8         }
9         match &self.right {
10            Some(node) => node.postorder(),
11            None => (),
12        }
13        println!("key: {:?}", &self.key);
14    }
15 }
16
17 // post-order: implemented externally [by recursion]
18 fn postorder<T: Clone + Ord + ToString + Debug>(bt: Link<T>) {
19     if !bt.is_none() {
20         postorder(bt.as_ref().unwrap().get_left());
21         postorder(bt.as_ref().unwrap().get_right());
22         println!("key: {:?}", bt.as_ref().unwrap().get_key());

```

```

23     }
24 }

```

The in-order traversal starts from the left subtree, followed by the root node, and finally the right subtree.

```

1 // binary_tree.rs
2
3 impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4     fn inorder(&self) {
5         if self.left.is_some() {
6             self.left.as_ref().unwrap().inorder();
7         }
8         println!("key: {:?}", &self.key);
9         if self.right.is_some() {
10            self.right.as_ref().unwrap().inorder();
11        }
12    }
13 }
14
15 // in-order: implemented externally [by recursion]
16 fn inorder<T: Clone + Ord + ToString + Debug>(bt: Link<T>) {
17     if !bt.is_none() {
18         inorder(bt.as_ref().unwrap().get_left());
19         println!("key: {:?}", bt.as_ref().unwrap().get_key());
20         inorder(bt.as_ref().unwrap().get_right());
21     }
22 }

```

To evaluate an arithmetic expression like $(1 + (2 * 3))$ that is stored in a tree, we need to retrieve the operator and operands in the correct order. While pre-order traversal starts as much as possible from the root node, it is not suitable for computing expressions as we need to start from the leaf nodes. Post-order traversal, on the other hand, is the correct method to retrieve data as it starts from the left subtree, followed by the right subtree, and finally the root node. By first retrieving the values of the left and right child nodes, then retrieving the operator of the root node, we can perform one operation and save the result in the position of the operator, and then continue post-order traversal by visiting the right child node until we calculate the final value.

Additionally, in-order traversal can be used on the tree that saves the arithmetic expression $(1 + (2 * 3))$ to retrieve the original expression $1 + 2 * 3$. However, since the tree does not save parentheses, the recovered expression is only in the correct order, but the priority may not be correct. To include parentheses in the output, we can modify the in-order traversal.

```

1 // binary_tree.rs
2
3 impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4     // form a expression: [internal implementation]
5     fn iexp(&self) -> String {
6         let mut exp = "".to_string();
7
8         exp += "(";
9         let exp_left = match &self.left {
10             Some(left) => left.iexp(),
11             None => "".to_string(),

```

```

12         };
13         exp += &exp_left;
14
15         exp += &self.get_key().to_string();
16
17         let exp_right = match &self.right {
18             Some(right) => right.iexp(),
19             None => "".to_string(),
20         };
21         exp += &exp_right;
22         exp += ")";
23
24         exp
25     }
26 }
27
28 // form a expression: [external implementation]
29 fn oexp<T>(bt: Link<T>) -> String
30 where T: Clone + Ord + ToString + Debug + Display
31 {
32     let mut exp = "".to_string();
33     if !bt.is_none() {
34         exp = "(" + &bt.as_ref().unwrap().get_left().iexp();
35         exp += &bt.as_ref().unwrap().get_key().to_string();
36         exp += &(oexp(bt.as_ref().unwrap().get_right()) + ")";
37     }
38 }
39
40     exp
41 }

```

In addition to the three traversal methods discussed earlier, there is another method called level-order traversal that visits the nodes layer by layer. As we have previously implemented the queue data structure required for level-order traversal, we can directly use it to implement this traversal method.

```

1 // binary_tree.rs
2
3 impl<T: Clone + Ord + ToString + Debug> BinaryTree<T> {
4     fn levelorder(&self) {
5         let size = self.size();
6         let mut q = Queue::new(size);
7
8         // enqueue the root node
9         let _r = q.enqueue(Box::new(self.clone()));
10        while !q.is_empty() {
11            // dequeue the first node, and output its value
12            let front = q.dequeue().unwrap();
13            println!("key: {:?}", front.get_key());
14
15            // enqueue the left/right child node
16            match front.get_left() {
17                Some(left) => {
18                    let _r = q.enqueue(left);

```

```

19         },
20         None => {},
21     }
22
23     match front.get_right() {
24         Some(right) => {
25             let _r = q.enqueue(right);
26         },
27         None => {},
28     }
29 }
30 }
31 }
32
33 // level-order: implemented externally [by recursion]
34 fn levelorder<T: Clone + Ord + ToString + Debug>(bt: Link<T>) {
35     if bt.is_none() { return; }
36
37     let size = bt.as_ref().unwrap().size();
38     let mut q = Queue::new(size);
39
40     let _r = q.enqueue(bt.as_ref().unwrap().clone());
41     while !q.is_empty() {
42         // dequeue the first node, and print its value
43         let front = q.dequeue().unwrap();
44         println!("key: {:?}", front.get_key());
45
46         match front.get_left() {
47             Some(left) => {
48                 let _r = q.enqueue(left);
49             },
50             None => {},
51         }
52
53         match front.get_right() {
54             Some(right) => {
55                 let _r = q.enqueue(right);
56             },
57             None => {},
58         }
59     }
60 }

```

Here are some use examples of binary trees.

```

1 // binary_tree.rs
2
3 fn main() {
4     basic();
5     order();
6
7     fn basic() {
8         let mut bt = BinaryTree::new(10usize);

```



```

9      let root = bt.get_key();
10     println!("root key: {:?}", root);
11
12     bt.set_key(11usize);
13     let root = bt.get_key();
14     println!("root key: {:?}", root);
15
16     bt.insert_left_tree(2usize);
17     bt.insert_right_tree(18usize);
18
19     println!("left child: {:?}", bt.get_left());
20     println!("right child: {:?}", bt.get_right());
21
22     println!("min key: {:?}", bt.min().unwrap());
23     println!("max key: {:?}", bt.max().unwrap());
24
25     println!("tree nodes: {}", bt.size());
26     println!("tree leaves: {}", bt.leaf_size());
27     println!("tree internals: {}", bt.none_leaf_size());
28     println!("tree depth: {}", bt.depth());
29     println!("tree contains '2': {}", bt.contains(&2));
30 }
31
32
33 fn order() {
34     let mut bt = BinaryTree::new(10usize);
35     bt.insert_left_tree(2usize);
36     bt.insert_right_tree(18usize);
37
38     println!("internal pre-in-post-level order");
39     bt.preorder();
40     bt.inorder();
41     bt.postorder();
42     bt.levelorder();
43
44     let nk = Some(Box::new(bt.clone()));
45     println!("outside pre-in-post-level order");
46     preorder(nk.clone());
47     inorder(nk.clone());
48     postorder(nk.clone());
49     levelorder(nk.clone());
50
51     println!("internal exp: {}", bt.iexp);
52     println!("outside exp: {}", oexp(nk));
53 }
54 }

```

```

root key: 10
root key: 11
left child: Some(
  BinaryTree {
    key: 2,

```

```

        left: None,
        right: None,
    },
)
right child: Some(
  BinaryTree {
    key: 18,
    left: None,
    right: None,
  },
)
min key: 2
max key: 18
tree nodes: 3
tree leaves: 2
tree internals: 1
tree depth: 2
tree contains '2': true
internal pre-in-post-level order:
key: 10
key: 2
key: 18
key: 2
key: 10
key: 18
key: 2
key: 18
key: 10
key: 10
key: 2
key: 18
outside pre-in-post-level order:
key: 10
key: 2
key: 18
key: 2
key: 10
key: 18
key: 2
key: 18
key: 10
key: 10
key: 2
key: 18
internal exp: ((2)10(18))
outside exp: ((2)10(18))

```

To simplify the descriptions of the three traversal orders, we can use abbreviations. Pre-order traversal can be abbreviated as "rt-l-r" to indicate that we first visit the root, then the left subtree, and finally the right subtree. Combining the three traversal orders, we have "rt-l-r" for pre-order traversal, "l-rt-r" for in-order traversal, and "l-r-rt" for post-order traversal. It's worth noting that there can also be a "rt-r-l" traversal order, but this is simply the mirror image of pre-order traversal. Since left and right are relative,

"l-rt-r" and "r-rt-l" can be regarded as mirror images of each other. Similarly, "r-l-rt" is the mirror image of post-order traversal, and "r-rt-l" is the mirror image of in-order traversal. The table below summarizes the traversal methods for easy reference:

Table 8.1: Pre-In-Post order and their mirror order method

No	transversal method	order	mirror-order	mirror transversal method
1	pre-order	rt-l-r	rt-r-l	mirror-pre-order
2	in-order	l-rt-r	r-rt-l	mirror-in-order
3	post-order	l-r-rt	r-l-rt	mirror-post-order
4	mirror-pre-order	rt-r-l	rt-l-r	pre-order
5	mirror-in-order	r-rt-l	l-rt-r	in-order
6	mirror-post-order	r-l-rt	l-r-rt	post-order

8.3 Binary Heap

In the previous sections, we learned about the queue, a first-in, first-out linear data structure. Another variation of the queue is the priority queue, which dequeues items based on priority rather than their order of addition. In this type of queue, the highest-priority item is dequeued first, regardless of when it was added. When a new item is added to a priority queue, it will move to the front if its priority is high enough. This movement is a way of sorting based on a certain metric, to ensure that high-priority items are at the front.

The priority queue is useful for tasks that involve priorities, such as scheduling processes in an operating system. In this scenario, a priority queue is used to obtain the priority of each process and sort them accordingly. For example, if you are listening to music and browsing news on your phone, and receive an incoming call, the system will give the call the highest priority and interrupt both the news and music to display the incoming call interface. This is an example of managing processes using a priority queue, and assigning a very high priority to the incoming call.

To implement a priority queue, we need to sort it based on some rule to place high-priority items at the front. However, inserting items into the queue has a complexity of $O(n)$, and sorting the queue has a complexity of at least $O(n\log(n))$. To achieve faster results, a binary heap can be used to sort the priority queue. A binary heap is essentially a complete binary tree and allows queuing and dequeuing in $O(\log(n))$ time, making it ideal for efficient scheduling systems.

Despite being defined as a binary tree, a linear data structure such as an array, slice, or Vec can be used to implement the binary heap. As long as the operations follow the definition of a binary heap, a linear data structure can implement its functionality and be used as a non-linear data structure. Additionally, there are two common forms of binary heaps: the min heap and the max heap, depending on whether the smallest or largest data item is at the top. This means that there are also two forms of priority queues.

8.3.1 The Binary Heap Abstract Data Type

A min-heap can be used as a priority queue, and its abstract data type includes the following methods:

- `new()` creates a new binary heap with no parameters and returns an empty heap.
- `push(k)` adds a new item with parameter value `k` to the heap and returns nothing.
- `pop()` returns and removes the minimum item from the heap and modifies the heap.
- `min()` returns the minimum item in the heap without modifying the heap.
- `size()` returns the number of items in the heap as a numerical value.
- `is_empty()` returns a Boolean value indicating whether the heap is empty or not.
- `build(arr)` constructs a new heap from an array or vector of data, with `arr` as the parameter.

If a binary heap 'h' has already been created as a priority queue, the table below shows the results of the heap operations performed, with the top item of the heap on the right. The priority value of an item is its value, with smaller values being higher priority and therefore appearing on the right.

Table 8.2: Operations on binary heap

operations	heap value	return value	operations	heap value	return value
h.is_empty()	[]	true	h.is_empty()	[8,6,3,2,1]	false
h.push(3)	[3]		h.pop()	[8,6,3,2]	1
h.push(8)	[8,3]		h.min()	[8,6,3,2]	2
h.min()	[8,3]	3	h.pop()	[8,6,3]	2
h.push(6)	[8,6,3]		h.pop()	[8,6]	3
h.size()	[8,6,3]	3	h.build([5,4])	[8,6,5,4]	
h.push(2)	[8,6,3,2]		h.build([1])	[8,6,5,4,1]	
h.push(1)	[8,6,3,2,1]		h.min()	[8,6,5,4,1]	1

8.3.2 Implementing a Binay Heap in Rust

To ensure efficient performance of a binary heap, it is crucial to leverage its logarithmic property. For a binary heap stored in a linear data structure, keeping it balanced is key to achieving logarithmic performance. A balanced binary heap has approximately equal numbers of nodes in its left and right subtrees, and it strives to fill each node's left and right child nodes, with at most one node having a non-full set of children in the worst case.

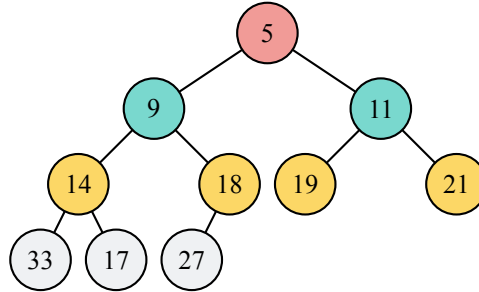


Figure 8.2: Binary Heap

To illustrate, let's consider a heap [0,5,9,11,14,18,19,21,33,17,27] stored using a Vec, and the corresponding tree structure displayed above. As the parent and child nodes are stored in a linear data structure, their relationship is straightforward to compute. If a node is at index p , then its left child node is at $2p$, and its right child node is at $2p + 1$. Here, p starts from index 1, and index 0 is not used for data, so it is set to 0 as a placeholder.

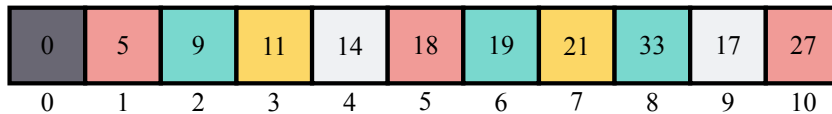


Figure 8.3: Heap in Vec

For instance, the index of 5 is $p = 1$, and its left child node is at index $2 * p = 2$, where the value is 9. Therefore, 9 is the left child node of 5 in the tree structure. Similarly, the parent node of any child node is located at index $p/2$. For example, if $p = 2$ for 9, then its parent node is at index $2/2=1$, and if $p = 3$

for the right child node 11, then its parent node is at index $3/2 = 1$ (the division result is rounded down). Hence, computing the parent node of any child node only requires the expression $p/2$, while child nodes are computed using $2p$ and $2p+1$. We previously defined macros for computing parent and child node indices, and we will continue to use them for calculation purposes.

```

1 // binary_heap.rs
2
3 // calculate parent node index
4 macro_rules! parent {
5     ($child:ident) => {
6         $child >> 1
7     };
8 }
9
10 // calculate left child node index
11 macro_rules! left_child {
12     ($parent:ident) => {
13         $parent << 1
14     };
15 }
16
17 // calculate right child node index
18 macro_rules! right_child {
19     ($parent:ident) => {
20         ($parent << 1) + 1
21     };
22 }

```

To begin with, the binary heap is defined as a data structure that includes a field representing the size of the heap. The size field does not include the first data item 0, which is considered a placeholder. The data saved in the heap is assumed to be `i32`. When initializing the heap, data is present at index 0, but the size is set to 0.

```

1 // binary_heap.rs
2
3 // Binary heap definision
4 // Implement Debug and Clone Trait
5 #[derive(Debug, Clone)]
6 struct BinaryHeap {
7     size: usize, // data count
8     data: Vec<i32>, // data container
9 }
10
11 impl BinaryHeap {
12     fn new() -> Self {
13         BinaryHeap {
14             size: 0,
15             data: vec![0] // first 0 not count in total
16         }
17     }
18
19     fn size(&self) -> usize {
20         self.size
21     }
22 }

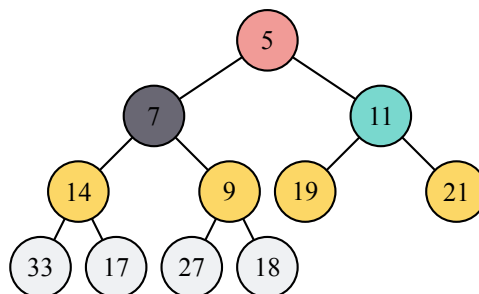
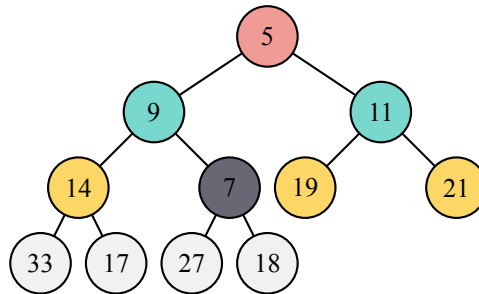
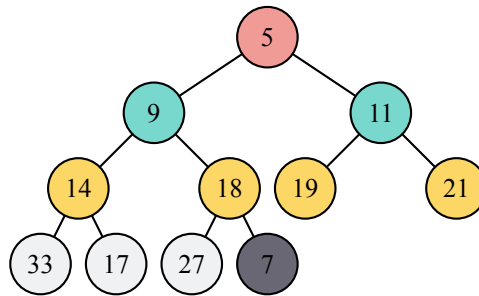
```

```

21     }
22
23     fn is_empty(&self) -> bool {
24         0 == self.size
25     }
26
27     // Get the minimum data in the heap
28     fn min(&self) -> Option<i32> {
29         if 0 == self.size {
30             None
31         } else {
32             // Some(self.data[1].clone());
33             // clone used for generic type
34             Some(self.data[1])
35         }
36     }
37 }

```

When adding data to the heap, adding it to the end of the heap will disrupt the balance, so the data needs to be moved up to maintain balance.



As each data item is added, the size of the heap is increased, and the data is moved up if necessary to maintain balance.

```

1 // binary_heap.rs
2
3 impl BinaryHeap {
4     // Add a data to the end and adjust the heap
5     fn push(&mut self, val: i32) {
6         self.data.push(val);
7         self.size += 1;
8         self.move_up(self.size);
9     }
10
11     // little data move up.
12     // c(child, current), p(parent)
13     fn move_up(&mut self, mut c: usize) {
14         loop {
15             // calculate the parent index of current node
16             let p = parent!(c);
17             if p <= 0 {
18                 break;
19             }
20
21             // If the current node's data is smaller than
22             // the parent node's data, swap them
23             if self.data[c] < self.data[p] {
24                 self.data.swap(c, p);
25             }
26
27             // The parent node becomes the current node
28             c = p;
29         }
30     }
31 }

```

To retrieve the minimum value from the heap, three cases need to be considered: when there is no data in the heap, return `None`; when there is only one data item, pop it directly; when there are multiple data items, swap the top and end data of the heap, adjust the heap, and then return the minimum value at the end. The `move_down` function is used to move elements down to maintain balance, and the `min_child` function is used to find the minimum child node.

```

1 // binary_heap.rs
2
3 impl BinaryHeap {
4     // pop out the top value
5     fn pop(&mut self) -> Option<i32> {
6         if 0 == self.size {
7             // no data, return None
8             None
9         } else if 1 == self.size {
10             self.size -= 1;
11
12             self.data.pop()

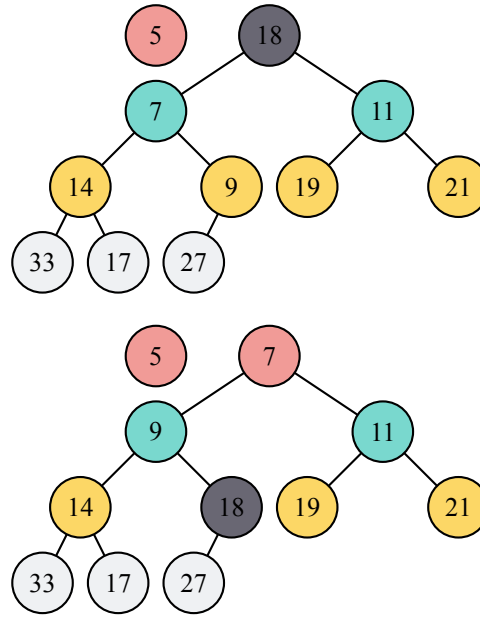
```

```

13         } else {
14             // swap data and then adjust the heap
15             self.data.swap(1, self.size);
16             let val = self.data.pop();
17             self.size -= 1;
18             self.move_down(1);
19
20             val
21         }
22     }
23
24     // bigger data move down
25     fn move_down(&mut self, mut c: usize) {
26         loop {
27             let lc = left_child!(c);
28             if lc > self.size { break; }
29
30             // the index of minimum child node of current node
31             let mc = self.min_child(c);
32             if self.data[c] > self.data[mc] {
33                 self.data.swap(c, mc);
34             }
35
36             // the minimum child node becomes current node
37             c = mc;
38         }
39     }
40
41     // Calculate the index of the minimum child node
42     fn min_child(&self, c: usize) -> usize {
43         let (lc, rc) = (left_child!(c), right_child!(c));
44
45         if rc > self.size {
46             // right child node is out of range,
47             // left child node is the minimum child node
48             lc
49         } else if self.data[lc] < self.data[rc] {
50             // left child node is smaller than right child node
51             lc
52         } else {
53             // right child node is smaller than left child node
54             rc
55         }
56     }
57 }

```

The process of deleting the minimum element from the heap involves taking out the element at the top of the heap and moving the last element to the top. The top element is not the minimum value at this point and does not meet the heap definition, so the heap needs to be rebuilt. To rebuild the heap, the top element needs to be moved down using the `move_down` function. This function uses a macro to calculate whether to swap with the left or right child node based on the node index, eventually swapping the top element with its minimum child node until the heap is restored.



Finally, data can be added to the heap in batches. For instance, a slice of data [0,5,4,3,1,2] can be added to the heap at once to avoid frequent calls to the push function. The original data in the heap can either be kept unchanged while adding the slice data one by one or deleted before adding the slice data. The latter approach results in a new heap that includes only the data in the slice.

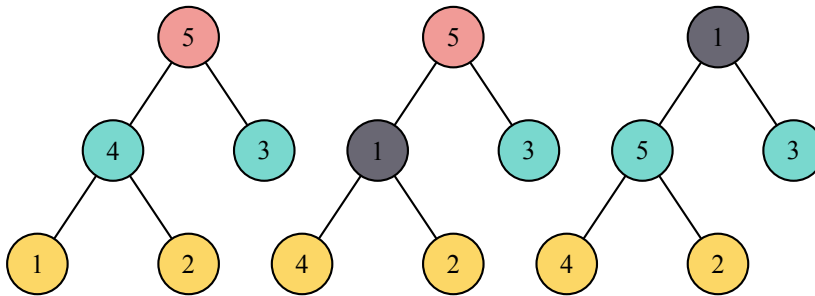


Figure 8.4: delete data and rebuild the heap

To implement the binary min heap, we define the `build_new` and `build_add` functions, as shown below.

```

1 // binary_heap.rs
2
3 impl BinaryHeap {
4     // build a new heap
5     fn build_new(&mut self, arr: &[i32]) {
6         // delete all data
7         for _i in 0..self.size {
8             let _rm = self.data.pop();
9         }
10
11         // add new data
12         for &val in arr {

```

```

13         self.data.push(val);
14     }
15
16     // change the size
17     self.size = arr.len();
18
19     // adjust the heap to make it a min-heap
20     let size = self.size;
21     let mut p = parent!(size);
22     while p > 0 {
23         self.move_down(p);
24         p -= 1;
25     }
26 }
27
28 // add slice data one by one
29 fn build_add(&mut self, arr: &[i32]) {
30     for &val in arr {
31         self.push(val);
32     }
33 }
34 }

```

With these functions, we have completed the construction of a binary min heap. The entire process should be easy to understand, and based on this, one can write the code for a binary max heap as well. Here's an example of using a binary heap.

```

1 // binary_heap.rs
2
3 fn main() {
4     let mut bh = BinaryHeap::new();
5     let nums = [-1,0,2,3,4];
6     bh.push(10); bh.push(9);
7     bh.push(8); bh.push(7); bh.push(6);
8
9     bh.build_add(&nums);
10    println!("empty: {:?}", bh.is_empty());
11    println!("min: {:?}", bh.min());
12    println!("pop min: {:?}", bh.pop());
13
14    bh.build_new(&nums);
15    println!("size: {:?}", bh.len());
16    println!("pop min: {:?}", bh.pop());
17 }

```

The following are the outputs after execution.

```

empty: false
min: Some(-1)
size: 10
pop min: Some(-1)
size: 5
pop min: Some(-1)

```

8.3.3 Analysis of Binary Heap

The binary heap stores data linearly in a Vec, but its sorting is based on a tree structure. As discussed earlier, the height of a tree is $O(n \log_2(n))$, and heap sorting moves from the bottom layer to the top layer of the tree, with the number of movements being equal to the number of layers. Hence, the time complexity of heap sorting is $O(n \log_2(n))$. Since constructing a heap requires processing all n items, the complexity is $O(n)$. Overall, the time complexity of the binary heap is $O(n \log_2(n)) + O(n) = O(n \log_2(n))$.

8.4 Binary Search Tree

However, linear data structures for trees are not suitable for large amounts of data as copying and moving data become very time-consuming. To address this, this section explores implementing trees using nodes. Specifically, this section focuses on binary trees, which have only two child nodes. To understand binary trees better, we study binary search trees for searching, which store data using key-value pairs, similar to HashMap.

8.4.1 The Binary Search Tree Abstract Data Type

The binary search tree provides the following abstract data types:

- `new()` creates a new tree, takes no parameters, and returns an empty tree.
- `insert(k, v)` stores the data (k, v) in the tree, takes the key k and value v , and returns nothing.
- `contains(&k)` searches the tree for the key k , takes a parameter $\&k$, and returns a boolean.
- `get(&k)` returns the value v of key k from the tree without deleting it, takes a parameter $\&k$.
- `max()` returns the maximum key k and its value v in the tree, takes no parameters.
- `min()` returns the minimum key k and its value v in the tree, takes no parameters.
- `len()` returns the number of data items in the tree, takes no parameters, and returns an integer of type `usize`.
- `is_empty()` tests whether the tree is empty, takes no parameters, and returns a boolean.
- `iter()` returns the tree in iterator form, takes no parameters, and does not change the tree.
- `preorder()` performs a pre-order traversal, takes no parameters, and outputs each k - v pairs.
- `inorder()` performs an in-order traversal, takes no parameters, and outputs each k - v pairs.
- `postorder()` performs a post-order traversal, takes no parameters, and outputs each k - v pairs.

The table below displays the binary search tree after different operations, represented by tuples for tree nodes and enclosed in square brackets, assuming `t` is a newly created empty tree.

Table 8.3: Operations on binary search tree

operation	tree value	return value
<code>t.is_empty()</code>	<code>[]</code>	<code>true</code>
<code>t.insert(1, 'a')</code>	<code>[(1, 'a')]</code>	
<code>t.insert(2, 'b')</code>	<code>[(1, 'a'), (2, 'b')]</code>	
<code>t.len()</code>	<code>[(1, 'a'), (2, 'b')]</code>	<code>2</code>
<code>t.get(&4)</code>	<code>[(1, 'a'), (2, 'b')]</code>	<code>None</code>
<code>t.get(&2)</code>	<code>[(1, 'a'), (2, 'b')]</code>	<code>Some('b')</code>
<code>t.min()</code>	<code>[(1, 'a'), (2, 'b')]</code>	<code>(Some(1), Some('a'))</code>
<code>t.max()</code>	<code>[(1, 'a'), (2, 'b')]</code>	<code>(Some(2), Some('b'))</code>
<code>t.contains(2)</code>	<code>[(1, 'a'), (2, 'b')]</code>	<code>true</code>
<code>t.insert(2, 'c')</code>	<code>[(1, 'a'), (2, 'c')]</code>	
<code>t.insert(3, 'd')</code>	<code>[(1, 'a'), (2, 'c'), (3, 'd')]</code>	
<code>t.contains(4)</code>	<code>[(1, 'a'), (2, 'c'), (3, 'd')]</code>	<code>false</code>
<code>t.get(&3)</code>	<code>[(1, 'a'), (2, 'c'), (3, 'd')]</code>	<code>Some('d')</code>

8.4.2 Implementing a Binary Search Tree in Rust

In contrast to heaps, where the left and right child nodes are not compared to their value, in a binary search tree, the key of the left child node is smaller than the key of the parent node, and the key of the right child node is greater than the key of the parent node. This rule of $\text{left} < \text{parent} < \text{right}$ is applied recursively to all subtrees.

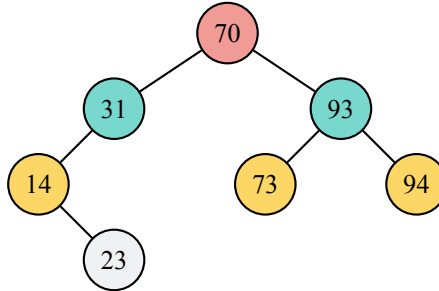


Figure 8.5: Binary search tree

In the example above, 70 is the root node, 31 is smaller and becomes the left node, and 93 is larger and becomes the right node. Then 14 is inserted, which is smaller than 70 and descends to 31, and as it is smaller than 31, it becomes the left node of 31. Similar steps are taken to insert other data, and finally, a binary search tree is formed. The inorder traversal of the tree is [14, 23, 31, 70, 73, 93, 94], which is sorted from small to large, so the binary search tree can also be used to sort data. Using inorder traversal, we obtain the ascending order sorting result, and using the mirrored inorder traversal, i.e., the "right-root-left" order traversal, we obtain the descending order sorting result.

To implement the binary search tree, we define it as a struct `BST` (`BinarySearchTree`), which includes key values and links to left and right child nodes. Following the definition of the abstract data type, the implementation of the binary search tree is shown below.

```

1 // bst.rs
2 use std::cmp::{max, Ordering::*};
3 use std::fmt::Debug;
4
5 // Binary search tree node link
6 type Link<T,U> = Option<Box<BST<T,U>>>;
7
8 // Definition of binary search tree
9 #[derive(Debug,Clone)]
10 struct BST<T,U> {
11     key: Option<T>,
12     val: Option<U>,
13     left: Link<T,U>,
14     right: Link<T,U>,
15 }
16
17 impl<T,U> BST<T,U>
18     where T: Copy + Ord + Debug,
19           U: Copy + Debug
20 {
21     fn new() -> Self {
22         Self {
23             key: None,

```

```

24         val: None,
25         left: None,
26         right: None,
27     }
28 }
29
30 fn is_empty(&self) -> bool {
31     self.key.is_none()
32 }
33
34 fn size(&self) -> usize {
35     self.calc_size(0)
36 }
37
38 // Recursively count the number of nodes
39 fn calc_size(&self, mut size: usize) -> usize {
40     if self.key.is_none() { return size; }
41
42     // Add current node count to total node count 'size'
43     size += 1;
44     // Count left and right child nodes
45     if !self.left.is_none() {
46         size = self.left.as_ref().unwrap().calc_size(size);
47     }
48     if !self.right.is_none() {
49         size = self.right
50             .as_ref().unwrap().calc_size(size);
51     }
52
53     size
54 }
55
56 // Count leaf nodes
57 fn leaf_size(&self) -> usize {
58     // If both left and right are empty,
59     // current node is a leaf node, return 1
60     if self.left.is_none() && self.right.is_none() {
61         return 1;
62     }
63
64     // Count leaf nodes in left and right subtree
65     let left_leaf = match &self.left {
66         Some(left) => left.leaf_size(),
67         None => 0,
68     };
69     let right_leaf = match &self.right {
70         Some(right) => right.leaf_size(),
71         None => 0,
72     };
73
74     // total sum of leaf nodes
75     left_leaf + right_leaf

```

```

76     }
77
78     // Count non-leaf nodes
79     fn none_leaf_size(&self) -> usize {
80         self.size() - self.leaf_size()
81     }
82
83     // Calculate tree depth
84     fn depth(&self) -> usize {
85         let mut left_depth = 1;
86         if let Some(left) = &self.left {
87             left_depth += left.depth();
88         }
89
90         let mut right_depth = 1;
91         if let Some(right) = &self.right {
92             right_depth += right.depth();
93         }
94
95         max(left_depth, right_depth)
96     }
97
98     fn insert(&mut self, key: T, val: U) {
99         // If no data, insert directly
100         if self.key.is_none() {
101             self.key = Some(key);
102             self.val = Some(val);
103         } else {
104             match &self.key {
105                 Some(k) => {
106                     // If key exists, update val
107                     if key == *k {
108                         self.val = Some(val);
109                         return;
110                     }
111
112                     // If no same key found,
113                     // Find the subtree to insert new node
114                     let child = if key < *k {
115                         &mut self.left
116                     } else {
117                         &mut self.right
118                     };
119
120                     // Recursively go down the tree
121                     // until insertion
122                     match child {
123                         Some(ref mut node) => {
124                             node.insert(key, val);
125                         },
126                         None => {
127                             let mut node = BST::new();

```

```

128             node.insert(key, val);
129             *child = Some(Box::new(node));
130         },
131     },
132 },
133     None => (),
134 },
135 },
136 }
137
138 // find the key in tree
139 fn contains(&self, key: &T) -> bool {
140     match &self.key {
141         None => false,
142         Some(k) => {
143             // Compare key value and determine
144             // whether to continue searching recursively
145             match k.cmp(key) {
146                 Equal => true, // find it
147                 Greater => { // search in left subtree
148                     match &self.left {
149                         Some(node) => node.contains(key),
150                         None => false,
151                     }
152                 },
153                 Less => { // search in right subtree
154                     match &self.right {
155                         Some(node) => node.contains(key),
156                         None => false,
157                     }
158                 },
159             }
160         },
161     }
162 }
163
164 // get the min/max node value
165 fn min(&self) -> (Option<&T>, Option<&U>) {
166     // miniumun value is on the left subtree
167     match &self.left {
168         Some(node) => node.min(),
169         None => match &self.key {
170             Some(key) => (Some(&key), self.val.as_ref()),
171             None => (None, None),
172         },
173     }
174 }
175
176 fn max(&self) -> (Option<&T>, Option<&U>) {
177     // maxiumun value is on the right subtree
178     match &self.right {
179         Some(node) => node.max(),

```

```

180         None => match &self.key {
181             Some(key) => (Some(&key), self.val.as_ref()),
182             None => (None, None),
183         },
184     }
185 }
186
187 // get subtree
188 fn get_left(&self) -> Link<T,U> {
189     self.left.clone()
190 }
191
192 fn get_right(&self) -> Link<T,U> {
193     self.right.clone()
194 }
195
196 // get a value reference with parameter key
197 fn get(&self, key: &T) -> Option<&U> {
198     match &self.key {
199         None => None,
200         Some(k) => {
201             match k.cmp(key) {
202                 Equal => self.val.as_ref(),
203                 Greater => {
204                     match &self.left {
205                         None => None,
206                         Some(node) => node.get(key),
207                     }
208                 },
209                 Less => {
210                     match &self.right {
211                         None => None,
212                         Some(node) => node.get(key),
213                     }
214                 },
215             }
216         },
217     }
218 }
219 }

```

Here are the implementation of preorder, inorder, postorder, and level-order traversals.

```

1 // bst.rs
2
3 impl<T,U> BST<T,U>
4     where T: Copy + Ord + Debug,
5           U: Copy + Debug
6 {
7     // Internal implementation
8     fn preorder(&self) {
9         println!("key: {:?}, val: {:?}",self.key, self.val);
10        match &self.left {

```



```

11         Some(node) => node.preorder(),
12         None => (),
13     }
14     match &self.right {
15         Some(node) => node.preorder(),
16         None => (),
17     }
18 }
19
20 fn inorder(&self) {
21     match &self.left {
22         Some(node) => node.inorder(),
23         None => (),
24     }
25     println!("key: {:?}, val: {:?}",self.key, self.val);
26     match &self.right {
27         Some(node) => node.inorder(),
28         None => (),
29     }
30 }
31
32 fn postorder(&self) {
33     match &self.left {
34         Some(node) => node.postorder(),
35         None => (),
36     }
37     match &self.right {
38         Some(node) => node.postorder(),
39         None => (),
40     }
41     println!("key: {:?}, val: {:?}",self.key, self.val);
42 }
43
44 fn levelorder(&self) {
45     let size = self.size();
46     let mut q = Queue::new(size);
47
48     let _r = q.enqueue(Box::new(self.clone()));
49     while !q.is_empty() {
50         let front = q.dequeue().unwrap();
51         println!("key: {:?}, val: {:?}",
52             front.key, front.val);
53
54         match front.get_left() {
55             Some(left) => { let _r = q.enqueue(left); },
56             None => (),
57         }
58         match front.get_right() {
59             Some(right) => { let _r = q.enqueue(right); },
60             None => (),
61         }
62     }

```

```

63     }
64 }
65
66 // External implementation
67 fn preorder<T, U>(bst: Link<T,U>)
68 where T: Copy + Ord + Debug,
69       U: Copy + Debug
70 {
71     if !bst.is_none() {
72         println!("key: {:?}, val: {:?}",
73                 bst.as_ref().unwrap().key.unwrap(),
74                 bst.as_ref().unwrap().val.unwrap());
75         preorder(bst.as_ref().unwrap().get_left());
76         preorder(bst.as_ref().unwrap().get_right());
77     }
78 }
79
80 fn inorder<T, U>(bst: Link<T,U>)
81 where T: Copy + Ord + Debug,
82       U: Copy + Debug
83 {
84     if !bst.is_none() {
85         inorder(bst.as_ref().unwrap().get_left());
86         println!("key: {:?}, val: {:?}",
87                 bst.as_ref().unwrap().key.unwrap(),
88                 bst.as_ref().unwrap().val.unwrap());
89         inorder(bst.as_ref().unwrap().get_right());
90     }
91 }
92
93 fn postorder<T, U>(bst: Link<T,U>)
94 where T: Copy + Ord + Debug,
95       U: Copy + Debug
96 {
97     if !bst.is_none() {
98         postorder(bst.as_ref().unwrap().get_left());
99         postorder(bst.as_ref().unwrap().get_right());
100        println!("key: {:?}, val: {:?}",
101                bst.as_ref().unwrap().key.unwrap(),
102                bst.as_ref().unwrap().val.unwrap());
103    }
104 }
105
106 fn levelorder<T, U>(bst: Link<T,U>)
107 where T: Copy + Ord + Debug,
108       U: Copy + Debug
109 {
110     if bst.is_none() { return; }
111
112     let size = bst.as_ref().unwrap().size();
113     let mut q = Queue::new(size);
114     let _r = q.enqueue(bst.as_ref().unwrap().clone());

```

```

115     while !q.is_empty() {
116         let front = q.dequeue().unwrap();
117         println!("key: {:?}, val: {:?}", front.key, front.val);
118
119         match front.get_left() {
120             Some(left) => { let _r = q.enqueue(left); },
121             None => {},
122         }
123
124         match front.get_right() {
125             Some(right) => { let _r = q.enqueue(right); },
126             None => {},
127         }
128     }
129 }

```

The following are the outputs after execution of binary search tree.

```

1  // bst.rs
2
3  fn main() {
4      basic();
5      order();
6
7      fn basic() {
8          let mut bst = BST::<i32, char>::new();
9          bst.insert(8, 'e'); bst.insert(6, 'c');
10         bst.insert(7, 'd'); bst.insert(5, 'b');
11         bst.insert(10, 'g'); bst.insert(9, 'f');
12         bst.insert(11, 'h'); bst.insert(4, 'a');
13
14         println!("bst is empty: {}", bst.is_empty());
15         println!("bst size: {}", bst.size());
16         println!("bst leaves: {}", bst.leaf_size());
17         println!("bst internals: {}", bst.none_leaf_size());
18         println!("bst depth: {}", bst.depth());
19
20         let min_kv = bst.min();
21         let max_kv = bst.max();
22         println!("min key-val: {:?}-{:?}", min_kv.0, min_kv.1);
23         println!("max key-val: {:?}-{:?}", max_kv.0, max_kv.1);
24         println!("bst contains 5: {}", bst.contains(&5));
25         println!("key: 5, val: {:?}", bst.get(&5).unwrap());
26     }
27
28     fn order() {
29         let mut bst = BST::<i32, char>::new();
30         bst.insert(8, 'e'); bst.insert(6, 'c');
31         bst.insert(7, 'd'); bst.insert(5, 'b');
32         bst.insert(10, 'g'); bst.insert(9, 'f');
33         bst.insert(11, 'h'); bst.insert(4, 'a');
34
35         println!("internal inorder, preorder, postorder: ");

```

```

36         bst.inorder();
37         bst.preorder();
38         bst.postorder();
39         bst.levelorder();
40         println!("outside inorder, preorder, postorder: ");
41         let nk = Some(Box::new(bst.clone()));
42         inorder(nk.clone());
43         preorder(nk.clone());
44         postorder(nk.clone());
45         levelorder(nk.clone());
46     }
47 }

```

The following are outputs after execution.

```

bst is empty: false
bst size: 8
bst leaves: 4
bst internals: 4
bst depth: 4
min key: Some(4), min val: Some('a')
max key: Some(11), max val: Some('h')
bst contains 5: true
key: 5, val: 'b'
internal inorder, preorder, postorder:
key: 4, val: 'a'
key: 5, val: 'b'
key: 6, val: 'c'
key: 7, val: 'd'
key: 8, val: 'e'
key: 9, val: 'f'
key: 10, val: 'g'
key: 11, val: 'h'
key: 8, val: 'e'
key: 6, val: 'c'
key: 5, val: 'b'
key: 4, val: 'a'
key: 7, val: 'd'
key: 10, val: 'g'
key: 9, val: 'f'
key: 11, val: 'h'
key: 4, val: 'a'
key: 5, val: 'b'
key: 7, val: 'd'
key: 6, val: 'c'
key: 9, val: 'f'
key: 11, val: 'h'
key: 10, val: 'g'
key: 8, val: 'e'
key: 8, val: 'e'
key: 6, val: 'c'
key: 10, val: 'g'
key: 5, val: 'b'

```

```

key: 7, val: 'd'
key: 9, val: 'f'
key: 11, val: 'h'
key: 4, val: 'a'
outside inorder, preorder, postorder:
key: 4, val: 'a'
key: 5, val: 'b'
key: 6, val: 'c'
key: 7, val: 'd'
key: 8, val: 'e'
key: 9, val: 'f'
key: 10, val: 'g'
key: 11, val: 'h'
key: 8, val: 'e'
key: 6, val: 'c'
key: 5, val: 'b'
key: 4, val: 'a'
key: 7, val: 'd'
key: 10, val: 'g'
key: 9, val: 'f'
key: 11, val: 'h'
key: 4, val: 'a'
key: 5, val: 'b'
key: 7, val: 'd'
key: 6, val: 'c'
key: 9, val: 'f'
key: 11, val: 'h'
key: 10, val: 'g'
key: 8, val: 'e'
key: 8, val: 'e'
key: 6, val: 'c'
key: 10, val: 'g'
key: 5, val: 'b'
key: 7, val: 'd'
key: 9, val: 'f'
key: 11, val: 'h'
key: 4, val: 'a'

```

When insert 76 into the tree, its search path is shown by the black nodes.

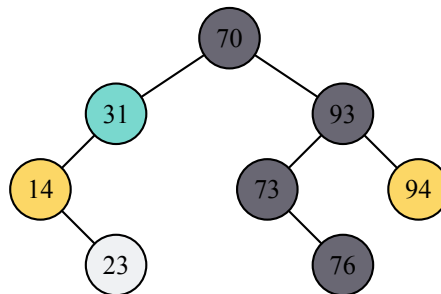


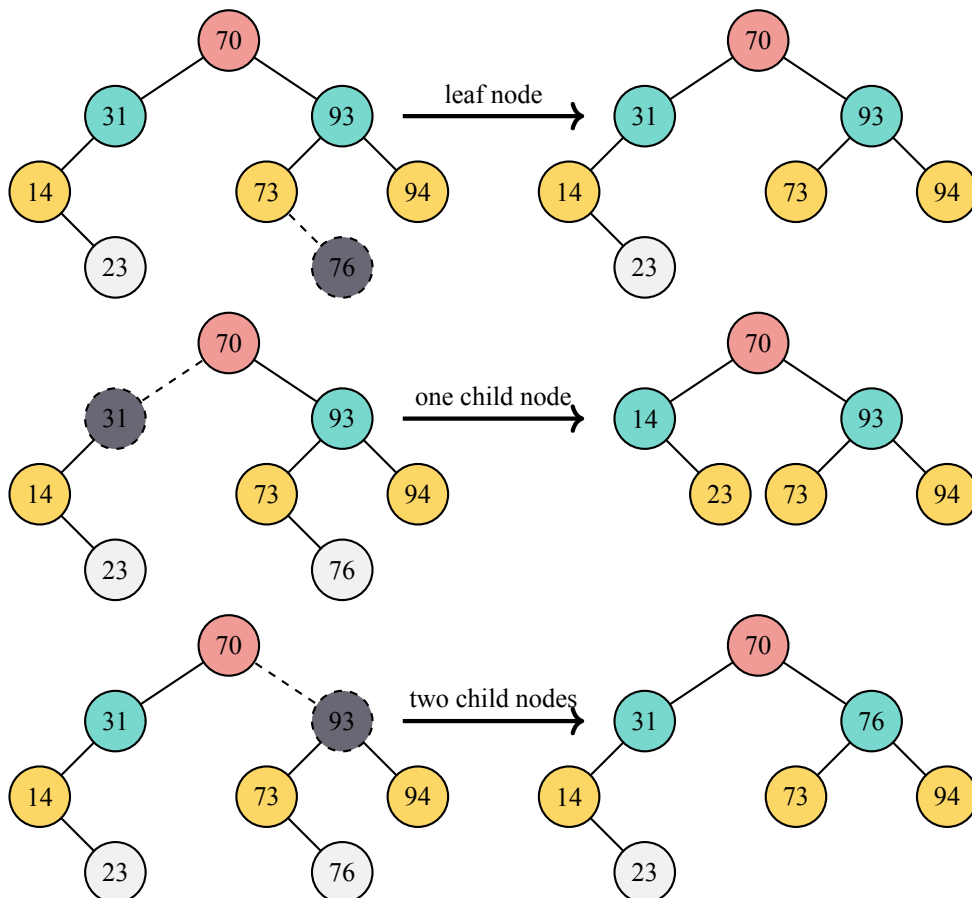
Figure 8.6: Insertion into a BST

Deleting a node in a binary search tree is a complex operation. The first step is to find the node to be deleted, which may not exist in the tree. Once the node is found, we need to determine if it has children, which can be one of three cases: no children, one child, or two children.

Table 8.4: Key deletion cases

No	Node	node k	delete method
1	No node	None	Directly return
2	One root node	None	Directly delete
3	Multiple nodes	None	Directly delete
4	Multiple nodes	One child	Replace k with child node
5	Multiple nodes	Two children	Replace k with the successor node

If the node is a leaf node with no children, we can simply remove its reference from its parent node. If it has one child, we modify the parent node's reference to point directly to the child node. The most challenging case is when the node has two children. In this case, we find the minimum node in the right subtree, called the successor node, and replace the node to be deleted with the successor node. The successor node may also have children, and we must adjust their relationships accordingly. The specific situations are shown in the following figure, where the dashed box represents the node k to be deleted, and the binary tree obtained after deleting the node is shown on the right.



Deleting a leaf node is the simplest case, while deleting an internal node with one child is straightforward. The most difficult case is when the node has two children, as this requires adjusting the relationships of multiple nodes.

8.4.3 Analysis of Binary Search Tree

Now that the binary search tree has been fully implemented, let's analyze the time complexity of each method. Traversals and `len()` both have a time complexity of $O(n)$ since they need to process all n elements. The `contains()` method compares data with the left and right child nodes and selects a branch based on the result. It walks the longest path from the root to the leaf node at most. The height of a binary tree is related to the total number of nodes.

$$\begin{aligned} 2^0 + 2^1 + 2^i \dots + 2^h &= n \\ h &= \log_2(n) \end{aligned} \quad (8.1)$$

Using the binary tree properties, we can approximate the maximum path length to be $h = \log_2(n)$. Thus, the time complexity of `contains()` is $O(\log_2(n))$. Insertion, deletion, and modification are based on search because the element must be located before processing can continue. Their time complexity depends only on search and can be completed in constant time. Therefore, the performance of `contains()`, `insert()`, and `get()` methods are all $O(\log_2(n))$. The height h of the binary tree is the limiting factor of their performance. If the inserted data is always in a sorted state, the binary tree may degenerate into a linear linked list, and the performance of `contains()`, `insert()`, and `remove()` methods will be $O(n)$. The figure below illustrates this scenario.

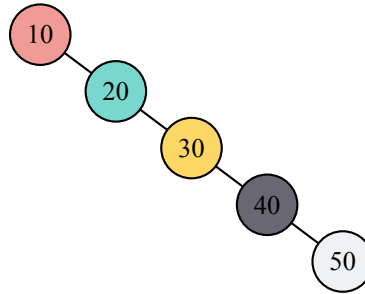


Figure 8.7: bst in a linear linked list

In this section, we did not implement the `remove()` function because it is not part of the abstract data type definition, and binary search trees are primarily used for data insertion and search, not deletion. However, interested readers can try to implement the `remove()` function as an exercise.

To improve performance by reducing the tree height, we can convert the binary tree into a multi-way tree such as B-trees and B+ trees. These trees have many child nodes, resulting in a short tree height and fast queries. They are commonly used in implementing databases and file systems. For instance, MySQL database uses B+ trees to store data, with nodes of 16K memory pages. If each data is 1k in size, one node can store 16 pieces of data. When used to store an index with a bigint key, 8 bytes are used, and the index is 6 bytes, for a total of 14 bytes. Thus, one node can store approximately $16 * 1024 / 14 = 1170$ indexes. With a height of 3, a B+ tree can store around $1170 * 1170 * 16 = 21902400$ indexes, which can hold about 20 million pieces of data. Retrieving data only requires at most two queries, which explains why database queries are fast. Readers interested in this topic can read MySQL-related books to learn more.

8.5 Balanced Binary Search Tree

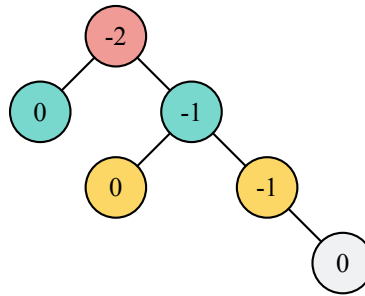
In the previous section, we constructed a binary search tree and learned that its performance can degrade to $O(n)$ in certain cases, such as when the tree is unbalanced with one side having significantly more nodes than the other. This inefficiency can lead to poor subsequent operations. To ensure efficient data processing, building a balanced binary tree is essential. In this section, we will discuss an AVL tree,

a balanced binary search tree that can automatically maintain balance. It is named after its inventors: G.M. Adelson-Velskii and E.M.Land.

The AVL tree is also an ordinary binary search tree, but with a difference in the way the tree operations are performed. AVL trees use a balance factor to determine if the tree is balanced during operations. The balance factor is the difference in height between the left and right subtrees of a node, and it is defined as:

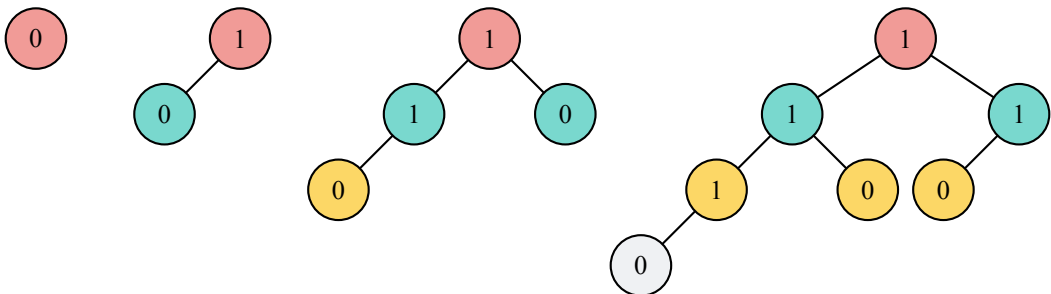
$$\text{balanceFactor} = \text{height}(\text{leftSubTree}) - \text{height}(\text{rightSubTree}) \quad (8.2)$$

Given this balance factor definition, if the balance factor is greater than zero, then the left subtree is heavy. If it is less than zero, then the right subtree is heavy, and if it is zero, then the tree is balanced. To implement an AVL tree efficiently, balance factors of -1, 0, and 1 are all considered balanced because the difference in height between the left and right subtrees is only 1 in these cases, which is essentially balanced. Once a node's balance factor is outside this range, such as 2 or -2, the tree needs to be rotated to maintain balance. The following figure illustrates the case of unbalanced left and right subtrees, and each node's balance factor is indicated by its value.



8.5.1 AVL Tree

To ensure a balanced binary tree, it is crucial to fulfill the balance factor condition, which allows for only three possible cases: left-heavy, balanced, or right-heavy. It is worth noting that a left-heavy or right-heavy tree that still meets the condition of having a balance factor of -1, 0, or 1 can also be considered balanced. The figure below depicts the most imbalanced left-heavy tree that meets the balance factor condition for trees of height 0, 1, 2, and 3.



By analyzing the total number of nodes in the tree, we can derive a formula for the number of nodes in a tree of height h , as follows: For a tree of height 0, there is only one node; for a tree of height 1, there are 2 nodes; for a tree of height 2, there are 4 nodes; for a tree of height 3, there are 7 nodes, and so on.

$$N_h = 1 + N_{h-1} + N_{h-2} \quad (8.3)$$

Remarkably, this formula resembles the Fibonacci sequence. With the number of nodes in the tree, we can obtain the height formula for an AVL tree using the Fibonacci equation, where the i th Fibonacci

number is given by the following equation:

$$\begin{aligned} F_0 &= 0 \\ F_1 &= 1 \\ F_i &= F_{i-1} + F_{i-2} \end{aligned} \tag{8.4}$$

To calculate the number of nodes in an AVL tree, we can use the following formula where $F_0 = 1$.

$$N_h = F_{h+2} - 1 \tag{8.5}$$

As i increases, the ratio F_i/F_{i-1} approaches the golden ratio $\Phi = (1 + \sqrt{5})/2$, so Φ can be used to represent F_i , which can be calculated as $F_i = \frac{\Phi^i}{\sqrt{5}}$.

$$N_h = \frac{\Phi^h}{\sqrt{5}} + 1 \tag{8.6}$$

Based on these formulas, we can derive a height formula for AVL trees.

$$\begin{aligned} \log(N_h - 1) &= \log\left(\frac{\Phi^h}{\sqrt{5}}\right) \\ \log(N_h - 1) &= h \log \Phi - \frac{1}{2} \log 5 \\ h &= \frac{\log(N_h - 1) + \frac{1}{2} \log 5}{\log \Phi} \\ h &= 1.44 \log(N_h) \end{aligned} \tag{8.7}$$

In this formula, h is the height of the AVL tree, and N_h is the number of nodes. This implies that the height of an AVL tree is at most 1.44 times the logarithm of the number of nodes, and the search complexity is $O(\log N)$, which is highly efficient.

8.5.2 Implementing a AVL Tree in Rust

To insert a new node into an AVL tree, we first add it as a leaf node, which automatically gives it a balance factor of 0. However, the balance factor of the parent node will need to be updated, depending on whether the new node is a left or right child. If it is a right child, the parent node's balance factor will be reduced by 1, and if it is a left child, the balance factor will be increased by 1. This update is propagated up the tree recursively, until we reach the root or a node whose balance factor becomes 0.

In our implementation, we represent the AVL tree as an enumeration with two cases: `Null`, which represents an empty tree, and `Tree`, which represents a node in the tree. The `AvlNode` type is used to store the data, left and right subtrees, and balance factor of each node.

```
1 // avl.rs
2
3 // Use a enum to define the Avl tree
4 #[derive(Clone, Debug, PartialEq)]
5 enum AvlTree<T> {
6     Null,
7     Tree(Box<AvlNode<T>>),
8 }
9
10 // Definish of Avl tree node
11 #[derive(Debug)]
12 struct AvlNode<T> {
```

```

13     key: T,
14     left: AvlTree<T>, // left subtree
15     right: AvlTree<T>, // right subtree
16     bfactor: i8,      // balance factor
17 }

```

To implement the AVL tree, we need to add two functions: insert and rebalance. To compare node data, we also introduce a comparison property called `Ord`. Additionally, we need the `replace` and `max` functions to update values and calculate the tree height.

```

1 // avl.rs
2
3 use std::cmp::{max, Ordering::*};
4 use std::fmt::Debug;
5 use std::mem::replace;
6 use AvlTree::*;
7
8 impl<T> AvlTree<T> where T : Clone + Ord + Debug {
9     // new tree is Empty
10    fn new() -> AvlTree<T> {
11        Null
12    }
13
14    fn insert(&mut self, key: T) -> (bool, bool) {
15        let ret = match self {
16            Null => {
17                // If there is no node, insert directly
18                let node = AvlNode {
19                    key: key,
20                    left: Null,
21                    right: Null,
22                    bfactor: 0,
23                };
24                *self = Tree(Box::new(node));
25
26                (true, true)
27            },
28            Tree(ref mut node) => match node.key.cmp(&key) {
29                // Compare the value of the node and determine
30                // which side to insert from
31                // inserted: whether insertion is performed
32                // deepened: whether the depth is increased
33
34                // If they are equal, no insertion is needed
35                Equal => (false, false),
36                // node value is smaller, insert to the right
37                Less => {
38                    let (inserted, deepened)
39                        = node.right.insert(key);
40                    if deepened {
41                        let ret = match node.bfactor {
42                            -1 => (inserted, false),
43                            0 => (inserted, true),

```

```

44             1 => (inserted, false),
45             _ => unreachable!(),
46         };
47         node.bfactor += 1;
48
49         ret
50     } else {
51         (inserted, deepened)
52     }
53 },
54 // node value is greater, insert to the left
55 Greater => {
56     let (inserted, deepened)
57     = node.left.insert(key);
58     if deepened {
59         let ret = match node.bfactor {
60             -1 => (inserted, false),
61             0 => (inserted, true),
62             1 => (inserted, false),
63             _ => unreachable!(),
64         };
65         node.bfactor -= 1;
66
67         ret
68     } else {
69         (inserted, deepened)
70     }
71 },
72 },
73 };
74 self.rebalance();
75
76     ret
77 }
78
79 // Adjust the balance factors of each node
80 fn rebalance(&mut self) {
81     match self {
82         // If there is no data, no adjustment is needed
83         Null => (),
84         Tree(_) => match self.node().bfactor {
85             // If the right subtree is heavy
86             -2 => {
87                 let lbf = self.node()
88                     .left
89                     .node()
90                     .bfactor;
91
92                 if lbf == -1 || lbf == 0 {
93                     let (a, b) = if lbf == -1 {
94                         (0, 0)
95                     } else {

```

```

96             (-1,1)
97         };
98
99         // Rotate and update the balance factor
100        self.rotate_right();
101        self.node().right.node().bfactor = a;
102        self.node().bfactor = b;
103    } else if lbf == 1 {
104        let (a, b) = match self.node()
105                        .left.node()
106                        .right.node()
107                        .bfactor
108        {
109            -1 => (1, 0),
110            0  => (0, 0),
111            1  => (0,-1),
112            _  => unreachable!(),
113        };
114
115        // First rotate left, then rotate right
116        // finally update the balance factor
117        self.node().left.rotate_left();
118        self.rotate_right();
119        self.node().right.node().bfactor = a;
120        self.node().left.node().bfactor = b;
121        self.node().bfactor = 0;
122    } else {
123        unreachable!()
124    }
125 },
126 // If the left subtree is heavy
127 2 => {
128     let rbf=self.node().right.node().bfactor;
129     if rbf == 1 || rbf == 0 {
130         let (a, b) = if rbf == 1 {
131             (0, 0)
132         } else {
133             (1,-1)
134         };
135
136         self.rotate_left();
137         self.node().left.node().bfactor = a;
138         self.node().bfactor = b;
139     } else if rbf == -1 {
140         let (a, b) = match self.node()
141                        .right.node()
142                        .left.node()
143                        .bfactor
144         {
145             1 => (-1,0),
146             0 => (0, 0),
147             -1 => (0, 1),

```

```

148         _ => unreachable!(),
149     };
150
151     // First rotate right, then rotate left
152     // finally update the balance factor
153     self.node().right.rotate_right();
154     self.rotate_left();
155     self.node().left.node().bfactor = a;
156     self.node().right.node().bfactor = b;
157     self.node().bfactor = 0;
158 } else {
159     unreachable!()
160 }
161 },
162 _ => (),
163 },
164 }
165 }
166 }

```

The insert function updates the balance factor recursively, while the rebalance function restores balance to the AVL tree. Effective rebalancing is crucial for maintaining proper AVL tree functionality without compromising performance. Rebalancing involves rotating the tree once or multiple times, which can be done through left or right rotations.

To perform a left rotation, we can follow the steps shown in Figure (8.8):

- (1) Promote the right child (B) as the new root of the subtree.
- (2) Move the old root (A) to the left child of the new root.
- (3) If the new root (B) already has a left child, make it the right child of the new left child (A).

Similarly, to perform a right rotation, we can follow these steps:

- (1) Promote the left child (B) as the new root of the subtree.
- (2) Move the old root (A) to the right child of the new root.
- (3) If the new root (B) already has a right child, make it the left child of the new right child (A).

The two trees in the figure are unbalanced, and we can rebalance them into the tree on the right by performing left and right rotations with A as the root.

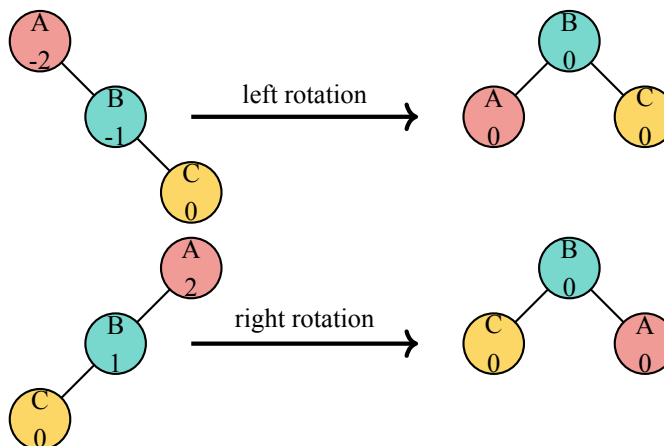
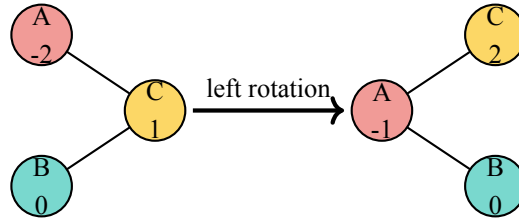


Figure 8.8: Unbalanced tree and its rotations

To perform rotations on a subtree, we can use left and right rotation rules. However, sometimes after performing one rotation, the balance may still be lost in the opposite direction.



In such cases, new rotation rules need to be used as follows:

- To balance a subtree with left rotation, we first check the balance factor of the right child node. If the right child is heavy, we perform a right rotation on it, and then a left rotation.
- Similarly, to balance a subtree with right rotation, we first check the balance factor of the left child node. If the left child is heavy, we perform a left rotation on it, and then a right rotation.

Although the process of subtree rotation is easy to understand conceptually, its code implementation can be complex because of the need to move nodes in the correct order while preserving all properties of the binary search tree. Additionally, ownership mechanisms in Rust make moving nodes and updating pointer relationships prone to errors. To implement rotation functions, we use node and subtree retrieval functions such as `left_subtree` and `right_subtree`.

```

1 // avl.rs
2
3 impl<T> AvlTree<T> where T : Ord {
4     // get a avlNode
5     fn node(&mut self) -> &mut AvlNode<T> {
6         match self {
7             Null => panic!("Empty tree"),
8             Tree(node) => node,
9         }
10    }
11
12    // get the left/right subtrees
13    fn left_subtree(&mut self) -> &mut Self {
14        match self {
15            Null => panic!("Error: Empty tree!"),
16            Tree(node) => &mut node.left,
17        }
18    }
19
20    fn right_subtree(&mut self) -> &mut Self {
21        match self {
22            Null => panic!("Error: Empty tree!"),
23            Tree(node) => &mut node.right,
24        }
25    }
26
27    fn rotate_left(&mut self) {
28        let mut n = replace(self, Null);
29        let mut right = replace(n.right_subtree(), Null);
30        let right_left = replace(right.left_subtree(), Null);
31        *n.right_subtree() = right_left;

```

```

32         *right.left_subtree() = n;
33         *self = right;
34     }
35
36     fn rotate_right(&mut self) {
37         let mut n = replace(self, Null);
38         let mut left = replace(n.left_subtree(), Null);
39         let left_right = replace(left.right_subtree(), Null);
40         *n.left() = left_right;
41         *left.right_subtree() = n;
42         *self = left;
43     }
44 }

```

To maintain the balance of the tree, we can perform rotation operations. However, we also need to implement methods to obtain important information about the tree, such as the number of nodes, node values, tree height, minimum and maximum values, and node queries. This requires implementing methods such as `size`, `leaf_size`, `depth`, `node`, `min`, `max`, and `contains` for the balanced binary tree.

```

1  // avl.rs
2
3  impl<T> AvlTree<T> where T : Ord {
4      // Calculate the nodes number: number of left and right
5      // child nodes + root node, calculated recursively
6      fn size(&self) -> usize {
7          match self {
8              Null => 0,
9              Tree(n) => 1 + n.left.size() + n.right.size(),
10         }
11     }
12
13     // Calculate the leaf nodes number: calculated recursively
14     fn leaf_size(&self) -> usize {
15         match self {
16             Null => 0,
17             Tree(node) => {
18                 if node.left == Null && node.right == null {
19                     return 1;
20                 }
21                 let left_leaf = match node.left {
22                     Null => 0,
23                     _ => node.left.leaf_size(),
24                 };
25                 let right_leaf = match node.right {
26                     Null => 0,
27                     _ => node.right.leaf_size(),
28                 };
29                 left_leaf + right_leaf
30             },
31         }
32     }
33
34     // Calculate the number of non-leaf nodes

```

```

35     fn none_leaf_size(&self) -> usize {
36         self.size() - self.leaf_size()
37     }
38
39     // The depth of the tree is the maximum depth of the left
40     // and right subtrees plus one, calculated recursively
41     fn depth(&self) -> usize {
42         match self {
43             Null => 0,
44             Tree(n) => max(n.left.depth(), n.right.depth()) + 1,
45         }
46     }
47
48     fn is_empty(&self) -> bool {
49         match self {
50             Null => true,
51             _ => false,
52         }
53     }
54
55     // Get the maximum and minimum node values of the tree
56     fn min(&self) -> Option<T> {
57         match self {
58             Null => None,
59             Tree(node) => {
60                 match node.left {
61                     Null => Some(&node.key),
62                     _ => node.left.min(),
63                 }
64             },
65         }
66     }
67
68     fn max(&self) -> Option<T> {
69         match self {
70             Null => None,
71             Tree(node) => {
72                 match node.right {
73                     Null => Some(&node.key),
74                     _ => node.right.min(),
75                 }
76             },
77         }
78     }
79
80     // determine if key is in the tree
81     fn contains(&self, key: &T) -> bool {
82         match self {
83             Null => false,
84             Tree(n) => {
85                 match n.key.cmp(&key) {
86                     Equal => { true },

```



```

87         Greater => {
88             match &n.left {
89                 Null => false,
90                 _ => n.left.contains(key),
91             }
92         },
93         Less => {
94             match &n.right {
95                 Null => false,
96                 _ => n.right.contains(key),
97             }
98         },
99     },
100 },
101 }
102 }
103 }

```

In addition to these methods, we can implement the four traversal methods commonly used for binary trees: *inorder*, *preorder*, *postorder*, and *levelorder*. These traversal methods allow us to visit and manipulate each node in the tree in a specific order, which can be useful in various scenarios such as printing the tree, searching for a specific node, or computing some statistics about the tree.

```

1 // avl.rs
2
3 impl<T> AVLTree<T> where T : Ord {
4     // Internal implementation of preorder, inorder, postorder,
5     // and level-order traversal
6     fn preorder(&self) {
7         match self {
8             Null => (),
9             Tree(node) => {
10                 println!("key: {:?}", node.key);
11                 node.left.preorder();
12                 node.right.preorder();
13             },
14         }
15     }
16
17     fn inorder(&self) {
18         match self {
19             Null => (),
20             Tree(node) => {
21                 node.left.inorder();
22                 println!("key: {:?}", node.key);
23                 node.right.inorder();
24             },
25         }
26     }
27
28     fn postorder(&self) {
29         match self {
30             Null => (),

```

```

31         Tree(node) => {
32             node.left.postorder();
33             node.right.postorder();
34             println!("key: {:?}", node.key);
35         },
36     }
37 }
38
39 fn levelorder(&self) {
40     let size = self.size();
41     let mut q = Queue::new(size);
42
43     let _r = q.enqueue(self);
44     while !q.is_empty() {
45         let front = q.dequeue().unwrap();
46         match front {
47             Null => (),
48             Tree(node) => {
49                 println!("key: {:?}", node.key);
50                 let _r = q.enqueue(&node.left);
51                 let _r = q.enqueue(&node.right);
52             },
53         }
54     }
55 }
56 }
57
58 // External implementation of preorder, inorder, postorder,
59 // and level-order traversal
60 fn preorder<T: Clone + Ord + Debug>(avl: &AvlTree<T>) {
61     match avl {
62         Null => (),
63         Tree(node) => {
64             println!("key: {:?}", node.key);
65             preorder(&node.left);
66             preorder(&node.right);
67         },
68     }
69 }
70
71 fn inorder<T: Clone + Ord + Debug>(avl: &AvlTree<T>) {
72     match avl {
73         Null => (),
74         Tree(node) => {
75             inorder(&node.left);
76             println!("key: {:?}", node.key);
77             inorder(&node.right);
78         },
79     }
80 }
81
82 fn postorder<T: Clone + Ord + Debug>(avl: &AvlTree<T>) {

```

```

83     match avl {
84         Null => (),
85         Tree(node) => {
86             postorder(&node.left);
87             postorder(&node.right);
88             println!("key: {:?}", node.key);
89         },
90     }
91 }
92
93 fn levelorder<T: Clone + Ord + Debug>(avl: &AvlTree<T>) {
94     let size = avl.size();
95     let mut q = Queue::new(size);
96     let _r = q.enqueue(avl);
97     while !q.is_empty() {
98         let front = q.dequeue().unwrap();
99         match front {
100             Null => (),
101             Tree(node) => {
102                 println!("key: {:?}", node.key);
103                 let _r = q.enqueue(&node.left);
104                 let _r = q.enqueue(&node.right);
105             },
106         }
107     }
108 }
109
110 fn main() {
111     basic();
112     order();
113
114     fn basic() {
115         let mut t = AvlTree::new();
116         for i in 0..5 { let (_r1, _r2) = t.insert(i); }
117
118         println!("empty:{},size:{},t.is_empty(),t.size()");
119         println!("leaves:{},depth:{},t.leaf_size(),t.depth()");
120         println!("internals:{},t.none_leaf_size()");
121         println!("min-max key:{:?}-{:?}", t.min(), t.max());
122         println!("contains 9:{},t.contains(&9)");
123     }
124
125     fn order() {
126         let mut avl = AvlTree::new();
127         for i in 0..5 { let (_r1, _r2) = avl.insert(i); }
128
129         println!("internal pre-in-post-level order");
130         avl.preorder(); avl.inorder();
131         avl.postorder(); avl.levelorder();
132         println!("outside pre-in-post-level order");
133         preorder(&avl); inorder(&avl);
134         postorder(&avl); levelorder(&avl);

```

```

135     }
136 }

```

The following are outputs after execution.

```

empty:false,size:5
leaves:3,depth:3
internals:2
min-max key:Some(0)-Some(4)
contains 9:false
internal pre-in-pos-level order
key: 1
key: 0
key: 3
key: 2
key: 4
key: 0
key: 1
key: 2
key: 3
key: 4
key: 0
key: 2
key: 4
key: 3
key: 1
key: 1
key: 0
key: 3
key: 2
key: 4
outside pre-in-pos-level order
key: 1
key: 0
key: 3
key: 2
key: 4
key: 0
key: 1
key: 2
key: 3
key: 4
key: 0
key: 2
key: 4
key: 3
key: 1
key: 1
key: 0
key: 3
key: 2
key: 4

```

8.5.3 Analysis of AVL Tree

An AVL balanced binary tree includes a rebalancing function and left and right rotation operations to maintain its balance, which enables its operations to maintain a relatively good performance with a worst-case complexity of $O(\log_2(n))$. However, it is important to note that AVL trees perform a significant number of rotation operations to maintain balance. A red-black tree is an optimized binary tree that requires fewer rotations compared to an AVL tree. It is a weakened version of the AVL tree and can be used in scenarios with frequent insertions, deletions, and modifications to achieve better performance.

8.6 Summary

This chapter introduced us to trees, an efficient data structure that enables us to implement a wide range of useful algorithms. Trees find extensive applications in storage, networking, and other domains. Throughout this chapter, we completed the following tasks using trees:

- Parse and evaluated expressions.
- Implemented binary heaps as priority queues.
- Implemented binary trees, binary search trees, and balanced binary search trees.

In the previous chapters, we learned about several abstract data types used to implement mapping relationships (Maps), including ordered tables, hash tables, binary search trees, and balanced binary search trees. The table below compares the worst-case performance of various operations supported by these data types. While red-black trees are an improvement over AVL trees, their complexity and performance remain similar to AVL trees, with only differences in coefficients. For further information, readers are encouraged to consult relevant resources.

Table 8.5: Performance of various abstract data types

Operation	Ordered table	Hash table	BST	AVL	Red-black tree
insert	$O(n)$	$O(1)$	$O(n)$	$O(\log(n))$	$O(\log(n))$
contains	$O(\log(n))$	$O(1)$	$O(n)$	$O(\log(n))$	$O(\log(n))$
delete	$O(n)$	$O(1)$	$O(n)$	$O(\log(n))$	$O(\log(n))$

Chapter 9

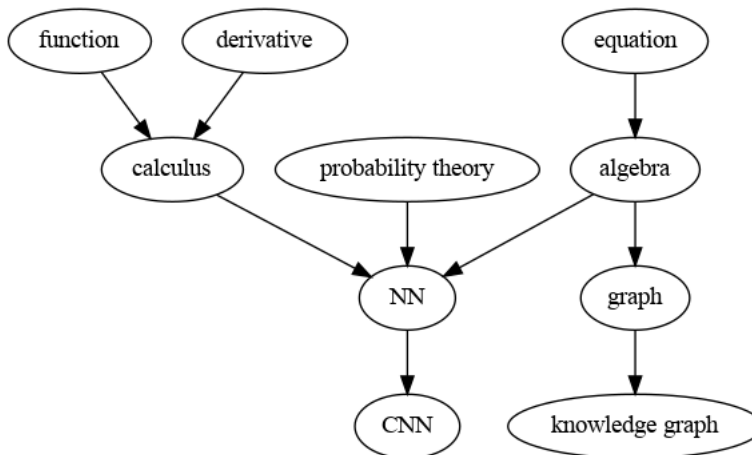
Graphs

9.1 Objectives

- Understand the concept and storage format of graphs.
- Implement several graph data structures in Rust.
- Learn two important graph search algorithms.
- Use graphs to solve various real-world problems.

9.2 What is Graph?

In the previous chapter, we discussed binary trees as a data structure, while this chapter focuses on graphs - a more generalized form of trees. Although trees can be viewed as simplified graphs or carefully selected graphs with certain constraints on their node relationships, graphs have no such limitations. While trees have a root node and a top-to-bottom direction, graphs lack both. Unlike trees, graphs can contain cycles. They consist of points, edges, and point-edge relationships, whereas trees consist of nodes and connections.



The figure above illustrates a graph with multiple nodes and connections, which demonstrates the vertical and horizontal relationships between different items. For example, calculus and linear algebra have no direct connection, but they can jointly incubate neural networks, forming a horizontal relationship. Derivatives, on the other hand, contribute to the construction of calculus, which is a part of calculus, and this forms a vertical relationship.

Graphs are versatile data structures that can be used to represent various real-world phenomena, such as flight maps, social network graphs, and course planning graphs. In this chapter, we will learn about the concept and storage format of graphs, and how to implement them in Rust. We will also explore two essential graph search algorithms that can be applied to solve different problems. Research on graphs, their various applications, and algorithms belong to a specialized discipline called graph theory.

9.2.1 Vocabularies and Definitions of Graph

A graph is a more general form of a tree and is defined by the following concepts:

- Vertex (or node): an element of the graph that has a name, or key, and may have additional information, or payload.
- Edge: an element that connects two vertices and indicates a relationship between them. An edge can be unidirectional or bidirectional, and may have a weight, which represents a measure such as distance, cost, time, or closeness.
- Weight: a measure of an edge.

A graph can be represented as $G = (V, E)$, where V is the set of vertices and E is the set of edges. The figure below shows an example of a weighted directed graph with a vertex set $V = (V0, V1, V2, V2, V4, V5)$ and an edge set E consisting of various connections and their weights, with a total of 9 edges.

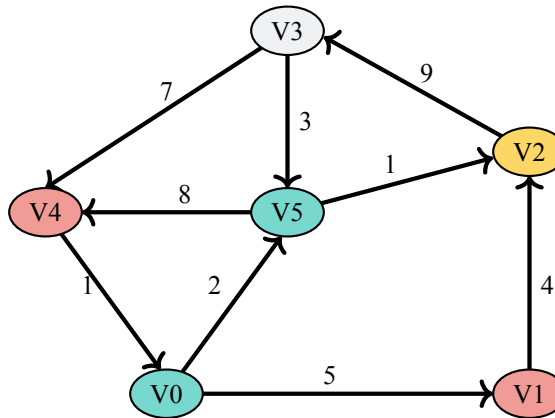


Figure 9.1: Graph

In addition to vertices, edges, and weights, paths are used to represent the order of points. A path is a vertex sequence that represents the order of point connections, such as (v, w, x, y, z) . Cyclic graphs, which have cycles, such as $V5 \rightarrow V2 \rightarrow V3 \rightarrow V5$, may also exist. If there are no cycles in the graph, it is called an acyclic graph or DAG graph. Many important problems can be represented using DAG graphs.

9.3 Graph Storage Format

To store a non-linear graph in a linear storage device like computer memory, there are two commonly used methods: adjacency matrix and adjacency list.

The adjacency matrix stores the nodes, edges, and weights of the graph in a two-dimensional matrix, which requires N^2 space for a graph with N nodes. However, this method may result in a large waste of space when the graph is sparse. In contrast, the adjacency list is similar to a hash table and saves all the edges related to each vertex by opening a chain for each vertex. Its storage space depends on the number of edges and is generally much smaller than N^2 . Therefore, computers use adjacency lists to store graph data structures.

9.3.1 Adjacency Matrix

An adjacency matrix is a straightforward way to store a graph. In this method, the vertices of the graph are represented by the rows and columns of a two-dimensional matrix. The value stored at the intersection of row v and column w indicates the presence of an edge between vertex v and vertex w , with the value indicating the weight of the edge. When two vertices have an edge between them, they are called adjacent. The graph shown in (9.1) can be stored using an adjacency matrix.

	V0	V1	V2	V3	V4	V5
V0		5				2
V1			4			
V2				9		
V3					7	3
V4	1					
V5			1		8	

Figure 9.2: Adjacency matrix

Although the adjacency matrix is easy to understand and analyze for small graphs, it suffers from the drawback of being very sparse. Most of the cells in the matrix are empty, leading to a significant waste of space. For a graph with N vertices, an adjacency matrix requires N^2 storage. Therefore, the adjacency matrix is not an efficient method to store large graphs.

9.3.2 Adjacency List

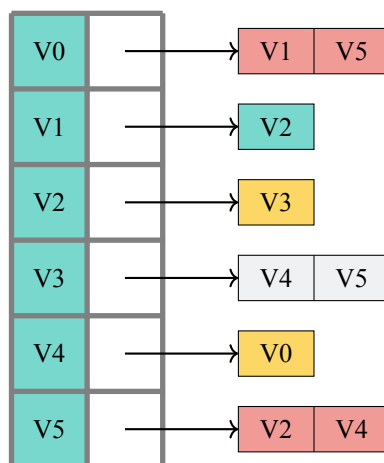


Figure 9.3: Adjacency list

An efficient way to save graphs is by using an adjacency list, as shown in the figure above. In this approach, an array stores all the vertices in the graph, and each vertex maintains a linked list that connects it to other vertices. By accessing the linked lists of various vertices, it is possible to know how many points it is connected to, which is similar to the chaining method used by HashMap to solve collisions.

The adjacency list stores the linked points in a structure similar to HashMap because edges have weights, while arrays can only save vertices. Graphs implemented with adjacency lists are compact and have no memory waste. This structure is also very convenient for storage. The importance of basic data structures is evident here, and the HashMap implemented in previous chapters is useful for implementing adjacency lists.

9.4 The Graph Abstract Data Type

The graph abstract data type can be defined based on the basic definition of a graph. At the core of the graph are vertices and edges, and operations are defined around them. The following operations are defined for a graph:

- `new()`: Creates a new empty graph with no parameters and returns an empty graph.
- `add_vertex(v)`: Adds a vertex with parameter `v` to the graph and has no return value.
- `add_edge(fv, tv, w)`: Adds a weighted directed edge with parameters `fv`, `tv`, and `w` to the graph and has no return value.
- `get_vertex(vk)`: Finds the vertex with key `vk` in the graph, requires parameter `vk`, and returns the vertex.
- `get_vertices()`: Returns a list of all vertices in the graph without parameters.
- `vert_nums()`: Returns the number of vertices in the graph without parameters.
- `edge_nums()`: Returns the number of edges in the graph without parameters.
- `contains(vk)`: Checks whether the vertex is in the graph with parameter `vk` and returns a Boolean value.
- `is_empty()`: Checks whether the graph is empty without parameters and returns a Boolean value.

Assuming `g` is a newly created empty graph, the table below shows the results of various graph operations. `[]` stores vertices, and `()` stores edges, where the first two values are points and the third value is the edge weight, such as `(1, 5, 2)`, which means that the edge weight between point 1 and point 5 is 2.

Table 9.1: Operations on Graph

Operation	Graph value	Return value
<code>g.is_empty()</code>	<code>[]</code>	<code>true</code>
<code>g.add_vertex(1)</code>	<code>[1]</code>	
<code>g.add_vertex(5)</code>	<code>[1,5]</code>	
<code>g.add_edge(1,5,2)</code>	<code>[1,5,(1,5,2)]</code>	
<code>g.get_vertex(5)</code>	<code>[1,5,(1,5,2)]</code>	<code>5</code>
<code>g.get_vertex(4)</code>	<code>[1,5,(1,5,2)]</code>	<code>None</code>
<code>g.edge_nums()</code>	<code>[1,5,(1,5,2)]</code>	<code>1</code>
<code>g.vert_nums()</code>	<code>[1,5,(1,5,2)]</code>	<code>2</code>
<code>g.contains(1)</code>	<code>[1,5,(1,5,2)]</code>	<code>true</code>
<code>g.get_vertices()</code>	<code>[1,5,(1,5,2)]</code>	<code>[1,5]</code>
<code>g.add_vertex(7)</code>	<code>[1,5,7,(1,5,2)]</code>	
<code>g.add_vertex(9)</code>	<code>[1,5,7,9,(1,5,2)]</code>	
<code>g.add_edge(7,9,8)</code>	<code>[1,5,7,9,(1,5,2),(7,9,8)]</code>	
<code>g.contains(3)</code>	<code>[1,5,7,9,(1,5,2),(7,9,8)]</code>	<code>false</code>
<code>g.vert_nums()</code>	<code>[1,5,7,9,(1,5,2),(7,9,8)]</code>	<code>4</code>
<code>g.edge_nums()</code>	<code>[1,5,7,9,(1,5,2),(7,9,8)]</code>	<code>2</code>

9.5 Implementing a Graph in Rust

To implement a graph based on an adjacency matrix in Rust, we need to first create vertices and edges, and then use a two-dimensional Vec to store the edge relationships.

```

1 // graph_matrix.rs
2
3 // Definition of graph vertex
4 #[derive(Debug)]
5 struct Vertex<'a> {
6     id: usize,
7     name: &'a str,
8 }
9
10 impl Vertex<'_> {
11     fn new(id: usize, name: &'static str) -> Self {
12         Self {
13             id,
14             name
15         }
16     }
17 }
```

We can represent edges using a Boolean value to indicate their existence, as they are simply relationships and do not need to be constructed as entities.

```

1 // graph_matrix.rs
2
3 // Definition of graph edge
4 #[derive(Debug, Clone)]
5 struct Edge {
6     edge: bool, // boolean value to show whethe edge exists
7 }
8
9 impl Edge {
10     fn new() -> Self {
11         Self { edge: false }
12     }
13
14     fn set_edge() -> Self {
15         Edge { edge: true }
16     }
17 }
```

Therefore, the edge relationships can be implemented and stored in a two-dimensional Vec.

```

1 // graph_matrix.rs
2
3 // Definition for Graph
4 #[derive(Debug)]
5 struct Graph {
6     nodes: usize,
7     graph: Vec<Vec<Edge>>, // store edges of a vertex
8 }
```

```

9
10 impl Graph {
11     fn new(nodes: usize) -> Self {
12         Self {
13             nodes,
14             graph: vec![vec![Edge::new(); nodes]; nodes],
15         }
16     }
17
18     fn is_empty(&self) -> bool {
19         0 == self.nodes
20     }
21
22     fn len(&self) -> usize { self.nodes }
23
24     // add a edge and set the edge property to true
25     fn add_edge(&mut self, n1: &Vertex, n2: &Vertex) {
26         if n1.id < self.nodes && n2.id < self.nodes {
27             self.graph[n1.id][n2.id] = Edge::set_edge();
28         } else {
29             println!("Error, vertex beyond the graph");
30         }
31     }
32 }
33
34 fn main() {
35     let mut g = Graph::new(4);
36     let n1 = Vertex::new(0, "n1"); let n2 = Vertex::new(1, "n2");
37     let n3 = Vertex::new(2, "n3"); let n4 = Vertex::new(3, "n4");
38     g.add_edge(&n1, &n2); g.add_edge(&n1, &n3);
39     g.add_edge(&n2, &n3); g.add_edge(&n2, &n4);
40     g.add_edge(&n3, &n4); g.add_edge(&n3, &n1);
41     println!("{:?}", g);
42     println!("graph empty: {}", g.is_empty());
43     println!("graph nodes: {}", g.len());
44 }

```

The following are outputs after execution.

```

Graph {
  nodes: 4,
  graph: [
    [
      Edge { edge: false, },
      Edge { edge: true, },
      Edge { edge: true, },
      Edge { edge: false, },
    ],
    [
      Edge { edge: false, },
      Edge { edge: false, },
      Edge { edge: true, },
      Edge { edge: true, },
    ],
  ],
}

```

```

    ],
    [
        Edge { edge: true, },
        Edge { edge: false, },
        Edge { edge: false, },
        Edge { edge: true, },
    ],
    [
        Edge { edge: false, },
        Edge { edge: false, },
        Edge { edge: false, },
        Edge { edge: false, },
    ],
],
}
graph empty: false
graph nodes: 4

```

Next, we will implement an adjacency list graph using `HashMap`. Since vertices are the core elements and edges are the relationships between vertices, we need to create a data structure `Vertex` to represent the vertex element. For `Vertex`, we need to define operations such as creating a new vertex, getting the value of the vertex itself, adding adjacent vertices, getting all adjacent vertices, and getting the weights of adjacent vertices. The `neighbors` variable is used to store all adjacent vertices of the current vertex.

```

1 // graph_adjlist.rs
2
3 use std::hash::Hash;
4 use std::collections::HashMap;
5
6 // Definision of Vertex
7 #[derive(Debug, Clone)]
8 struct Vertex<T> {
9     key: T,
10    neighbors: Vec<(T, i32)>, // store adjacent vertices
11 }
12
13 impl<T: Clone + PartialEq> Vertex<T> {
14     fn new(key: T) -> Self {
15         Self {
16             key: key,
17             neighbors: Vec::new()
18         }
19     }
20
21     // Check if a point is adjacent to the current point
22     fn adjacent_key(&self, key: &T) -> bool {
23         for (nbr, _wt) in self.neighbors.iter() {
24             if nbr == key {
25                 return true;
26             }
27         }
28     }

```

```

29         false
30     }
31
32     fn add_neighbor(&mut self, nbr: T, wt: i32) {
33         self.neighbors.push((nbr, wt));
34     }
35
36     // Get the set of adjacent points
37     fn get_neighbors(&self) -> Vec<&T> {
38         let mut neighbors = Vec::new();
39         for (nbr, _wt) in self.neighbors.iter() {
40             neighbors.push(nbr);
41         }
42
43         neighbors
44     }
45
46     // Return the edge weight to the adjacent point
47     fn get_nbr_weight(&self, key: &T) -> &i32 {
48         for (nbr, wt) in self.neighbors.iter() {
49             if nbr == key {
50                 return wt;
51             }
52         }
53
54         &0
55     }
56 }

```

Graph is a data structure used to implement graphs, which includes a HashMap that maps vertex names to vertex objects.

```

1 // graph_adjlist.rs
2
3 // Definish of Graph
4 #[derive(Debug, Clone)]
5 struct Graph <T> {
6     vertnums: u32, // count of vertices
7     edgenums: u32, // count of edges
8     vertices: HashMap<T, Vertex<T>>,
9 }
10
11 impl<T: Hash + Eq + PartialEq + Clone> Graph<T> {
12     fn new() -> Self {
13         Self {
14             vertnums: 0,
15             edgenums: 0,
16             vertices: HashMap::<T, Vertex<T>>::new(),
17         }
18     }
19
20     fn is_empty(&self) -> bool { 0 == self.vertnums }
21 }

```

```

22     fn vertex_num(&self) -> u32 { self.vertnums }
23
24     fn edge_num(&self) -> u32 { self.edgenums }
25
26     fn contains(&self, key: &T) -> bool {
27         for (nbr, _vertex) in self.vertices.iter() {
28             if nbr == key { return true; }
29         }
30
31         false
32     }
33
34     fn add_vertex(&mut self, key: &T) -> Option<Vertex<T>> {
35         let vertex = Vertex::new(key.clone());
36         self.vertnums += 1;
37         self.vertices.insert(key.clone(), vertex)
38     }
39
40     fn get_vertex(&self, key: &T) -> Option<&Vertex<T>> {
41         if let Some(vertex) = self.vertices.get(key) {
42             Some(&vertex)
43         } else {
44             None
45         }
46     }
47
48     // Get the keys of all nodes
49     fn vertex_keys(&self) -> Vec<T> {
50         let mut keys = Vec::new();
51         for key in self.vertices.keys() {
52             keys.push(key.clone());
53         }
54
55         keys
56     }
57
58     // Delete a point (and its edges)
59     fn remove_vertex(&mut self, key: &T) -> Option<Vertex<T>> {
60         let old_vertex = self.vertices.remove(key);
61         self.vertnums -= 1;
62
63         // Delete edges from the current point
64         self.edgenums -= old_vertex.clone()
65             .unwrap()
66             .get_neighbors()
67             .len() as u32;
68
69         // Delete edges to the current point
70         for vertex in self.vertex_keys() {
71             if let Some(vt) = self.vertices.get_mut(&vertex) {
72                 if vt.adjacent_key(key) {
73                     vt.neighbors.retain(|(k, _)| k != key);

```

```

74             self.edgenums -= 1;
75         }
76     }
77 }
78
79     old_vertex
80 }
81
82 fn add_edge(&mut self, from: &T, to: &T, wt: i32) {
83     // If the point doesn't exist, add it first
84     if !self.contains(from) {
85         let _fv = self.add_vertex(from);
86     }
87     if !self.contains(to) {
88         let _tv = self.add_vertex(to);
89     }
90
91     // Add an edge
92     self.edgenums += 1;
93     self.vertices.get_mut(from)
94         .unwrap()
95         .add_neighbor(to.clone(), wt);
96 }
97
98 // Determine if two vertices are adjacent
99 fn adjacent(&self, from: &T, to: &T) -> bool {
100     self.vertices.get(from).unwrap().adjacent_key(to)
101 }
102 }

```

Using Graph, we can create the vertices V0-V5 and their edges as shown in the graph (9.1).

```

1 // graph_adjlist.rs
2
3 fn main() {
4     let mut g = Graph::new();
5
6     for i in 0..6 { g.add_vertex(&i); }
7     println!("graph empty: {}", g.is_empty());
8
9     let vertices = g.vertex_keys();
10    for vtx in vertices { println!("Vertex: {:#?}", vtx); }
11
12    g.add_edge(&0,&1,5); g.add_edge(&0,&5,2);
13    g.add_edge(&1,&2,4); g.add_edge(&2,&3,9);
14    g.add_edge(&3,&4,7); g.add_edge(&3,&5,3);
15    g.add_edge(&4,&0,1); g.add_edge(&4,&4,8);
16    println!("vert nums: {}", g.vertex_num());
17    println!("edge nums: {}", g.edge_num());
18    println!("contains 0: {}", g.contains(&0));
19
20    let vertex = g.get_vertex(&0).unwrap();
21    println!("key: {}, to nbr 1 weight: {}",

```

```

22         vertex.key, vertex.get_nbr_weight(&1));
23
24     let keys = vertex.get_neighbors();
25     for nbr in keys { println!("nighbor: {nbr}"); }
26
27     for (nbr, wt) in vertex.neighbors.iter() {
28         println!("0 nighbor: {nbr}, weight: {wt}");
29     }
30
31     let res = g.adjacent(&0, &1);
32     println!("0 adjacent to 1: {res}");
33     let res = g.adjacent(&3, &2);
34     println!("3 adjacent to 2: {res}");
35
36     let rm = g.remove_vertex(&0).unwrap();
37     println!("remove vertex: {}", rm.key);
38     println!("left vert nums: {}", g.vertex_num());
39     println!("left edge nums: {}", g.edge_num());
40     println!("contains 0: {}", g.contains(&0));
41 }

```

The following are outputs after execution.

```

graph empty: false
Vertex: 3
Vertex: 4
Vertex: 1
Vertex: 2
Vertex: 5
Vertex: 0
vert nums: 6
edge nums: 8
contains 0: true
key: 0, to nbr 1 weight: 5
nighbor: 1
nighbor: 5
0 nighbor: 1, weight: 5
0 nighbor: 5, weight: 2
0 is adjacent to 1: true
3 is adjacent to 2: false
remove vertex: 0
left vert nums: 5
left edge nums: 5
contains 0: false

```

9.5.1 The Word Ladder Problem

The graph data structure enables us to solve practical problems, such as the word ladder puzzle. In this puzzle, we must transform one word into another by changing one letter at a time, (FOOL -> SAGE), but the resulting word must exist in the dictionary. This game was invented by Lewis Carroll, the author of "Alice's Adventures in Wonderland," in 1878. Multiple solutions exist for this problem, as shown in the word sequence below.

a	b	c	d	e	f	g
FOOL	FOOL	FOOL	FOOL	FOOL	FOOL	FOOL
POOL	FOIL	FOIL	COOL	COOL	FOUL	FOUL
POLL	FAIL	FAIL	POOL	POOL	FOIL	FOIL
POLE	FALL	FALL	POLL	POLL	FALL	FALL
PALE	PALL	PALL	POLE	PALL	FALL	FALL
SALE	PALE	PALE	PALE	PALE	PALL	PALL
SAGE	PAGE	SALE	SALE	SALE	POLL	POLL
	SAGE	SAGE	SAGE	SAGE	PALE	PALE
					PAGE	SALE
					SAGE	SAGE

Our goal is to use graph algorithms to find the shortest path between the starting and ending words, as shown in column a of the figure above. To achieve this, we first convert the words into vertices and link together the words that can be transformed into each other. If two words differ by only one letter, we create a bidirectional edge between them, as shown in the figure below. Finally, we use a graph search algorithm to find the shortest path between the starting and ending vertices.

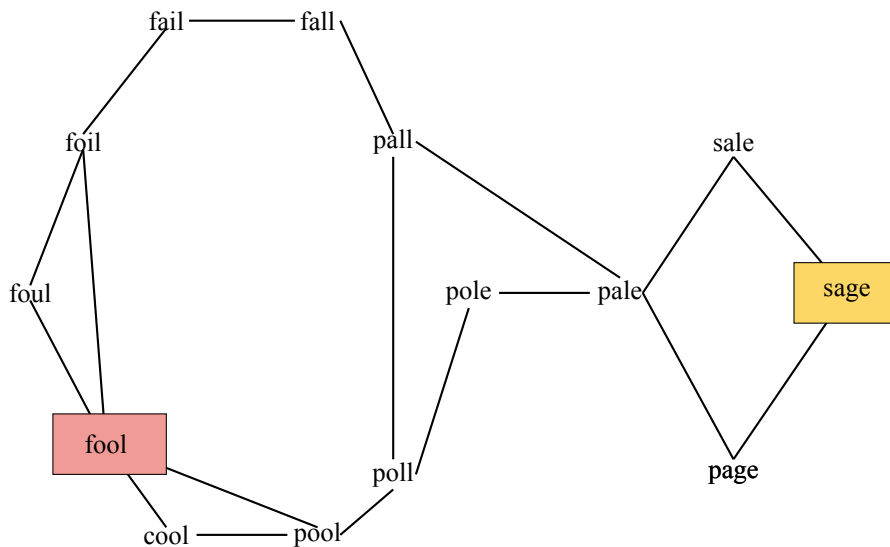


Figure 9.4: The word ladder

To create a graph model for the word ladder problem, there are various methods available. When dealing with a list of words with the same length, a vertex can be created for each word in the list. To connect the words, each word in the list must be compared to the others. If two words being compared have only one letter different, an edge can be created between them in the graph. This method is feasible for small word lists, but for larger lists such as the CET-4 vocabulary list with 3000 words or the CET-6 list with 5000 words, the comparison is inefficient, requiring millions of comparisons.

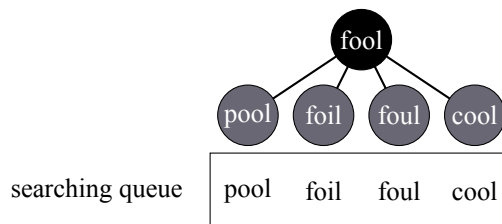
An alternative approach is to consider the letter positions of the words and group them based on similar patterns. This way, we can search for similar words at each position and group them together. For instance, if we remove the first letter of SOPE and keep the rest "_OPE," all four-letter words ending in "OPE" should be connected to SOPE and collected into a set. The same process is used for "S_PE" and all the words matching this pattern are collected. Finally, all the words matching this pattern are collected, as shown in the figure below.

_OPE	P_PE	PO_E	POP_
POPE	POPE	POPE	POPE
ROPE	PIPE	POLE	POPS
NOPE	PAPE	PORE	
HOPE		POSE	
LOPE		POKE	
COPE			

This solution can be implemented using a HashMap. The set mentioned earlier stores words with the same pattern as keys in the HashMap, which stores similar sets of words. Once the word sets are established, the graph can be created. To start the graph, create a vertex for each word in the HashMap, and then create edges between all vertices found under the same key in the dictionary. Once the graph is implemented, the ladder search task can be completed.

9.6 Breadth First Search(BFS)

The study of graphs and graph theory is a complex subject matter with various excellent algorithms developed for graph study. However, the Breadth-First Search (BFS) and Depth-First Search (DFS) algorithms are the most commonly used ones. In the case of the ladder problem, BFS is used to find the shortest path since graphs are non-linear and traditional binary and linear search algorithms cannot be used for this task.



The BFS algorithm explores the edges in the graph to find all vertices in G (assuming the distance between points is 1) that have a path starting from a given vertex s . It does this by first finding all vertices that are one unit away from s , then all vertices that are two units away, and so on, until all vertices are found. The search is performed layer by layer, which is why it is called breadth-first search. The points connected to the starting vertex are considered as one layer, put into a queue first, and then the algorithm finds the next layer of connecting points for those in the queue and repeats the process until the search is complete.

During the search, vertices can be colored to indicate their status. Initially, all vertices are white. As the search progresses, vertices connected to the current search vertex are set to gray. The algorithm checks each gray vertex, and if it is not the target value, it is set to black, and the search continues for white vertices connected to the gray vertex. This process continues until the search task is completed or the entire graph search is complete. This search method is similar to the garbage collection mechanism of some languages, such as the tricolor garbage collection in the Go language.

9.6.1 Implementing a BFS

To implement a simple BFS algorithm, we first need to create a graph. The specific form of the graph depends on the purpose of the algorithm. Here, we will demonstrate the BFS algorithm using a simple graph, where nodes contain data and a link to the next node. We will store only the first node and the link to the next node in the graph.

```

1 // bfs.rs
2
3 use std::rc::Rc;
4 use std::cell::RefCell;
5
6 // Node has multiple shared links,
7 // Box cannot be shared, only Rc can be shared.
8 // Rc is immutable, RefCell with internal mutability
9 // is used to wrap it.
10 type Link = Option<Rc<RefCell<Node>>>>;
11
12 // Definition for a Node
13 struct Node {
14     data: usize,
15     next: Link,
16 }
17
18 impl Node {
19     fn new(data: usize) -> Self {
20         Self {
21             data: data,
22             next: None
23         }
24     }
25 }
26
27 // Definition for a Graph
28 struct Graph {
29     first: Link,
30     last: Link,
31 }
32
33 impl Graph {
34     fn new() -> Self {
35         Self { first: None, last: None }
36     }
37
38     fn is_empty(&self) -> bool {
39         self.first.is_none()
40     }
41
42     fn get_first(&self) -> Link {
43         self.first.clone()
44     }
45
46     // Print node
47     fn print_node(&self) {
48         let mut curr = self.first.clone();
49         while let Some(val) = curr {
50             print!("[{}]", &val.borrow().data);
51             curr = val.borrow().next.clone();

```

```

52     }
53
54     print!("\n");
55 }
56
57 // Insert node, RefCell is used to borrow_mut to modify
58 fn insert(&mut self, data: usize) {
59     let node = Rc::new(RefCell::new(Node::new(data)));
60
61     if self.is_empty() {
62         self.first = Some(node.clone());
63         self.last = Some(node);
64     } else {
65         self.last.as_mut()
66             .unwrap()
67             .borrow_mut()
68             .next = Some(node.clone());
69         self.last = Some(node);
70     }
71 }
72 }

```

The code above shows the implementation of the graph. The following code uses this graph to implement the basic breadth-first search algorithm. The `build_graph` function constructs the graph and encapsulates it into a tuple, which is then saved to a vector. The second value of the tuple indicates whether the node has been visited, with 0 indicating not visited and 1 indicating visited.

```

1 // bfs.rs
2
3 // Build the graph based on data
4 fn build_graph(data: [[usize;2];20]) -> Vec<(Graph, usize)> {
5     let mut graphs: Vec<(Graph, usize)> = Vec::new();
6     for _ in 0..9 { graphs.push((Graph::new(), 0)); }
7     for i in 1..9 {
8         for j in 0..data.len() {
9             if data[j][0] == i {
10                 graphs[i].0.insert(data[j][1]);
11             }
12         }
13         print!("[{i}]->");
14         graphs[i].0.print_node();
15     }
16     graphs
17 }
18
19 fn bfs(graph: Vec<(Graph, usize)>) {
20     let mut gp = graph;
21     let mut nodes = Vec::new();
22     gp[1].1 = 1;
23     let mut curr = gp[1].0.get_first().clone();
24
25     // Print the graph
26     print!("[1]->");

```

```

27     while let Some(val) = curr {
28         nodes.push(val.borrow().data);
29         curr = val.borrow().next.clone();
30     }
31
32     // Print the breadth-first graph
33     loop {
34         if 0 == nodes.len() {
35             break;
36         } else {
37             // Pop the first node from nodes, mimicking a queue
38             let data = nodes.remove(0);
39             // If the node has not been visited,
40             // add it to nodes and change its visit status to 1
41             if 0 == gp[data].1 {
42                 gp[data].1 = 1;
43                 // Print the value of the current node
44                 print!("{data}->");
45                 // Add nodes connected to current node to nodes
46                 let mut curr = gp[data].0.get_first().clone();
47                 while let Some(val) = curr {
48                     nodes.push(val.borrow().data);
49                     curr = val.borrow().next.clone();
50                 }
51             }
52         }
53     }
54     println!();
55 }
56
57 fn main() {
58     let data = [
59         [1,2],[2,1],[1,3],[3,1],[2,4],[4,2],[2,5],
60         [5,2],[3,6],[6,3],[3,7],[7,3],[4,5],[5,4],
61         [6,7],[7,6],[5,8],[8,5],[6,8],[8,6]
62     ];
63     let gp = build_graph(data);
64     bfs(gp);
65 }

```

The output of the code, which prints the adjacent nodes of each vertex and all vertices, is shown below.

```

[1]->[2][3]
[2]->[1][4][5]
[3]->[1][6][7]
[4]->[2][5]
[5]->[2][4][8]
[6]->[3][7][8]
[7]->[3][6]
[8]->[5][6]
1->2->3->4->5->6->7->8->

```

This algorithm uses `Vec` as a queue to search for the shortest transformation path in a word ladder. In addition to printing the values of each node's connections, it also prints all nodes in the order they are searched. To better represent node colors, an enum representing colors needs to be defined and node colors need to be updated accordingly. A distance value needs to be added to the node to represent the distance from the starting point in order to calculate the shortest distance.

```

1 // word_ladder.rs
2
3 // Enum for colors used to determine
4 // if a node has been searched
5 #[derive(Clone, Debug, PartialEq)]
6 enum Color {
7     White, // White: not yet explored
8     Gray,  // Gray: currently being explored
9     Black, // Black: exploration complete
10 }
11
12 // Definition for a Vertex
13 #[derive(Debug, Clone)]
14 struct Vertex<T> {
15     color: Color,
16     distance: u32, // Minimum distance from the starting node
17                   // i.e. minimum number of transformations
18     key: T,
19     neighbors: Vec<(T, u32)>, // store all adjacent vertices
20 }
21
22 impl<T: Clone + PartialEq> Vertex<T> {
23     fn new(key: T) -> Self {
24         Self {
25             color: Color::White,
26             distance: 0,
27             key: key,
28             neighbors: Vec::new(),
29         }
30     }
31
32     fn add_neighbor(&mut self, nbr: T, wt: u32) {
33         self.neighbors.push((nbr, wt));
34     }
35
36     // Get adjacent nodes
37     fn get_neighbors(&self) -> Vec<&T> {
38         let mut neighbors = Vec::new();
39
40         for (nbr, _wt) in self.neighbors.iter() {
41             neighbors.push(nbr);
42         }
43
44         neighbors
45     }
46 }

```

The code above, providing a color and node, is more complex than the basic BFS algorithm. To add nodes to the queue, the Queue implementation from Chapter 4 is introduced. The graph definition for solving the word ladder problem includes the number of nodes, edges, and a hashMap for storing node values and their structs.

```

1  // word_ladder.rs
2
3  use std::collections::HashMap;
4  use std::hash::Hash;
5
6  // Definition of Graph
7  #[derive(Debug, Clone)]
8  struct Graph<T> {
9      vertnums: u32,
10     edgenums: u32,
11     vertices: HashMap<T, Vertex<T>>,
12 }
13
14 impl<T: Hash + Eq + PartialEq + Clone> Graph<T> {
15     fn new() -> Self {
16         Self {
17             vertnums: 0,
18             edgenums: 0,
19             vertices: HashMap::<T, Vertex<T>>::new(),
20         }
21     }
22
23     fn contains(&self, key: &T) -> bool {
24         for (nbr, _vertex) in self.vertices.iter() {
25             if nbr == key { return true; }
26         }
27
28         false
29     }
30
31     // add a vertice
32     fn add_vertex(&mut self, key: &T) -> Option<Vertex<T>> {
33         let vertex = Vertex::new(key.clone());
34         self.vertnums += 1;
35         self.vertices.insert(key.clone(), vertex)
36     }
37
38     // add an edge
39     fn add_edge(&mut self, from: &T, to: &T, wt: u32) {
40         // Add node if it doesn't exist
41         if !self.contains(from) {
42             let _fvert = self.add_vertex(from);
43         }
44         if !self.contains(to) {
45             let _tvert = self.add_vertex(to);
46         }
47     }

```

```

48         self.edgenums += 1;
49         self.vertices
50             .get_mut(from)
51             .unwrap()
52             .add_neighbor(to.clone(), wt);
53     }
54 }

```

With this graph, a word ladder graph can be constructed by arranging the words according to their patterns.

```

1  // word_ladder.rs
2
3  // Construct the graph based on words and patterns
4  fn build_word_graph(words: Vec<&str>) -> Graph<String> {
5      let mut hmap: HashMap<String,Vec<String>> = HashMap::new();
6
7      // Build a word-pattern hashMap
8      for word in words {
9          for i in 0..word.len() {
10             let pattn = word[..i].to_string()
11                 + "_"
12                 + &word[i + 1..];
13             if hmap.contains_key(&pattn) {
14                 hmap.get_mut(&pattn)
15                     .unwrap()
16                     .push(word.to_string());
17             } else {
18                 hmap.insert(pattn, vec![word.to_string()]);
19             }
20         }
21     }
22
23     // Double-ended graph with a distance of 1
24     let mut word_graph = Graph::new();
25     for word in hmap.keys() {
26         for w1 in &hmap[word] {
27             for w2 in &hmap[word] {
28                 if w1 != w2 {
29                     word_graph.add_edge(w1, w2, 1);
30                 }
31             }
32         }
33     }
34
35     word_graph
36 }

```

The BFS algorithm is then used to search for the shortest path. Note that although three colors are defined, only white nodes will be added to the queue, so gray nodes can be set to black or left unset. This BFS function constructs the breadth-first search tree, and the search queue is emptied when the algorithm finds the result "sage".


```

1 // word_ladder.rs
2
3 // Word ladder graph - breadth-first search
4 fn word_ladder(g: &mut Graph<String>,
5               start: Vertex<String>,
6               end: Vertex<String>,
7               len: usize) -> u32
8 {
9     // Check if the starting point exists
10    if !g.vertices.contains_key(&start.key) {
11        return 0;
12    }
13
14    if !g.vertices.contains_key(&end.key) {
15        return 0;
16    }
17
18    // Prepare the queue and add the starting point
19    let mut vertex_queue = Queue::new(len);
20    let _r = vertex_queue.enqueue(start);
21
22    while vertex_queue.len() > 0 {
23        // Dequeue the node
24        let curr = vertex_queue.dequeue().unwrap();
25
26        for nbr in curr.get_neighbors() {
27            // Clone to avoid conflicts with data in the graph
28            // Vertices in the Graph are wrapped with RefCell
29            // and do not need to be cloned
30            let mut nbv = g.vertices.get(nbr).unwrap().clone();
31
32            if end.key != nbv.key {
33                // Only white nodes can be added to the queue;
34                // other colors have already been processed
35                if Color::White == nbv.color {
36                    // Update the node's color and distance and
37                    // add it to the queue
38                    nbv.color = Color::Gray;
39                    nbv.distance = curr.distance + 1;
40
41                    // The color and distance of the node
42                    // in the graph also need to be updated
43                    g.vertices.get_mut(nbr)
44                        .unwrap()
45                        .color = Color::Gray;
46                    g.vertices.get_mut(nbr)
47                        .unwrap()
48                        .distance = curr.distance + 1;
49
50                    // Add white nodes to the queue
51                    let _r = vertex_queue.enqueue(nbv);

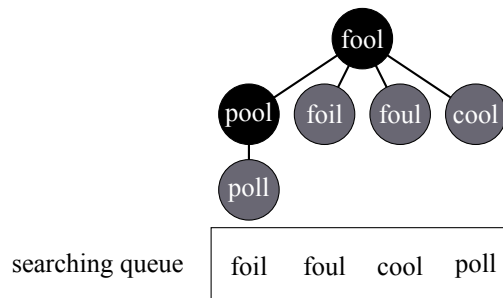
```

```

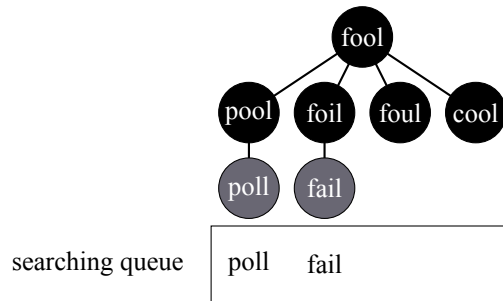
52         }
53     } else {
54         // Since the neighbor of curr contains end,
55         // one more transformation is enough
56         return curr.distance + 1;
57     }
58 }
59 }
60
61 0
62 }
63
64 fn main() {
65     let words = [
66         "FOOL", "COOL", "POOL", "FOUL", "FOIL", "FAIL", "FALL",
67         "POLL", "PALL", "POLE", "PALE", "SALE", "PAGE", "SAGE",
68     ];
69
70     let len = words.len();
71     let mut g = build_word_graph(words);
72
73     // The starting node is added to the queue and is
74     // being explored, so its color changes to gray
75     g.vertices.get_mut("FOOL").unwrap().color = Color::Gray;
76
77     // Retrieve the first and last points
78     let start = g.vertices.get("FOOL").unwrap().clone();
79     let end = g.vertices.get("SAGE").unwrap().clone();
80
81     // Calculate the minimum number of transformations,
82     // which is the distance.
83     let distance = word_ladder(&mut g, start, end, len);
84     println!("the shortest distance: {distance}");
85     // the shortest distance: 6
86 }

```

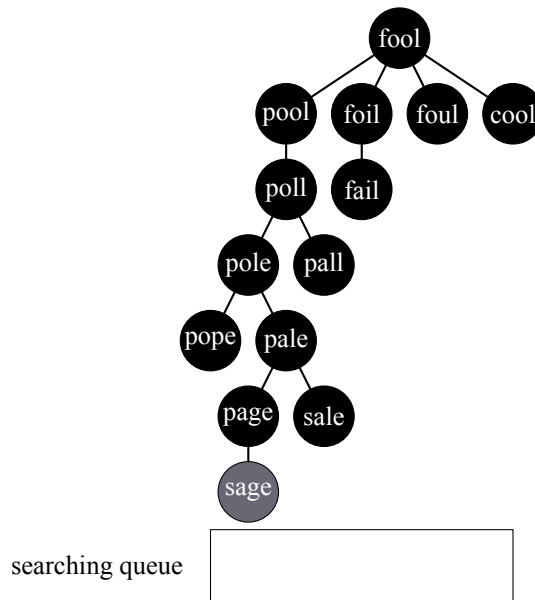
During the search process, the breadth-first search tree is constructed, and the node coloring during the search process is shown in the figure. At the beginning, all nodes adjacent to "fool", including "pool", "foil", "foul", and "cool", are taken, and these nodes are added to the queue for searching.



When bfs checks the node "cool", it finds that it is gray, indicating that there is a shorter path to "cool". When checking "pool", a new node "poll" is added.



The next vertex on the queue is "foil", and the only new node that can be added to the tree is "fail". When bfs continues to process the queue, the next two nodes do not add any new points to the queue.



Finally, the shortest path is the number of levels from "fool" to "sage", which is 6 levels(shortest path is 6). Readers can test this code with different words.

9.6.2 Analysis of BFS

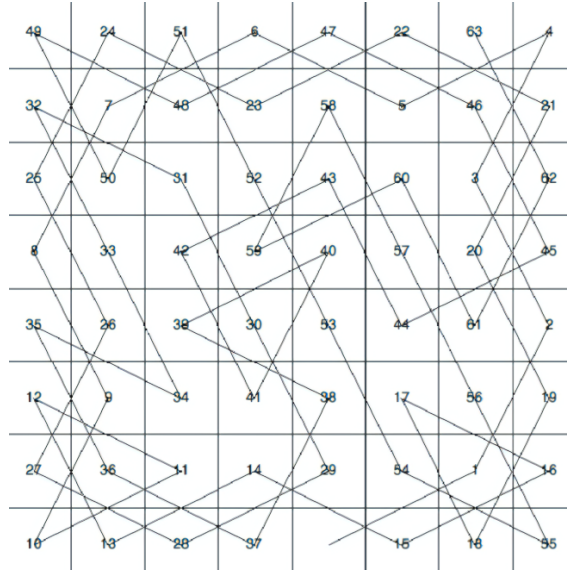
The time complexity of breadth-first search (BFS) depends on the number of vertices and edges in the graph. To create a graph in BFS, we need to process at least V vertices, resulting in a time complexity of $O(V)$. During the search, BFS processes one edge at a time, resulting in a time complexity of $O(E)$. Thus, the total time complexity of BFS is $O(V + E)$.

If all vertices are connected to the starting node, the queue needs to store at least the number of graph nodes, leading to a space complexity of $O(V)$.

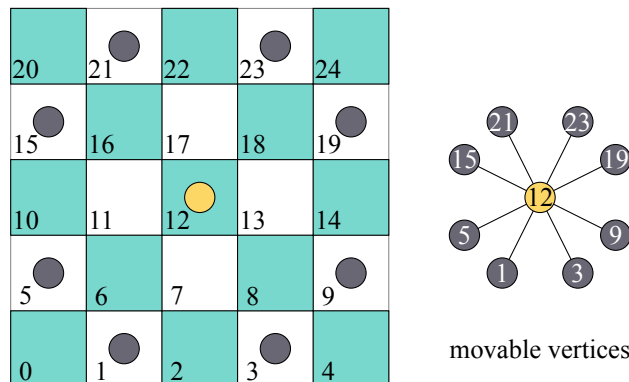
Interestingly, a graph can also be seen as a polyhedron split open, and its vertex-edge-face count must satisfy Euler's formula $V - E + F = 2$. By rearranging this formula, we get $V + E = 2E - F + 2$, which implies that the complexity of BFS is $O(V + E) = O(2E - F + 2)$. In the case of the word ladder graph(Figure 9.5.1), the vertex, edge, and face counts are 18, 14, and 6 respectively, and the formula is verified by substituting these values. Here, what I want to say is that the BFS algorithm for graphs seems to be extended to the geometric field which is a big inspiration for me.

9.6.3 The Knight's Tour Problem

The Knight's tour is a classic puzzle where a knight chess piece moves on a chessboard to visit every square exactly once, forming a tour. This problem has fascinated many people including players, mathematicians, and computer scientists for many years, and there are various algorithms to solve it. The graph search is the most straightforward solution, which requires two steps: representing the knight's moves and finding a path of length $rows \times columns - 1$, where the starting square is not counted. An 8×8 chessboard has a maximum of 1.305×10^{35} possible tours.



To represent the Knight's tour problem as a graph, each square on the chessboard is represented as a node, and each legal move of the knight is represented as an edge in the graph. The knight is represented as a yellow square in the figure below.



To construct a complete graph of size $n \times n$, the knight's eight possible moves must be considered. By traversing the entire graph, a move list can be created for each position, all of which can be converted into edges in the graph.

```
1 // knight_tour.rs
2
3 use std::collections::HashMap;
4 use std::hash::Hash;
```

```

5 use std::fmt::Display;
6
7 // bread-width
8 const BDSIZE: u32 = 8;
9
10 // Color enum used to determine if a node has been searched
11 #[derive(Debug, Clone, PartialEq)]
12 enum Color {
13     White, // White: not yet explored
14     Gray,  // Gray: currently being explored
15 }
16
17 // Definition of vertice
18 #[derive(Debug, Clone)]
19 struct Vertex<T> {
20     key: T,
21     color: Color,
22     neighbors: Vec<T>,
23 }
24
25 impl<T: PartialEq + Clone> Vertex<T> {
26     fn new(key: T) -> Self {
27         Self {
28             key: key,
29             color: Color::White,
30             neighbors: Vec::new(),
31         }
32     }
33
34     fn add_neighbor(&mut self, nbr: T) {
35         self.neighbors.push(nbr);
36     }
37
38     fn get_neighbors(&self) -> Vec<&T> {
39         let mut neighbors = Vec::new();
40         for nbr in self.neighbors.iter() {
41             neighbors.push(nbr);
42         }
43         neighbors
44     }
45 }

```

Once all the movable edges of each node have been found, the Knight's tour graph can be constructed.

```

1 // knight_tour.rs
2
3 // Knight's tour graph definition
4 #[derive(Debug, Clone)]
5 struct Graph<T> {
6     vertnums: u32,
7     edgenums: u32,
8     vertices: HashMap<T, Vertex<T>>,
9 }

```

```

10
11 impl<T: Eq + PartialEq + Clone + Hash> Graph<T> {
12     fn new() -> Self {
13         Self {
14             vertnums: 0,
15             edgenums: 0,
16             vertices: HashMap::<T, Vertex<T>>::new(),
17         }
18     }
19
20     fn add_vertex(&mut self, key: &T) -> Option<Vertex<T>> {
21         let vertex = Vertex::new(key.clone());
22         self.vertnums += 1;
23         self.vertices.insert(key.clone(), vertex)
24     }
25
26     fn add_edge(&mut self, src: &T, des: &T) {
27         if !self.vertices.contains_key(src) {
28             let _fv = self.add_vertex(src);
29         }
30         if !self.vertices.contains_key(des) {
31             let _tv = self.add_vertex(des);
32         }
33
34         self.edgenums += 1;
35         self.vertices.get_mut(src)
36             .unwrap()
37             .add_neighbor(des.clone());
38     }
39 }
40
41 // Destination coordinates that can be moved to
42 fn legal_moves(x: u32, y: u32, bdsz: u32) -> Vec<(u32, u32)>
43 {
44     // Knight moves like a horse, which is a "L" shape move
45     // Horizontal and vertical coordinates of the horse will
46     // increase or decrease accordingly in eight directions
47     let move_offsets = [
48         (-1, 2), (1, 2),
49         (-2, 1), (2, 1),
50         (-2, -1), (2, -1),
51         (-1, -2), (1, -2),
52     ];
53
54     // A closure function to check if the new coordinate
55     // is valid (within the chessboard range)
56     let legal_pos = |a: i32, b: i32| { a >= 0 && a < b };
57
58     let mut legal_positions = Vec::new();
59     for (x_offset, y_offset) in move_offsets.iter() {
60         let new_x = x as i32 + x_offset;
61         let new_y = y as i32 + y_offset;

```

```

61
62     // Check the coordinate and add it to
63     // the set of movable points
64     if legal_pos(new_x, bdsz as i32)
65         && legal_pos(new_y, bdsz as i32) {
66         legal_positions.push((new_x as u32, new_y as u32));
67     }
68 }
69
70 // Return the set of movable points
71 legal_positions
72 }
73
74 // Build the graph of movable paths
75 fn build_knight_graph(bdsz: u32) -> Graph<u32> {
76     // A closure function to calculate the point value
77     // Point range is [0, 63]
78     let calc_point = |row: u32, col: u32, size: u32| {
79         (row % size) * size + col
80     };
81
82     // Set edges between points
83     let mut knight_graph = Graph::new();
84     for row in 0..bdsz {
85         for col in 0..bdsz {
86             let dests = legal_moves(row, col, bdsz);
87             for des in dests {
88                 let src_p = calc_point(row, col, bdsz);
89                 let des_p = calc_point(des.0, des.1, bdsz);
90                 knight_graph.add_edge(&src_p, &des_p);
91             }
92         }
93     }
94
95     knight_graph
96 }

```

The search algorithm used to solve the Knight's tour problem is called depth-first search. While breadth-first search explores vertices on the same level as widely as possible, depth-first search explores multiple levels of the tree as deeply as possible. Various strategies can be employed using depth-first search to solve the problem. The first strategy prohibits nodes from being visited multiple times, while the second strategy allows nodes to be visited multiple times during tree construction. When the depth-first search algorithm encounters a dead (i.e., a point with no movable points) end, it backtracks to the previous deepest point and continues the exploration.

```

1 // depth: length of the path traveled,
2 // curr: current node, path: saves visited points
3 fn knight_tour<T>(
4     kg: &mut Graph<T>,
5     curr: Vertex<T>,
6     path: &mut Vec<String>,
7     depth: u32) -> bool
8     where T: Eq + PartialEq + Clone + Hash + Display

```

```

9  {
10     // Add the string value of the current node to path
11     path.push(curr.key.to_string());
12
13     let mut done = false;
14     if depth < BDSIZE * BDSIZE - 1 {
15         let mut i = 0;
16         let nbrs = curr.get_neighbors();
17
18         // Knight travels between neighboring points
19         while i < nbrs.len() && !done {
20             // Avoid multiple mutable references
21             let nbr = kg.vertices.get(nbrs[i]).unwrap().clone();
22
23             if Color::White == nbr.color {
24                 // Update the corresponding point color to gray
25                 kg.vertices.get_mut(nbrs[i])
26                     .unwrap()
27                     .color = Color::Gray;
28                 // Search for the next suitable point
29                 done = knight_tour(kg, nbr, path, depth + 1);
30                 if !done {
31                     // If not found, remove the current point
32                     // from path and restore the color of the
33                     // corresponding point to white
34                     let _rm = path.pop();
35                     kg.vertices.get_mut(nbrs[i])
36                         .unwrap()
37                         .color = Color::White;
38                 }
39             }
40
41             // Explore the next neighboring point
42             i += 1;
43         }
44     } else {
45         done = true;
46     }
47
48     done
49 }
50
51 fn main() {
52     // Build the knight's tour graph
53     let mut kg: Graph<u32> = build_knight_graph(BDSIZE);
54
55     // Choose a starting point and update the color
56     // of the corresponding point in the graph
57     let point = 0;
58     kg.vertices.get_mut(&point).unwrap().color = Color::Gray;
59     let start = kg.vertices.get(&point).unwrap().clone();
60

```



```

61 // Start the knight's tour path saves all visited points
62 let mut path = Vec::new();
63 let succeeded = knight_tour(&mut kg, start, &mut path, 0);
64
65 // Format and output the result.
66 if succeeded {
67     for row in 0..BDSIZE {
68         let row_s = ((row % BDSIZE) * BDSIZE) as usize;
69         let row_e = row_s + BDSIZE as usize;
70         let row_str = path[row_s..row_e].join("\t");
71         println!("{row_str}");
72     }
73 }
74 }

```

The knight's tour pathes can obtained using the `knight_tour` algorithm.

0	10	4	14	31	46	63	53
47	30	15	5	22	7	13	23
6	21	38	55	45	39	54	37
20	3	18	12	29	44	61	51
36	19	2	8	25	35	41	56
50	60	43	28	11	1	16	26
9	24	34	40	57	42	59	49
32	17	27	33	48	58	52	62

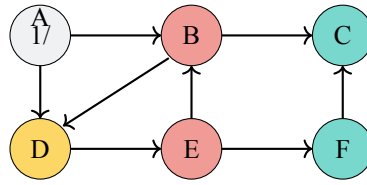
The knight's tour problem can be solved using the recursive version of depth-first search algorithm called the `knight_tour` algorithm. This algorithm searches for a path of length 64, which is the number of squares on the chessboard. The algorithm explores the graph by moving to the next vertex until it reaches a dead end or exhausts the entire graph, and then backtracks. To track the visited vertices, depth-first search uses colors where unvisited vertices are white and visited vertices are gray.

Backtracking occurs when the `knight_tour` function returns false, indicating that a dead end has been reached. Unlike breadth-first search, which uses a queue to track the next vertex to be visited, depth-first search is recursive and has an implicit stack that helps the algorithm backtrack.

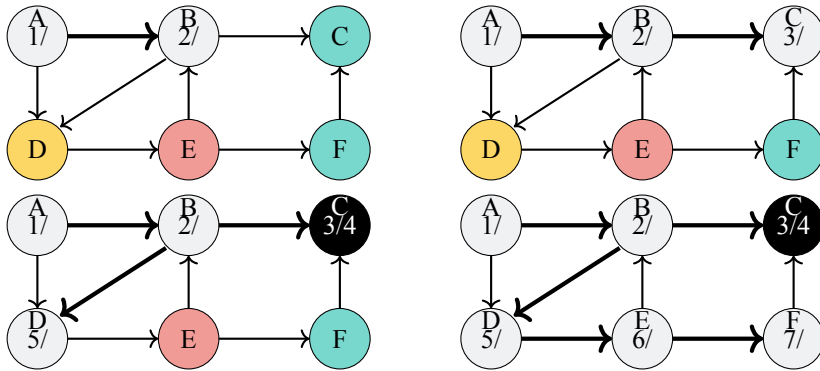
9.7 Depth First Search(DFS)

The Knight's Tour problem is a specific case of the depth-first search algorithm, which aims to create the deepest possible tree. Depth-first search is generally used to search graphs with multiple branches, with the goal of searching as deeply as possible, connecting as many nodes as possible, and branching when necessary. In some cases, depth-first search may create multiple trees, which are called a depth-first forest. Like breadth-first search, depth-first search uses predecessor links to construct trees. To better understand depth-first search, the vertex class can be expanded with two variables indicating the start and end times of the node's search, where the start time also indicates the number of steps before encountering the vertex, and the end time is the number of steps before the vertex is colored black.

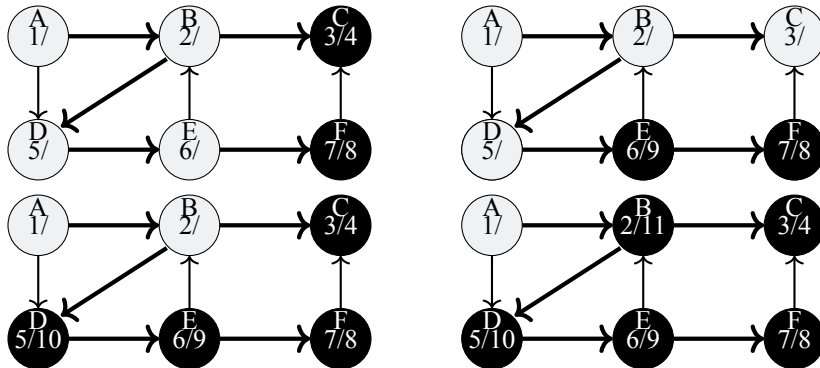
The following figure shows a small graph and the depth-first search algorithm applied to it. The thick lines indicate the edges checked, with the node on the other end of the edge already added to the depth-first tree. The search starts from vertex A, and since all vertices are colored at the start of the search, the algorithm randomly starts visiting vertex A. When visiting a vertex, the first step is to set its color to gray, indicating that the vertex is being explored, and to set the discovery time to 1. Since vertex A has two adjacent vertices (B and D), each vertex needs to be visited, and we visit adjacent vertices in alphabetical order.



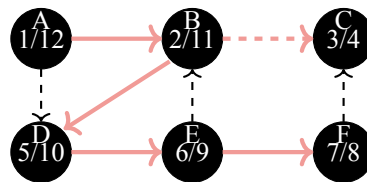
Next, we visit vertex B, which is adjacent to vertices C and D. Vertex C leads us to the end of a branch, so it can be colored black, and the end time set to 4. Now, return to vertex B and continue explore D, which leads us to vertex E. Vertex E has two adjacent vertices, B and F. Since B is already colored black, we can skip it and visit Vertex F.



Vertex F has only one adjacent vertex, C, already colored black, the algorithm has reached the end of the branch. Therefore, the algorithm must backtrack and continue to search until it encounters a gray vertex or exits.



Eventually, the algorithm backtracks to original vertex and exits the search, obtaining the visited path: $A \rightarrow B \rightarrow D \rightarrow E \rightarrow F$.



9.7.1 Implementing a DFS

From the previous discussion and illustration, we can present a basic implementation of the depth-first search algorithm that does not include color markings but implements essential exploration functionality.

```

1 // dfs.rs
2 use std::rc::Rc;
3 use std::cell::RefCell;
4
5 // Node link
6 type Link = Option<Rc<RefCell<Node>>>;
7
8 struct Node {
9     data: usize,
10    next: Link,
11 }
12
13 impl Node {
14     fn new(data: usize) -> Self {
15         Self {
16             data: data,
17             next: None
18         }
19     }
20 }
21
22 // Definition of Graph
23 struct Graph {
24     first: Link,
25     last: Link,
26 }
27
28 impl Graph {
29     fn new() -> Self {
30         Self { first: None, last: None }
31     }
32
33     fn is_empty(&self) -> bool {
34         self.first.is_none()
35     }
36
37     fn get_first(&self) -> Link {
38         self.first.clone()
39     }
40
41     fn print_node(&self) {
42         let mut curr = self.first.clone();
43         while let Some(val) = curr {
44             print!("[{}]", &val.borrow().data);
45             curr = val.borrow().next.clone();
46         }
47         print!("\n");

```

```

48     }
49
50     // insert data
51     fn insert(&mut self, data: usize) {
52         let node = Rc::new(RefCell::new(Node::new(data)));
53         if self.is_empty() {
54             self.first = Some(node.clone());
55             self.last = Some(node);
56         } else {
57             self.last.as_mut()
58                 .unwrap()
59                 .borrow_mut()
60                 .next = Some(node.clone());
61             self.last = Some(node);
62         }
63     }
64 }
65
66 // build a graph
67 fn build_graph(data: [[usize;2];20]) -> Vec<(Graph, usize)> {
68     let mut graphsVec<(Graph, usize)> = Vec::new();
69     for _ in 0..9 {
70         graphs.push((Graph::new(), 0));
71     }
72
73     for i in 1..9 {
74         for j in 0..data.len() {
75             if data[j][0] == i {
76                 graphs[i].0.insert(data[j][1]);
77             }
78         }
79
80         print!("[{i}]->");
81         graphs[i].0.print_node();
82     }
83
84     graphs
85 }
86
87 fn dfs(graph: Vec<(Graph, usize)>) {
88     let mut gp = graph;
89     let mut nodes: Vec<usize> = Vec::new();
90     let mut temp: Vec<usize> = Vec::new();
91
92     gp[1].1 = 1;
93     let mut curr = gp[1].0.get_first().clone();
94
95     // print graph
96     print!("[1]->");
97     while let Some(val) = curr {
98         nodes.insert(0, val.borrow().data);
99         curr = val.borrow().next.clone();

```

```

100     }
101
102     // print dfs graph
103     loop{
104         if 0 == nodes.len() {
105             break;
106         }else{
107             let data = nodes.pop().unwrap();
108             if 0 == gp[data].1 { // Unvisited
109                 // Change state to visited
110                 gp[data].1 = 1;
111                 print!("{data}->");
112
113                 // Add node to temp and perform depth-first
114                 // search on it
115                 let mut curr = gp[data].0.get_first().clone();
116                 while let Some(val) = curr {
117                     temp.push(val.borrow().data);
118                     curr = val.borrow().next.clone();
119                 }
120
121                 while !temp.is_empty(){
122                     nodes.push(temp.pop().unwrap());
123                 }
124             }
125         }
126     }
127
128     println!("");
129 }
130
131 fn main() {
132     let data = [
133         [1,2],[2,1],[1,3],[3,1],[2,4],[4,2],[2,5],
134         [5,2],[3,6],[6,3],[3,7],[7,3],[4,5],[5,4],
135         [6,7],[7,6],[5,8],[8,5],[6,8],[8,6]
136     ];
137     let gp = build_graph(data);
138     dfs(gp);
139 }

```

Here is the output that displays the adjacent nodes of each vertex and the deepest path.

```

[1]->[2][3]
[2]->[1][4][5]
[3]->[1][6][7]
[4]->[2][5]
[5]->[2][4][8]
[6]->[3][7][8]
[7]->[3][6]
[8]->[5][6]
1->2->4->5->8->6->3->7->

```

9.7.2 Analysis of DFS

The time complexity of depth-first search (DFS) can be calculated from both the vertices and edges perspectives. In `build_graph`, each vertex needs to be processed once to construct the graph, resulting in a complexity of $O(V)$. During the search, DFS starts from a vertex and explores all adjacent vertices as deeply as possible, processing each edge with a time complexity of $O(E)$. Therefore, the total time complexity of DFS is $O(V + E)$. If all vertices form a chain during the DFS, the deepest search will include all nodes, resulting in a space complexity of $O(V)$.

It is important to note that trees are a special case of graphs. Therefore, depth-first search and breadth-first search algorithms for graphs can also be used for searching trees. The breadth-first search we learned in the chapter on trees is actually a level-order traversal. DFS and BFS are the foundation for many other important graph algorithms, making them essential for searching graphs. In the following sections, we will use BFS and DFS of the graph to solve practical problems to deepen our understanding.

9.7.3 Topological Sorting

Real-world problems can often be modeled as graphs and solved using graph algorithms. For example, consider the problem of designing a four-year course plan for a major, which involves ensuring correct course dependencies and a feasible sequence of courses. This problem can be complex, but with the graph data structure, we can use graph algorithms to solve it.

Suppose we have four courses with their dependencies shown in the graph, a feasible course study order is [functions, derivatives, systems of equations, calculus, linear algebra, probability theory, convolutional neural networks]. To determine the order of the courses, we can use an algorithm to sort the course dependencies, which is known as topological sorting.

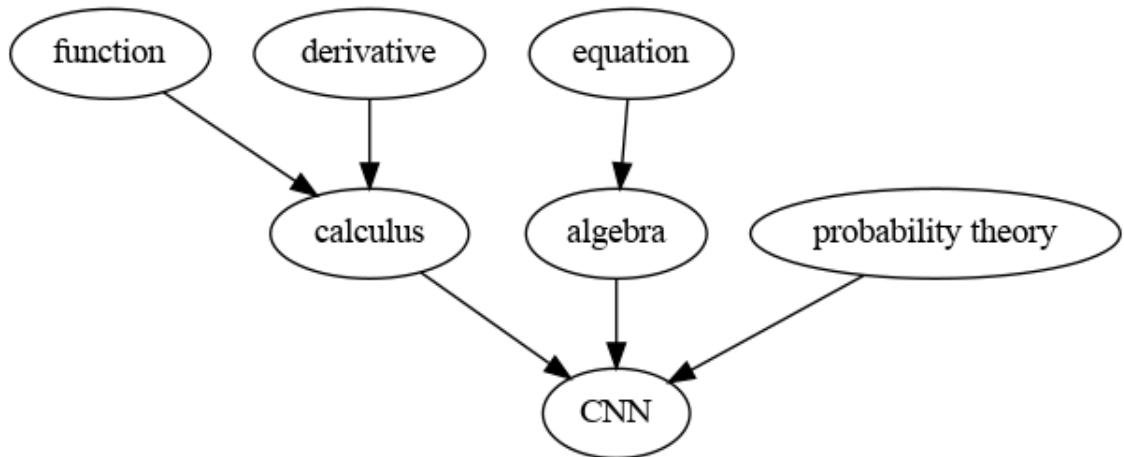


Figure 9.5: Courses Topological Relationship

Topological sorting is a variant of depth-first search that produces a linear ordering of all vertices in a directed acyclic graph. This resulting order can be used to indicate event priorities, set project schedules, generate priority graphs for database queries, and so on. The basic ideas of topological sorting are as follows:

- Call `dfs()` on the graph `g` using depth-first search to find suitable vertices.
- Store the vertices in a stack, with the last one at the bottom.
- Return the data in the stack as the result of the topological sort.

Following these steps, we can transform a graph into a linear relationship, such as the course plan mentioned above, which can be converted into a topological sequence.

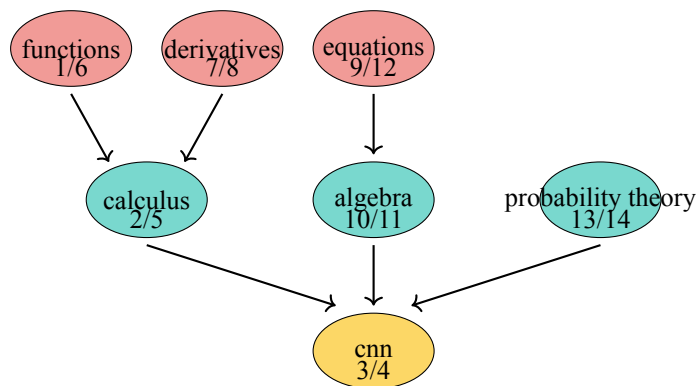


Figure 9.6: Course scheduling

The final topological sort result is presented below. Courses with the same color can be taken in any order, while courses with different colors must be taken in the order specified by the topological sort.

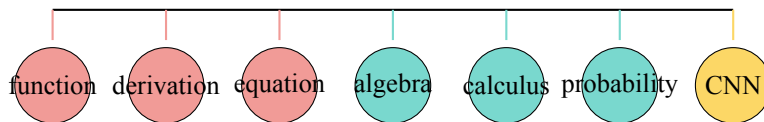


Figure 9.7: Topological sequence of course scheduling

To implement the graph algorithm for course scheduling, we first define an enumeration of colors to represent the state of node exploration.

```

1 // course_topological_sort.rs
2 use std::collections::HashMap;
3 use std::hash::Hash;
4 use std::fmt::Display;
5
6 // Color enum used to determine if a node has been searched
7 #[derive(Debug, Clone, PartialEq)]
8 enum Color {
9     White, // White: not yet explored
10    Gray,  // Gray: currently being explored
11    Black, // Black: exploration complete
12 }
  
```

Then, we define the node and graph as we have done many times before.

```

1 // course_topological_sort.rs
2
3 // course vertex definition
4 #[derive(Debug, Clone)]
5 struct Vertex<T> {
6     key: T,
7     color: Color,
8     neighbors: Vec<T>,
9 }
10
  
```

```

11 impl<T: PartialEq + Clone> Vertex<T> {
12     fn new(key: T) -> Self {
13         Self {
14             key: key,
15             color: Color::White,
16             neighbors: Vec::new(),
17         }
18     }
19
20     fn add_neighbor(&mut self, nbr: T) {
21         self.neighbors.push(nbr);
22     }
23 }
24
25 // coueses relation graph definition
26 #[derive(Debug, Clone)]
27 struct Graph<T> {
28     vertnums: u32,
29     edgenums: u32,
30     vertices: HashMap<T, Vertex<T>>, // all vertex
31     edges: HashMap<T, Vec<T>>, // all edges
32 }
33
34 impl<T: Eq + PartialEq + Clone + Hash> Graph<T> {
35     fn new() -> Self {
36         Self {
37             vertnums: 0,
38             edgenums: 0,
39             vertices: HashMap::<T, Vertex<T>>::new(),
40             edges: HashMap::<T, Vec<T>>::new(),
41         }
42     }
43
44     fn add_vertex(&mut self, key: &T) -> Option<Vertex<T>> {
45         let vertex = Vertex::new(key.clone());
46         self.vertnums += 1;
47         self.vertices.insert(key.clone(), vertex)
48     }
49
50     fn add_edge(&mut self, src: &T, des: &T) {
51         if !self.vertices.contains_key(src) {
52             let _sv = self.add_vertex(src);
53         }
54
55         if !self.vertices.contains_key(des) {
56             let _dv = self.add_vertex(des);
57         }
58
59         // add vertex
60         self.edgenums += 1;
61         self.vertices.get_mut(src)
62             .unwrap()

```



```

63             .add_neighbor(des.clone());
64
65         // add edge
66         if !self.edges.contains_key(src) {
67             let _eg = self.edges
68                 .insert(src.clone(), Vec::new());
69         }
70
71         self.edges.get_mut(src).unwrap().push(des.clone());
72     }
73 }

```

With the graph in place, we can start building the course dependency graph. When exploring all course nodes in the graph, we use the color attribute to indicate whether the node has been visited or not. The schedule variable stores the result of the course topological sorting. To prevent scheduling errors, such as circular dependencies, we add a "has_circle" variable here to control the search process. When encountering a cycle, the program exits, indicating that the input data is incorrect.

```

1  // course_topological_sort.rs
2
3  // Build the course dependency graph.
4  fn build_course_graph<T>(pre_requisites:Vec<Vec<T>>)->Graph<T>
5      where T: Eq + PartialEq + Clone + Hash
6  {
7      // Create edge relationships for dependent courses.
8      let mut course_graph = Graph::new();
9      for v in pre_requisites.iter() {
10         let prev = v.first().unwrap();
11         let last = v.last().unwrap();
12         course_graph.add_edge(prev, last);
13     }
14
15     course_graph
16 }
17
18 // Course scheduling.
19 fn course_scheduling<T>(
20     cg: &mut Graph<T>,
21     course: Vertex<T>,
22     schedule: &mut Vec<String>,
23     mut has_circle: bool)
24 where T: Eq + PartialEq + Clone + Hash + Display
25 {
26     // Clone to prevent conflicts with mutable references.
27     let edges = cg.edges.clone();
28
29     // Explore dependent courses.
30     let dependencies = edges.get(&course.key);
31
32     if !dependencies.is_none() {
33         for dep in dependencies.unwrap().iter() {
34             let course = cg.vertices.get(dep)
35                 .unwrap().clone();

```

```

36         if Color::White == course.color {
37             cg.vertices.get_mut(dep)
38                 .unwrap()
39                 .color = Color::Gray;
40
41             course_scheduling(cg, course,
42                             schedule, has_circle);
43
44             // Exit when encountering a cycle.
45             if has_circle {
46                 return;
47             }
48         } else if Color::Gray == course.color {
49             has_circle = true;
50             return;
51         }
52     }
53 }
54
55 // Change node color and add to schedule.
56 cg.vertices.get_mut(&course.key)
57     .unwrap()
58     .color = Color::Black;
59 schedule.push(course.key.to_string());
60 }
61
62 fn find_topological_order<T>(
63     course_num: usize,
64     pre_requisites: Vec<Vec<T>>)
65 where T: Eq + PartialEq + Clone + Hash + Display
66 {
67     // Build the course dependency graph.
68     let mut cg = build_course_graph(pre_requisites);
69
70     // Get all course nodes into "courses".
71     let vertices = cg.vertices.clone();
72     let mut courses = Vec::new();
73     for key in vertices.keys() {
74         courses.push(key);
75     }
76
77     // Save feasible course schedules.
78     let mut schedule = Vec::new();
79
80     // Determine if there is a cycle.
81     let has_circle = false;
82
83     // Perform topological sorting of courses.
84     for i in 0..course_num {
85         let course = cg.vertices.get(&courses[i])
86             .unwrap()
87             .clone();

```

```

88
89     // Only explore courses that are not explored
90     // and do not have cycles.
91     if !has_circle && Color::White == course.color {
92         // Change the color of the course node to indicate
93         // that it is currently being explored.
94         cg.vertices.get_mut(&courses[i])
95             .unwrap()
96             .color = Color::Gray;
97
98         course_scheduling(&mut cg, course,
99             &mut schedule, has_circle);
100     }
101 }
102
103 if !has_circle {
104     println!("{:?}", schedule);
105 }
106 }
107
108 fn main() {
109     let course_num = 7;
110
111     // Build course dependency relationships.
112     let mut pre_requisites = Vec::<Vec<&str>>::new();
113     pre_requisites.push(vec!["calculus", "function"]);
114     pre_requisites.push(vec!["calculus", "derivation"]);
115     pre_requisites.push(vec!["linear algebra", "equation"]);
116     pre_requisites.push(vec!["CNN", "calculus"]);
117     pre_requisites.push(vec!["CNN", "probability theory"]);
118     pre_requisites.push(vec!["CNN", "linear algebra"]);
119
120     // Find the topological sorting result, which is a
121     // reasonable course learning order.
122     find_topological_order(course_num, pre_requisites);
123 }

```

Here is a feasible course learning order.

```

[
    "function",
    "derivation",
    "calculus",
    "equation",
    "linear algebra",
    "probability theory",
    "CNN",
]

```

In addition to course selection, cooking can also be abstracted as a topological sort. For example, consider making pancakes, as shown in the figure below. The recipe is simple: 1 egg, 1 cup of pancake mix, 1 tablespoon of oil, and 3/4 cup of milk. To make pancakes, you must first turn on the stove and heat the pan. Then mix all the ingredients together with a spoon and pour the mixture into the pan. When

bubbles begin to form, flip the pancake over until both sides are golden brown. Before eating, you can add sauce.

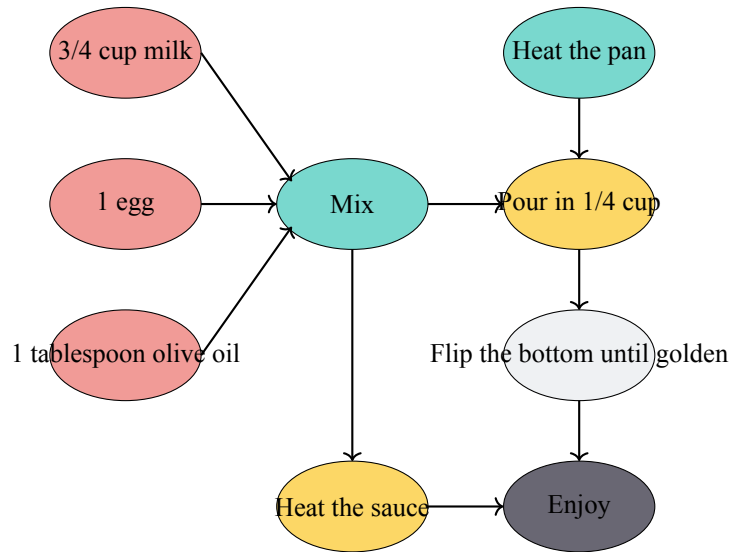


Figure 9.8: Pancake Cooking

By using the topological sort algorithm, the above figure can be simplified into a standard cooking step diagram. In this diagram, steps with the same color can be taken in any order. This step diagram can help kitchen novices make pancakes successfully. However, making them taste good is another matter altogether.

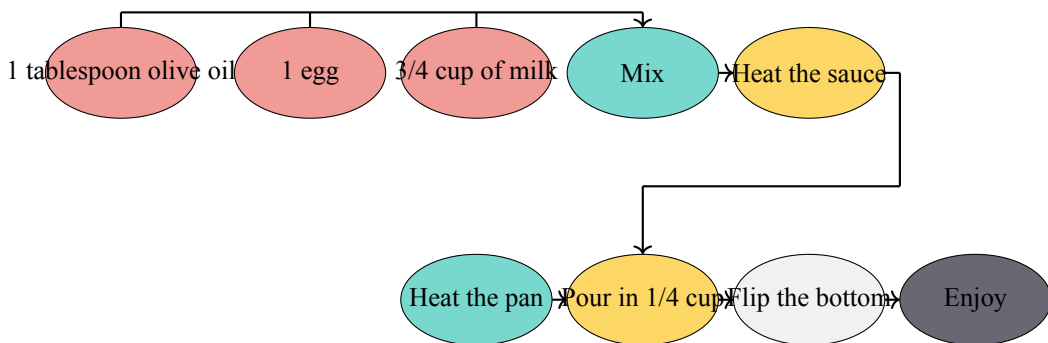


Figure 9.9: Pancake cooking process topological relationship

Both cooking and course selection can be abstracted as process arrangements, and therefore, the same code can theoretically be used to handle them. The implementation of the cooking process topological sort algorithm is the same as the course planning topological sort algorithm. Therefore, we will not include most of the code here, but it can be found in the source code files of this book.

```

1 // cooking_topological_sort.rs
2
3 // Implementation omitted as it is identical to
4 // course_topological_sort
5 fn main() {
6     let operation_num = 9;
  
```

```

7
8 // constructe a cooking process dependency relationship
9 let mut pre_requisites = Vec::<Vec<&str>>::new();
10 pre_requisites.push(vec!["Mix", "3/4 cup of milk"]);
11 pre_requisites.push(vec!["Mix", "1 egg"]);
12 pre_requisites.push(vec!["Mix", "1 tablespoon olive oil"]);
13 pre_requisites.push(vec!["Pour in 1/4 cup", "Mix"]);
14 pre_requisites.push(vec!["Pour in 1/4 cup",
15                          "Heat the pan"]);
16 pre_requisites.push(vec!["Flip the bottom until golden",
17                          "Pour in 1/4 cup"]);
18 pre_requisites.push(vec!["Enjoy",
19                          "Flip the bottom until golden"]);
20 pre_requisites.push(vec!["Enjoy", "Heat the sauce"]);
21
22 // Find a feasible topological result
23 // which is a reasonable cooking order
24 find_topological_order(operation_num, pre_requisites);
25 }

```

Here are two feasible cooking sequences.

```

[
  "3/4 cup of milk",
  "1 egg",
  "1 tablespoon olive oil",
  "Mix",
  "Heat the pan",
  "Pour in 1/4 cup",
  "Flip the bottom until golden",
  "Heat the sauce",
  "Enjoy",
]
[
  "Heat the pan",
  "3/4 cup of milk",
  "1 egg",
  "1 tablespoon olive oil",
  "Mix",
  "Pour in 1/4 cup",
  "Flip the bottom until golden",
  "Heat the sauce",
  "Enjoy",
]

```

9.8 Strongly Connected Components

Large graphs, like the links between web pages, are commonly represented as graphs, and search engines such as Google uses massive directed graphs to store the links on the World Wide Web. In this type of graph, pages are vertices, and hyperlinks are edges connecting them. Figure 9.10 depicts the web pages linked to Google's main site, highlighting that the entire internet can be seen as a graph.

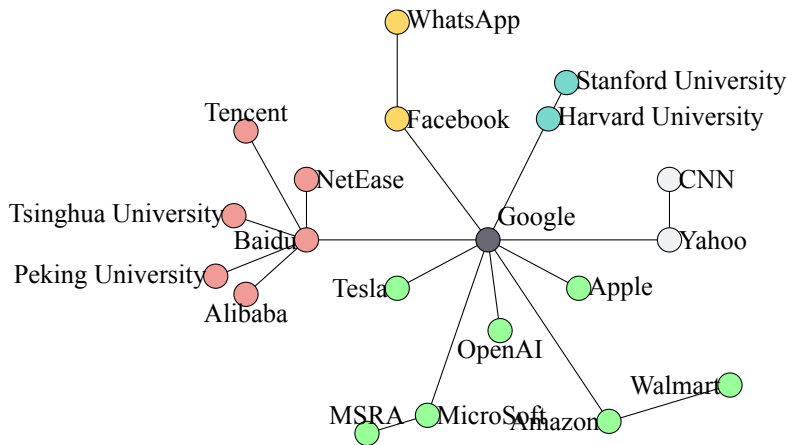


Figure 9.10: Internet connection

One significant feature of these graphs is that certain vertices have an exceptionally high number of edges. For example, Google is connected to nearly every website globally, while websites like Baidu and Tencent have numerous links within China. However, many vertices have very few or just one edge, forming clusters of interconnected nodes. These clusters are known as connected components, and if a connected component can be reached by any node in it in a finite number of steps, it is called a strongly connected component. A directed graph is strongly connected if and only if every two vertices in it are mutually reachable. Strongly connected graphs are similar to nested loops and must have loops. A directed graph may have multiple strongly connected components, and each of these components is called a strongly connected component. Figure(9.11) illustrates an example of strongly connected components in a directed graph.

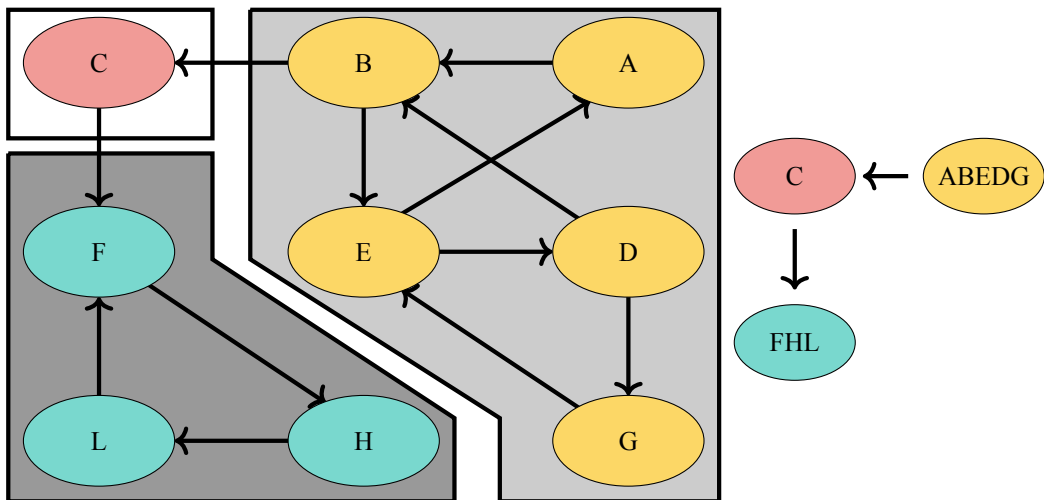
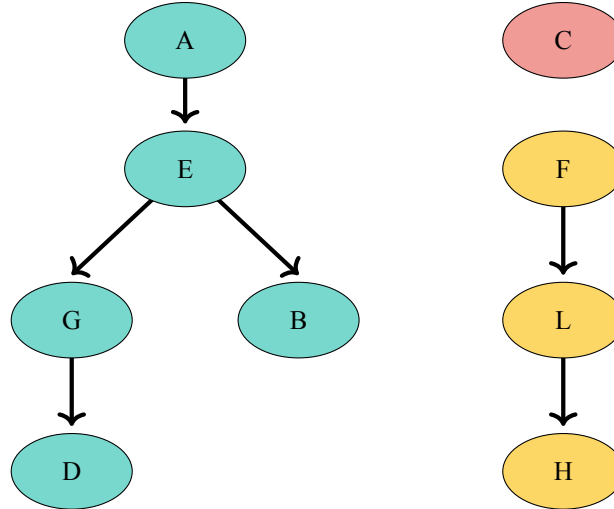


Figure 9.11: Strongly connected components and its simplification

The strongly connected components $C \in G$, where vertices v and w are connected, can be represented as a point in a simplified graph (shown in the left tricolored areas). To identify strongly connected components, the Tarjan algorithm, which is based on depth-first search, is commonly used.

Applying the strongly connected component algorithm to a connected graph produces three trees that correspond to three connected components, as shown in the figure below. The independent area C

consists of only itself as a node, while F, L, and H form one connected component, and A, E, G, D, and B form another connected component. The three trees obtained by the strongly connected component algorithm represent the three connected components. By shifting the problem from the node level to the connected component level, the algorithm simplifies the problem and facilitates subsequent analysis and processing.



9.8.1 The BFS Strongly Connected Components Algorithm

The provincial partition problem is an example of a strongly connected component. In China, cities are categorized into provincial administrative regions. To solve this problem, a large amount of information about cities and their interconnections is provided, and an algorithm is used to divide cities into partitions and determine the number of provinces to which these cities belong.

Assuming there are n cities, some of which are connected to each other and some are not, Chengdu and Yibin both belong to Sichuan Province, so the relationship between them can be represented as ["Chengdu", "Yibin"] and ["Yibin", "Chengdu"]. Another example may be ["Guangzhou", "Shenzhen"], ["Guangzhou", "Dongguan"]... Here, cities are abstracted as points in the graph, and relationships are seen as edges. The number of provinces can be determined by finding the number of strongly connected components, which can be achieved by the breadth-first search algorithm.

For each city node in the graph, its color is initially white, and when visited, it is colored gray. The breadth-first search starts from that city until all the cities in the same connected component have been visited (colored gray), indicating a province. Traversing all connected components of cities, all provinces can be found. In the case of the provincial partition problem, the answer should be 2, the number of provinces.

Below is an implementation of the strongly connected component algorithm using breadth-first search. It contains only the most basic functions for point and graph implementation, and unused ones are removed. The edges variable in the graph definition saves all edge relationships for exploring edge relationships.

```

1 // find_province_num_bfs.rs
2
3 use std::collections::HashMap;
4 use std::hash::Hash;
5
6 // Enum for colors
7 #[derive(Debug, Clone, PartialEq)]

```

```

8  enum Color {
9      White, // White: not yet explored
10     Gray,  // Gray: currently being explored
11 }
12
13 // Definition of cities
14 #[derive(Debug, Clone)]
15 struct Vertex<T> {
16     key: T,
17     color: Color,
18     neighbors: Vec<T>,
19 }
20 impl<T: PartialEq + Clone> Vertex<T> {
21     fn new(key: T) -> Self {
22         Self {
23             key: key,
24             color: Color::White,
25             neighbors: Vec::new(),
26         }
27     }
28
29     fn add_neighbor(&mut self, nbr: T) {
30         self.neighbors.push(nbr);
31     }
32
33     fn get_neighbors(&self) -> Vec<&T> {
34         let mut neighbors = Vec::new();
35         for nbr in self.neighbors.iter() {
36             neighbors.push(nbr);
37         }
38         neighbors
39     }
40 }
41
42 // Definition of province
43 #[derive(Debug, Clone)]
44 struct Graph<T> {
45     vertnums: u32,
46     edgenums: u32,
47     vertices: HashMap<T, Vertex<T>>,
48     edges: HashMap<T, Vec<T>>,
49 }
50
51 impl<T: Eq + PartialEq + Clone + Hash> Graph<T> {
52     fn new() -> Self {
53         Self {
54             vertnums: 0,
55             edgenums: 0,
56             vertices: HashMap::<T, Vertex<T>>::new(),
57             edges: HashMap::<T, Vec<T>>::new(),
58         }
59     }

```



```

60
61     fn add_vertex(&mut self, key: &T) -> Option<Vertex<T>> {
62         let vertex = Vertex::new(key.clone());
63         self.vertnums += 1;
64         self.vertices.insert(key.clone(), vertex)
65     }
66
67     fn add_edge(&mut self, src: &T, des: &T) {
68         if !self.vertices.contains_key(src) {
69             let _fv = self.add_vertex(src);
70         }
71         if !self.vertices.contains_key(des) {
72             let _tv = self.add_vertex(des);
73         }
74
75         // add a vertex (city)
76         self.edgenums += 1;
77         self.vertices.get_mut(src)
78             .unwrap()
79             .add_neighbor(des.clone());
80         // add an edge
81         if !self.edges.contains_key(src) {
82             let _ = self.edges.insert(src.clone(), Vec::new());
83         }
84         self.edges.get_mut(src).unwrap().push(des.clone());
85     }
86 }

```

Given the graph and node definitions, we can build a city connection graph. We continually change the color of each city to gray until a group of cities is fully explored, at which point we find a strongly connected component, i.e., a province.

```

1 // find_province_num_bfs.rs
2
3 / Build the city connection graph
4 fn build_city_graph<T>(connected: Vec<Vec<T>>) -> Graph<T>
5     where T: Eq + PartialEq + Clone + Hash {
6     // Set edges between nodes with relationships
7     let mut city_graph = Graph::new();
8     for v in connected.iter() {
9         let src = v.first().unwrap();
10        let des = v.last().unwrap();
11        city_graph.add_edge(src, des);
12    }
13
14    city_graph
15 }
16
17 fn find_province_num_bfs<T>(connected: Vec<Vec<T>>) -> u32
18     where T: Eq + PartialEq + Clone + Hash {
19     let mut cg = build_city_graph(connected);
20
21     // Get keys of all main city nodes

```

```

22     let mut cities = Vec::new();
23     for key in cg.edges.keys() { cities.push(key.clone()); }
24
25     // Process each strongly connected component
26     let mut province_num = 0;
27     let mut q = Queue::new(cities.len());
28     for ct in &cities {
29         let city = cg.vertices.get(ct).unwrap().clone();
30         if Color::White == city.color {
31             // Change the current node's color and enqueue it
32             cg.vertices.get_mut(ct)
33                 .unwrap()
34                 .color = Color::Gray;
35             q.enqueue(city);
36             // Process a strongly connected component
37             // of a province
38             while !q.is_empty() {
39                 // Get a node and its neighbors
40                 let q_city = q.dequeue().unwrap();
41                 let nbrs = q_city.get_neighbors();
42                 // Process each neighbor
43                 for nbr in nbrs {
44                     let nbrc = cg.vertices.get(nbr)
45                         .unwrap()
46                         .clone();
47                     if Color::White == nbrc.color {
48                         // Enqueue the unexplored nodes
49                         cg.vertices.get_mut(nbr)
50                             .unwrap()
51                             .color = Color::Gray;
52                         q.enqueue(nbrc);
53                     }
54                 }
55             }
56             // Finish processing a strongly connected component
57             // of a province
58             province_num += 1;
59         }
60     }
61
62     province_num
63 }

```

```

1 fn main() {
2     // Build the city dependency relationship
3     let mut connected = Vec:::<Vec<&str>>::new();
4     connected.push(vec!["Chengdu", "Zigong"]);
5     connected.push(vec!["Chengdu", "Mianyang"]);
6     connected.push(vec!["Chengdu", "Deyang"]);
7     connected.push(vec!["Chengdu", "Luzhou"]);
8     connected.push(vec!["Chengdu", "Neijiang"]);
9     connected.push(vec!["Chengdu", "Leshan"]);

```

```

10     connected.push(vec!["Chengdu", "Yibin"]);
11     connected.push(vec!["Zigong", "Chengdu"]);
12
13     connected.push(vec!["Guangzhou", "Shenzhen"]);
14     connected.push(vec!["Guangzhou", "Dongguan"]);
15     connected.push(vec!["Guangzhou", "Zhuhai"]);
16     connected.push(vec!["Guangzhou", "Zhongshan"]);
17     connected.push(vec!["Guangzhou", "Shantou"]);
18     connected.push(vec!["Guangzhou", "Foshan"]);
19     connected.push(vec!["Guangzhou", "Zhanjiang"]);
20     connected.push(vec!["Shenzhen", "Guangzhou"]);
21
22     connected.push(vec!["Wuhan", "Jingzhou"]);
23     connected.push(vec!["Wuhan", "Yichang"]);
24     connected.push(vec!["Wuhan", "Xiangyang"]);
25     connected.push(vec!["Wuhan", "Jingmen"]);
26     connected.push(vec!["Wuhan", "Xiaogan"]);
27     connected.push(vec!["Wuhan", "Huanggang"]);
28     connected.push(vec!["Jingzhou", "Wuhan"]);
29
30     // Find all strongly connected components, there are three
31     // provinces: Sichuan, Guangdong, and Hubei
32     let province_num = find_province_num_bfs(connected);
33     println!("province number: {province_num}");
34     // province number: 3
35 }

```

The time complexity of this algorithm is $O(n^2)$ because we need to process n city nodes, and each city node may have a relationship with all remaining cities in the graph (when there is only one province). Breadth-first search uses a queue that can hold up to all n city nodes, so the space complexity is $O(n)$.

9.8.2 The DFS Strongly Connected Components Algorithm

The province partition problem can also be solved using the depth-first search algorithm. The idea of depth-first search is to traverse all city nodes, and for each city, if it has not been visited yet (colored white), color it gray, and then start a depth-first search from that city. By using the depth-first search algorithm, we can obtain the cities directly connected to that city. These cities and the city itself belong to the same connected component. We can continue performing depth-first search on these cities until all cities in the same connected component have been visited (colored gray), indicating a province has been found. Traversing all cities allows us to obtain the total number of connected components, which is the total number of provinces.

```

1 // find_province_num_dfs.rs
2 use std::collections::HashMap;
3 use std::hash::Hash;
4
5 // Implementation details omitted for brevity,
6 // please add them yourself
7
8 // Search for neighboring nodes of current node 'city'
9 fn search_city<T>(cg: &mut Graph<T>, city: Vertex<T>)
10     where T: Eq + PartialEq + Clone + Hash
11 {

```

```

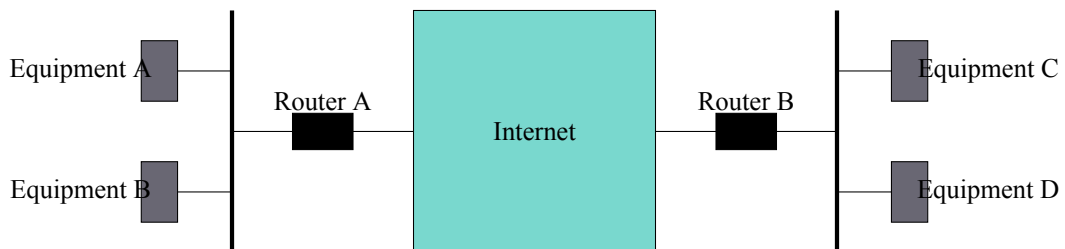
12     // Traverse the neighboring nodes of current node 'city'
13     for ct in city.get_neighbors() {
14         let city = cg.vertices.get(ct).unwrap().clone();
15         if Color::White == city.color {
16             // Change the color of current node
17             cg.vertices.get_mut(ct)
18                 .unwrap()
19                 .color = Color::Gray;
20             // Continue to search for neighboring nodes
21             search_city(cg, city);
22         }
23     }
24 }
25
26 fn find_province_num_dfs<T>(city_connected: Vec<Vec<T>>) -> u32
27     where T: Eq + PartialEq + Clone + Hash
28 {
29     let mut cg = build_city_graph(city_connected);
30     let mut cities = Vec::new();
31
32     // Get the key of each main city node
33     for key in cg.edges.keys() { cities.push(key.clone()); }
34
35     let mut province_num = 0;
36     // Process connected components of a province one by one
37     for ct in &cities {
38         let city = cg.vertices.get(ct).unwrap().clone();
39         if Color::White == city.color {
40             // Change the color of current node
41             cg.vertices.get_mut(ct)
42                 .unwrap()
43                 .color = Color::Gray;
44             // Search for neighboring nodes of current node
45             search_city(&mut cg, city);
46             // Finish processing a province connected component
47             province_num += 1;
48         }
49     }
50
51     province_num
52 }
53
54 fn main() {
55     // Build city dependency relationship
56
57     let mut city_connected = Vec::<Vec<String>>::new();
58
59     // Same as in BFS, omitted for brevity
60     let province_num = find_province_num_dfs(city_connected);
61     println!("province nummber: {province_num}");
62     // province nummber: 3
63 }

```

In terms of complexity, it is similar to BFS since it also needs to process n city nodes, and each city node may have a relationship with all the remaining cities in the graph (when there is only one province). Therefore, the time complexity is still $O(n^2)$. Depth-first search uses a stack that can hold up to n city nodes, so the space complexity is $O(n)$.

9.9 Shortest Path Problem

When accessing online content or remote computing resources, such as watching videos, sending emails, or logging into a lab computer from off-campus, network communication is required. Understanding how information is transmitted from one computer to another via the internet is a fundamental aspect of computer networks.



The diagram above illustrates the process of internet communication. When a user requests a webpage from a server using a browser, the request first travels through the local area network and then through a router to access the internet. The request then propagates through the internet until it reaches the local area network router where the server is located. The requested webpage is then sent back to the user's browser via the same router. The `tracert` command can be used to identify the path from a user's computer to a specific link. For instance, tracing the website `xxx.cn` may involve passing through 13 routers, with the first two being gateway routers of the user's network group.

```

1?: [LOCALHOST]                pmtu 1500
1:  _gateway                    4.523 ms
1:  _gateway                    3.495 ms
2:  10.253.0.22                 2.981 ms
3:  no response
4:  ???                         6.166 ms
5:  202.115.254.237             558.609 ms
6:  no response
7:  no response
8:  101.4.117.54                48.822 ms asymm 16
9:  no response
10: 101.4.112.37                48.171 ms asymm 14
11: no response
12: 101.4.114.74                44.981 ms
13: 202.97.15.89                49.560 ms

```

Each router on the internet is connected to one or more routers. Thus, running `tracert` at different times of the day may yield different results since the connections between routers have a cost that varies depending on network traffic. The network connections can be viewed as a weighted graph, with connections being adjusted based on the network situation.

The goal is to find the path with the minimum total weight to transmit the message. This problem is similar to the word ladder problem discussed earlier since both involve finding the minimum value, but the weights in the word ladder problem are all the same.

9.9.1 Dijkstra's Algorithm

Predecessors have proposed various algorithms for studying the shortest path in network graphs, and among them, Dijkstra's algorithm is a good option. It is a greedy iterative algorithm that finds the shortest path from a specific starting node to all other nodes in the graph, similar to breadth-first search.

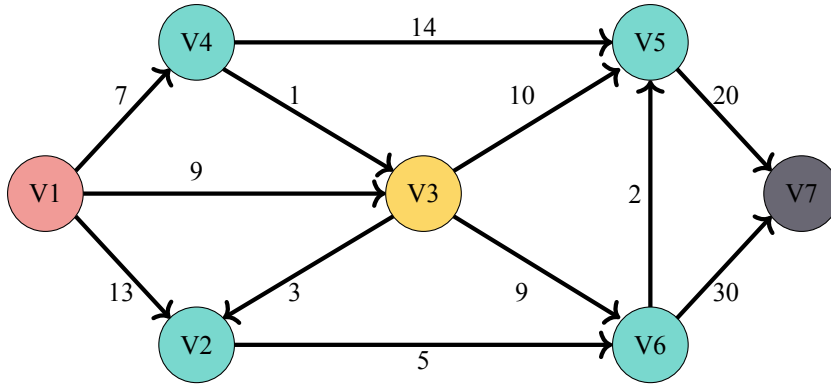


Figure 9.12: Dijkstra

In the figure above, we want to find the shortest path from V1 to V7. By visually exploring the graph, we can see that there are two shortest paths: [V1->V4->V3->V2->V6->V5->V7] and [V1->V4->V3->V5->V7], with a shortest distance of 38. If we were to calculate this using Dijkstra's algorithm, we would need to track and calculate various distances and then add them up.

To keep track of the total distance from the starting node to each target node, we will use the `dist` instance variable in the graph vertex. This variable contains the total weight of the path from the start to the target node. Dijkstra's algorithm iterates over each node in the graph, with the iteration order controlled by a priority queue, and the `dist` value is used to determine the order of objects in the queue. When a node is first created, `dist` is set to 0. Theoretically, it is possible to set `dist` to infinity, but in practice, setting it to 0 is feasible, or setting it to a value larger than any real distance, such as the distance that light travels in one second, which is approximately equal to the distance between the earth and the moon, because no two points on earth are so far apart.

9.9.2 Implementing the Dijkstra's Algorithm

Dijkstra's algorithm is an incremental approach to finding the shortest path step-by-step, using a priority queue to process vertices based on their priority value. The algorithm selects the closest unvisited vertex to the starting point from all vertices that haven't yet had their shortest path found, and continues the process. However, using this algorithm on the internet poses a challenge because it requires a complete graph representation of the entire network. This means that every router must have a map of all routers in the Internet, which is impractical.

```

1 // dijkstra.rs
2
3 use std::cmp::Ordering;
4 use std::collections::{BinaryHeap, HashMap, HashSet};
5
6 #[derive(Debug, Copy, Clone, PartialEq, Eq, Hash)]
7 struct Vertex<'a> {
8     name: &'a str,
9 }
  
```

```

10
11 impl<'a> Vertex<'a> {
12     fn new(name: &'a str) -> Vertex<'a> {
13         Vertex {
14             name
15         }
16     }
17 }
18
19 // Vertices visited
20 #[derive(Debug)]
21 struct Visited<V> {
22     vertex: V,
23     distance: usize,
24 }
25
26 // Add total ordering functionality to Visited
27 impl<V> Ord for Visited<V> {
28     fn cmp(&self, other: &Self) -> Ordering {
29         other.distance.cmp(&self.distance)
30     }
31 }
32 impl<V> PartialOrd for Visited<V> {
33     fn partial_cmp(&self, other: &Self) -> Option<Ordering> {
34         Some(self.cmp(other))
35     }
36 }
37
38 impl<V> Eq for Visited<V> {}
39 impl<V> PartialEq for Visited<V> {
40     fn eq(&self, other: &Self) -> bool {
41         self.distance.eq(&other.distance)
42     }
43 }
44
45 // Shortest path algorithm
46 fn dijkstra<'a>(
47     start: Vertex<'a>,
48     adj_list: &HashMap<Vertex<'a>,
49     Vec<(Vertex<'a>, usize)>>) -> HashMap<Vertex<'a>, usize>
50 {
51     let mut distances = HashMap::new();
52     let mut visited = HashSet::new(); // Visited vertices
53     let mut to_visit = BinaryHeap::new(); // Unvisited vertices
54
55     // Set the starting point and
56     // initial distances to all points
57     distances.insert(start, 0);
58     to_visit.push(Visited {
59         vertex: start,
60         distance: 0,
61     });

```

```

62
63     while let Some(Visited { vertex, distance }) =
64         to_visit.pop() {
65         // If visited, continue with the next point
66         if !visited.insert(vertex) { continue; }
67
68         if let Some(nbrs) = adj_list.get(&vertex) {
69             for (nbr, cost) in nbrs {
70                 let new_dist = distance + cost;
71                 let is_shorter =
72                     distances.get(&nbr)
73                         .map_or(true,
74                             |&curr| new_dist < curr);
75                 // If the distance is shorter, insert the new
76                 // distance and neighbor
77                 if is_shorter {
78                     distances.insert(*nbr, new_dist);
79                     to_visit.push(Visited {
80                         vertex: *nbr,
81                         distance: new_dist,
82                     });
83                 }
84             }
85         }
86     }
87
88     distances
89 }
90
91 fn main() {
92     let v1 = Vertex::new("V1");
93     let v2 = Vertex::new("V2");
94     let v3 = Vertex::new("V3");
95     let v4 = Vertex::new("V4");
96     let v5 = Vertex::new("V5");
97     let v6 = Vertex::new("V6");
98     let v7 = Vertex::new("V7");
99
100     let mut adj_list = HashMap::new();
101     adj_list.insert(v1, vec![(v4, 7), (v2, 13)]);
102     adj_list.insert(v2, vec![(v6, 5)]);
103     adj_list.insert(v3, vec![(v2, 3), (v6, 9), (v5, 10)]);
104     adj_list.insert(v4, vec![(v3, 1), (v5, 14)]);
105     adj_list.insert(v5, vec![(v7, 20)]);
106     adj_list.insert(v6, vec![(v5, 2), (v7, 30)]);
107
108     // Find the shortest path from V1 to any point
109     let distances = dijkstra(v1, &adj_list);
110     for (v, d) in &distances {
111         println!("{}", min distance: {d}", v1.name, v.name);
112     }
113 }

```


The following outputs are shortest path between V1 and any point(including itself).

```
V1-V5, min distance: 18
V1-V2, min distance: 11
V1-V6, min distance: 16
V1-V3, min distance: 8
V1-V7, min distance: 38
V1-V1, min distance: 0
V1-V4, min distance: 7
```

To solve this problem, other algorithms such as the Distance Vector Routing Protocol^[15] and Link State Routing Protocol^[16] are used for network information transmission. These algorithms allow routers to discover the network maps saved by other routers, which contain information about interconnected nodes. This approach allows for real-time updates of the network map content and is more efficient, greatly reducing network capacity requirements.

9.9.3 Analysis of Dijkstra's Algorithm

Let's analyze the time complexity of Dijkstra's algorithm. Firstly, building the priority queue takes $O(V)$ time. Once the queue is constructed, the while loop is executed once for each vertex. Since vertices are added at the beginning and removed only after that, calling pop in the loop takes $O(\log V)$ time. We can consider this loop and the calls to pop as $O(V \log V)$. The for loop iterates once for each edge in the graph, and calling decrease_key takes $O(E \log V)$ time. Therefore, the total time complexity is $O((V + E) \log V)$, and the space complexity is $O(V)$.

9.10 Summary

In this chapter, we have explored the abstract data type of graphs and their implementation. Graphs are widely used in various fields such as course scheduling, networks, transportation, computers, knowledge graphs, and databases. Graphs can be utilized to solve many problems, as long as we can transform the original problem into a graph representation. Graphs have numerous applications in the following areas:

- Strongly connected components can be used to simplify graphs.
- Depth-first search can be utilized to explore deep branches of a graph.
- Topological sorting is useful in clarifying complex graph connections.
- Dijkstra's algorithm can be employed to search for the shortest path in a weighted graph.
- Breadth-first search can be used to search for the shortest path in an unweighted graph.

Chapter 10

Practices

10.1 Objectives

- Utilize Rust data structures and algorithms to accomplish diverse practical projects.
- Comprehend and implement data structures and algorithms employed in practical projects.

10.2 Edit Distance

The edit distance measures the similarity between two sequences by calculating the minimum number of operations required to transform one word, W_1 , into another word, W_2 , through insertion, deletion, or substitution. These operations are commonly used in daily editing, which is why it is called edit distance. Notably, the number of edits performed by a single operation is counted as an edit distance.

There are two main types of edit distance: the Hamming distance and the Levenshtein distance. The Hamming distance only considers substitution operations, while the Levenshtein distance simultaneously takes into account insertion, deletion, and substitution operations. Readers may ponder if it is possible to develop an edit distance that involves only two of these operations.

10.2.1 The Hammig Distance

The Hamming distance is the number of differing symbols at the same positions between two sequences of equal length or the number of differing bits in a binary sequence. It measures the minimum number of substitution operations required to convert one sequence into another. For instance, in the figure below, converting "trust" to "rrost" only requires substituting two characters, resulting in a Hamming distance of 2.

t	r	u	s	t
r	r	o	s	t

Figure 10.1: Hamming distance

The Hamming distance is commonly used for error correction in coding. In Hamming codes^[17], the algorithm for computing the distance is the Hamming distance. To simplify the code, separate algorithms are implemented for calculating the Hamming distance of numbers and characters. Computing the Hamming distance of numbers is straightforward as bitwise operations can compare numbers for similarity and difference. The code for calculating the Hamming distance of numbers is shown below.

```

1 // hamming_distance.rs
2
3 fn hamming_distance1(source: u64, target: u64) -> u32 {
4     let mut count = 0;
5     let mut xor = source ^ target;
6     while xor != 0 {
7         count += xor & 1;
8         xor >>= 1;
9     }
10    count as u32
11 }
12
13 fn main() {
14     let source = 1;
15     let target = 2;
16     let distance = hamming_distance1(source, target);
17     println!("the hamming distance is {distance}");
18     // the hamming distance is 2
19
20     let source = 3;
21     let target = 4;
22     let distance = hamming_distance1(source, target);
23     println!("the hamming distance is {distance}");
24     // the hamming distance is 3
25 }

```

The XOR operation sets the same bits in "source" and "target" numbers to 0 and different bits to 1. If the result is nonzero, different bits exist. Thus, we calculate the different bits step by step from the last bit using XOR and AND operations with 1. After computing each bit, we remove it to compare the previous bits by adding a right shift operation. Note that the implementation requires manually counting the number of 1's in the binary system. However, Rust provides a `count_ones()` function for counting the number of 1's in its numbers, simplifying the above code to the following concise form.

```

1 // hamming_distance.rs
2
3 fn hamming_distance2(source: u64, target: u64) -> u32 {
4     (source ^ target).count_ones()
5 }

```

With this foundation, we now implement the character version of the Hamming distance.

```

1 // hamming_distance.rs
2
3 fn hamming_distance_str(source: &str, target: &str) -> u32 {
4     let mut count = 0;
5     let mut source = source.chars();
6     let mut target = target.chars();
7
8     // Compare two strings character by character
9     loop {
10         match (source.next(), target.next()) {
11             // Four situations may arise
12             (Some(cs), Some(ct)) if cs != ct => count += 1,

```

```

13             (Some(_), None) | (None, Some(_)) =>
14                 panic!("Must have the same length"),
15             (None, None) => break,
16             _ => continue,
17         }
18     }
19
20     count as u32
21 }
22
23 fn main() {
24     let source = "abce";
25     let target = "edcf";
26     let distance = hamming_distance_str(source, target);
27     println!("the hamming distance is {distance}");
28     // the hamming distance is 3
29 }

```

The character version of the Hamming distance algorithm takes two parameters, "source" and "target", and compares Unicode characters using the `chars` method. Unicode is preferred over ASCII because characters may include not only letters but also other characters, such as Chinese, Japanese, and Korean characters, with each character corresponding to multiple ASCII values. The `"if c1 != c2"` is an additional condition check outside of the pattern matching. It only enters the matching branch if both "source" and "target" have the next character, and the two characters are not equal. If either character is "None" and the other is "Some," it means the input strings have different lengths, and the function can return directly. If there are no more characters left, the comparison ends. In other cases, if the two characters are the same, the function continues to compare the next character. Since the Hamming distance requires calculating all characters, the time complexity is $O(n)$, and the space complexity is $O(1)$.

10.2.2 The Levenshtein Distance

The Levenshtein distance, also known as edit distance, is an algorithm used to measure the difference between two strings by representing the minimum number of edit operations required to transform one string into another, including insertion, deletion, and substitution. The concept of edit distance is easy to understand, and the operations involved are simple, making it useful for basic character correction. For instance, calculating the edit distance between the words "sitting" and "kitten" using the Levenshtein distance algorithm involves the following steps:

- sitting -> kitting, replace "s" with "k".
- kitting -> kitteng, replace "i" with "e".
- kitteng -> kitten, delete "g".

Since it took three operations, making the edit distance three. However, how can we prove that three is the minimum number of edit operations required? This is because there are only three types of operations that can be performed to transform one string to another: deletion, insertion, and substitution.

One way to transform an empty string into a string of length s is to insert s characters, making the edit distance between the two strings equal to the length of s . For example, transforming an empty string into "abc" requires inserting three characters, so the edit distance is 3. The figure (10.2.2) on next page illustrates the extreme cases required to transform "sitting" into "kitten". It also shows that the maximum edit distance is the length of the longer string. This can be expressed mathematically using the formula:

$$edi_{a,b}(i, j) = \max(i, j) \quad \text{if } \min(i, j) = 0 \quad (10.1)$$

Where i and j are the lengths of the two strings, $\min(i, j) = 0$ means that there is no common substring.

		s	i	t	t	i	n	g
	0	1	2	3	4	5	6	7
k	1							
i	2							
t	3							
t	4							
e	5							
n	6							

Figure 10.2: Edit distances (Converting empty string to a string)

Except for cases where the strings have no common substrings, the edit distance can be increased through three types of editing operations, each of which increases the edit distance by 1. Thus, we can calculate the edit distance obtained by each of the three editing operations and then take the minimum value.

$$edi_{a,b}(i, j) = \min \begin{cases} edi_{a,b}(i-1, j) + 1 \\ edi_{a,b}(i, j-1) + 1 \\ edi_{a,b}(i-1, j-1) + 1_{a \neq b} \end{cases} \quad (10.2)$$

The formula $edi_{a,b}(i-1, j) + 1$ means that one character must be deleted to transform string a into string b , increasing the edit distance by 1. The formula $edi_{a,b}(i, j-1) + 1$ means that one character must be inserted to transform string a into string b , also increasing the edit distance by 1. The formula $edi_{a,b}(i-1, j-1) + 1_{a \neq b}$ means that one character must be replaced to transform string a into string b , increasing the edit distance by 1 only if a and b are not equal. These functions are recursively defined, and their space complexity is $O(3^{m+n-1})$, where m and n are the lengths of the strings.

We have learned that dynamic programming is a good way to solve big problem by split it into several small ones. Therefore, dynamic programming is used to handle recursion in this case. The algorithm works by storing the edit distance in a matrix, which represents the state of the problem after various operations. The most basic case is when an empty string is transformed into a string of varying lengths, as illustrated in the figure (10.2.2) on next page.

The edit distance between character k and s is calculated next, which can be divided into three cases:

- The accumulated edit distance of 1 deletion above the red cell, plus a deletion operation, results in an edit distance of 2.
- The accumulated edit distance of 1 insertion to the left of the red cell, plus an insertion operation, results in an edit distance of 2.
- The accumulated edit distance of 0 substitutions along the diagonal of the red cell, plus a substitution operation, results in an edit distance of 1.

All the values being processed in this step are from the yellow region in the diagram. Starting from the top left corner, the edit distance in the red cell is computed by taking the minimum of the three values in the yellow region and adding 1. This yields the new edit distance.

		s	i	t	t	i	n	g
	0	1	2	3	4	5	6	7
k	1	1						
i	2							
t	3							
t	4							
e	5							
n	6							

Figure 10.3: State transition matrix

Based on the above description and diagram, we can develop the following algorithm to calculate the edit distance between two strings.

```

1 // edit_distance.rs
2
3 use std::cmp::min;
4
5 fn edit_distance(source: &str, target: &str) -> usize {
6     // Extreme case: transformation from empty string
7     // to a non-empty string
8     if source.is_empty() {
9         return target.len();
10    } else if target.is_empty() {
11        return source.len();
12    }
13
14    // Establish a matrix to store process values
15    let source_c = source.chars().count();
16    let target_c = target.chars().count();
17    let mut distance = vec![vec![0; target_c+1]; source_c+1];
18    (1..=source_c).for_each(|i| {
19        distance[i][0] = i;
20    });
21    (1..=target_c).for_each(|j| {
22        distance[0][j] = j;
23    })
24
25    // Save the minimum number of steps(insert, delete, modify)
26    for (i, cs) in source.chars().enumerate() {
27        for (j, ct) in target.chars().enumerate() {

```

```

28         let ins = distance[i+1][j] + 1;
29         let del = distance[i][j+1] + 1;
30         let sub = distance[i][j] + (cs != ct) as usize;
31         distance[i+1][j+1] = min(min(ins, del), sub);
32     }
33 }
34
35 // Return the value in the last row and last column
36 *distance.last().and_then(|d| d.last()).unwrap()
37 }

```

```

1 fn main() {
2     let source = "abce";
3     let target = "adcf";
4     let dist = edit_distance(source, target);
5     println!("distance between {source} and {target}: {dist}");
6     // distance between abce and adcf: 2
7
8     let source = "bdfc";
9     let target = "adcf";
10    let dist = edit_distance(source, target);
11    println!("distance between {source} and {target}: {dist}");
12    // distance between bdfc and adcf: 3
13 }

```

To select the three values that need to be calculated, we can gradually move the yellow area. Once we have these values, we take the minimum calculated result and fill it into the lower right corner of the current yellow area. This new value represents the edit distance for that particular section. The final calculation result can be seen in the figure below.

		s	i	t	t	i	n	g
	0	1	2	3	4	5	6	7
k	1	1	2	3	4	5	6	7
i	2	2	1	2	3	4	5	6
t	3	3	2	1	2	3	4	5
t	4	4	3	2	1	2	3	4
e	5	5	4	3	2	2	3	4
n	6	6	5	4	3	3	2	3

The entire edit distance state transition matrix can be used to calculate the value in the last row and last column, which represents the overall edit distance. It is important to note that the matrix is two-

dimensional and requires careful use of subscripts. One approach is to place each row of the matrix in an array, forming a large array with $m \times n$ values but smaller dimensions. However, this approach can result in a lot of intermediate values that waste memory. To optimize the algorithm, we can repeatedly use a single array to calculate and save values. This approach reduces the matrix to an array with a length of $m + 1$ or $n + 1$. The optimized code for the edit distance algorithm is shown below.

```

1 // edit_distance.rs
2
3 fn edit_distance2(source: &str, target: &str) -> usize {
4     if source.is_empty() {
5         return target.len();
6     } else if target.is_empty() {
7         return source.len();
8     }
9
10    // The "distances" variable stores the edit distances
11    // to various strings
12    let target_c = target.chars().count();
13    let mut distances = (0..=target_c).collect::<Vec<_>>();
14    for (i, cs) in source.chars().enumerate() {
15        let mut substt = i;
16        distances[0] = substt + 1;
17        // Combinations are continuously calculated
18        // to obtain the distances
19        for (j, ct) in target.chars().enumerate() {
20            let dist = min(
21                min(distances[j], distances[j+1]) + 1,
22                substt + (cs != ct) as usize);
23            substt = distances[j+1];
24            distances[j+1] = dist;
25        }
26    }
27
28    // The last distance value is the answer
29    distances.pop().unwrap()
30 }
31
32 fn main() {
33     let source = "abced";
34     let target = "adcf";
35     let dist = edit_distance2(source, target);
36     println!("distance between {source} and {target}: {dist}");
37     // distance between abced and adcf: 3
38 }

```

The optimized edit distance algorithm has a worst-case time complexity of $O(mn)$ and a worst-case space complexity that has been reduced from $O(mn)$ for the matrix to $O(\min(m, n))$ for the array. This represents a significant improvement.

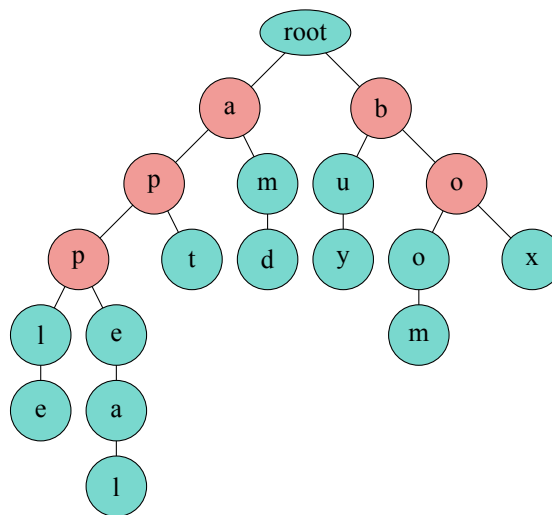
Microsoft Word software uses a spell check function that does not rely on edit distance. Instead, it uses a hash table to store commonly used words, allowing it to quickly look up each word as it is typed. If the word is not found in the hash table, an error is reported. Because hash tables are very fast, and several hundred thousand words only require a few megabytes of memory, this approach is highly efficient.

10.3 Trie

Trie, also known as a radix tree or prefix tree, is a tree data structure used to search for words or prefixes. It has a wide range of applications, including auto-completion, spell checking, typing prediction, and more.

While balanced trees and hash tables can also be used to search for words, they cannot quickly find all words with the same prefix or enumerate all stored words in lexicographical order. Hash tables can find a word in $O(1)$ time, but as the number of words grows, the hash table may become too large and increase the time complexity to $O(n)$ due to collisions. In contrast, Trie uses less space to store multiple words with the same prefix, and its time complexity is only $O(m)$, where m is the length of the word. The time complexity of searching for a word in a balanced tree is $O(m \log(n))$.

The structure of Trie is shown in the figure below. Storing words only requires processing 26 letters, and words with the same prefix share the same prefix, which saves storage space. For example, "apple" and "appeal" share "app," and "boom" and "box" share "bo."



To implement Trie, we first need to abstract the node, similar to the node in the above figure. The node stores the reference to its child nodes and the state of the current node. The state indicates whether this node is the end of a word ("end"), which can be used to determine whether the word ends when searching. In addition, the root node is the entrance to Trie and used to represent the entire Trie.

```

1 // trie.rs
2
3 // Definition of Trie
4 #[derive(Default, Debug)]
5 struct Trie {
6     root: Node,
7 }
8
9 // Definition of Trie node
10 #[derive(Default)]
11 struct Node {
12     end: bool,
13     children: [Option<Box<Node>>; 26], // List of letter nodes
14 }
15
16 impl Trie {

```

```

17     fn new() -> Self {
18         Self::default()
19     }
20
21     // Insert word
22     fn insert(&mut self, word: &str) {
23         let mut node = &mut self.root;
24
25         // Insert character by character
26         for c in word.as_bytes() {
27             let index = (c - b'a') as usize;
28             let next = &mut node.children[index];
29             node = next.get_or_insert_with(
30                 Box::<Node>::default);
31         }
32
33         node.end = true;
34     }
35
36     fn contains(&self, word: &str) -> bool {
37         self.word_node(word)
38             .map_or(false, |n| n.end)
39     }
40
41     // Check if there exists a word with a certain prefix
42     fn start_with(&self, prefix: &str) -> bool {
43         self.word_node(prefix)
44             .is_some()
45     }
46
47     // Prefix string (wps: word_prefix_string)
48     fn word_node(&self, wps: &str) -> Option<&Node> {
49         let mut node = &self.root;
50         for c in wps.as_bytes() {
51             let index = (c - b'a') as usize;
52             match &node.children[index] {
53                 None => return None,
54                 Some(next) => node = next.as_ref(),
55             }
56         }
57
58         Some(node)
59     }
60 }
61
62 fn main() {
63     let mut trie = Trie::new();
64     trie.insert("box");
65     trie.insert("insert");
66     trie.insert("apple");
67     trie.insert("appeal");
68

```

```

69     let res1 = trie.contains("apple");
70     let res2 = trie.contains("apples");
71     let res3 = trie.start_with("ins");
72     let res4 = trie.start_with("ina");
73     let res5 = trie.word_node("box");
74
75     println!("word 'apple' in Trie: {res1}");
76     println!("word 'apples' in Trie: {res2}");
77     println!("prefix 'ins' in Trie: {res3}");
78     println!("prefix 'ina' in Trie: {res4}");
79     println!("{:?}", res5);
80 }

```

The following are outputs after execution.

```

word 'apple' in Trie: true
word 'apples' in Trie: false
prefix 'ins' in Trie: true
prefix 'ina' in Trie: false
Some(Node { end: true, children: [None, ..., None] })

```

10.4 Filters

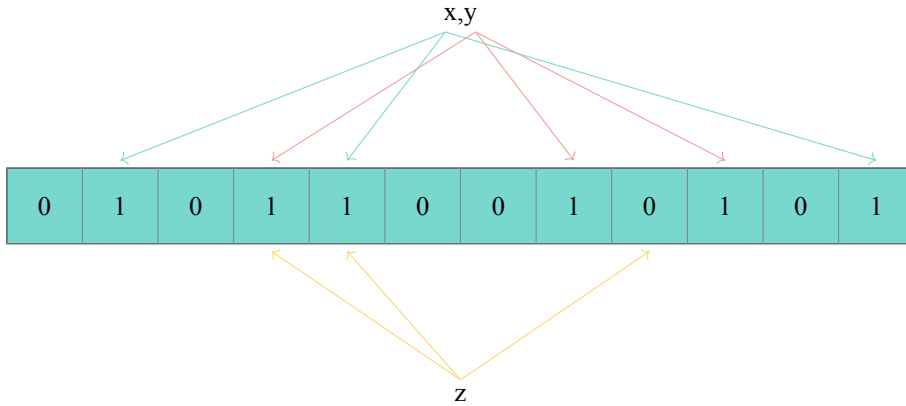
In many software projects, determining whether an element exists in a collection is a common task. For instance, in word processing software, it is essential to verify if an English word is spelled correctly by checking if it exists in a known dictionary. Similarly, in the FBI, it is necessary to determine if a suspect's name appears on the suspect list to issue an FBI Warning. Likewise, in a web crawler, it is crucial to determine if a URL has been visited. To achieve this, the most straightforward approach is to store all the elements in the collection in the computer and compare them directly with the new element encountered. Usually, computer sets use a hash table to store the collection, which is fast and accurate but consumes more space.

For small data collections, this approach is adequate. However, when the collection is extensive, the low storage efficiency of hash tables becomes apparent. For example, email providers such as Yahoo or Gmail need to filter spam regularly. One way to accomplish this is to keep a record of the email addresses of those who send spam. However, since spammers keep registering new addresses, storing all of them would require numerous network servers. Storing 100 million email addresses in a hash table requires about 1.6 GB of memory, which is a considerable amount. Furthermore, hash tables have a load factor that may not allow the full utilization of space. Additionally, if the data set is stored on a remote server and input is accepted locally, it may not be feasible to construct a hash table since the dataset is too large to read into memory at once.

10.4.1 The Bloom Filter

When dealing with problems of identifying elements in a set or collection, Bloom Filters are a useful data structure to consider. These probabilistic data structures were introduced by Burton Howard Bloom in 1970 and consist of a long binary vector and random mapping functions. They are particularly useful in terms of space and query efficiency, although they do have a certain error rate and deleting elements can be challenging.

A Bloom Filter, shown below, is composed of a binary vector (bit array) that can store n data elements and k hash functions. Two basic operations are supported: insert and query. However, the number of values that can be inserted is fixed at design time, so the size of the Bloom Filter must be specifically designed for the problem at hand.



Initially, all positions in the Bloom Filter are set to 0. When data is inserted, k hash functions are used to determine the position of the data in the filter and set the corresponding positions to 1. For instance, when $k = 3$, three hash values are calculated as indices, and their corresponding positions are set to 1. When querying, k hash functions are also used to generate k hash values as indices. If all indices correspond to values that are 1, then the element may exist in the set. In the given figure, x and y are stored in the Bloom Filter, but the last hash value for z is 0, so it definitely does not exist in the filter. To experience a Bloom Filter, visit [bloomfilter](#).

To determine the optimal number of elements m to be stored in a Bloom filter of length n with a tolerable error rate of ϵ , use the formula:

$$m = -\frac{n \ln \epsilon}{(\ln 2)^2} \quad (10.3)$$

To calculate the required number of hash functions k , use the formula:

$$k = -\frac{\ln \epsilon}{\ln 2} = -\log_2 \epsilon \quad (10.4)$$

For example, if the tolerable error rate ϵ is 8%, then k would be 3, and a larger k value would indicate a higher error rate. It is possible to simulate k hash functions by combining two basic hash functions and the number of iterations, without changing the fault-tolerance rate, using the following formula:

$$g_i(x) = h_1(x) + i h_2(x) \quad (10.5)$$

To implement a Bloom filter, a struct can be used to encapsulate all the necessary information, including the set of storage bits and hash functions. Since Bloom filter only determines the existence of a value and not the value itself, it must be able to accommodate data of any type using generics. The Boolean values of 1 and 0 can be used to represent the storage bits, and they can be saved to a Vec for easy implementation. During the judgment process, the Boolean value can indicate whether the value exists or not.

```
1 // bloom_filter.rs
2
3 use std::collections::hash_map::DefaultHasher;
4
5 // Definition of bloom filter
6 struct BloomFilter<T> {
7     bits: Vec<bool>,           // Bit bucket
8     hash_fn_count: usize,      // Number of hash functions
9     hashers: [DefaultHasher; 2], // Two hash functions
10 }
```

The code above cannot compile since the generic type `T` is not used by any field, which violates Rust's rules. To make it compile, we can use Rust's phantom data to occupy the space of `T`. Phantom data can pretend to use `T` without actually using it, deceiving the compiler into passing the code. Additionally, to support data with an undefined size at compile time, we add the `?Sized` trait to make the filter work with such data types. We prefix the unused field with `_phantom` to indicate that it is not used and occupies 0 bytes. However, it includes `T` to deceive the compiler. We also simulate `k` hash functions using two random hash functions.

```

1 // bloom_filter.rs
2
3 use std::collections::hash_map::DefaultHasher;
4 use std::marker::PhantomData;
5
6 // Definition of a new bloom filter
7 struct BloomFilter<T: ?Sized> {
8     bits: Vec<bool>,
9     hash_fn_count: usize,
10    hashers: [DefaultHasher; 2],
11    _phantom: PhantomData<T>,
12    // T is a placeholder
13    // for deceiving the compiler
14 }

```

To implement the filter's functionality, we need to implement three functions: the initialization function `new`, the function to add elements `insert`, and the function to check contains. We also need auxiliary functions to implement the first three functions. The `new` function calculates the size of `m` based on the fault tolerance rate and the approximate storage scale, and initializes the filter.

```

1 // bloom_filter.rs
2
3 use std::hash::{BuildHasher, Hash, Hasher};
4 use std::collections::hash_map::RandomState;
5
6 impl<T: ?Sized + Hash> BloomFilter<T> {
7     fn new(cap: usize, ert: f64) -> Self {
8         let ln22 = std::f64::consts::LN_2.powf(2f64);
9
10        // Calculate the size of the bit bucket and
11        // the number of hash functions
12        let bits_count = -1f64 * cap as f64 * ert.ln() / ln22;
13        let hash_fn_count = -1f64 * ert.log2();
14
15        // Random hash function
16        let hashers = [
17            RandomState::new().build_hasher(),
18            RandomState::new().build_hasher(),
19        ];
20
21        Self {
22            bits: vec![false; bits_count.ceil() as usize],
23            hash_fn_count: hash_fn_count.ceil() as usize,
24            hashers: hashers,
25            _phantom: PhantomData,

```

```

26     }
27 }
28
29 // Set the corresponding bit of the bit bucket to true
30 // according to hash_fn_count
31 fn insert(&mut self, elem: &T) {
32     let hashes = self.make_hash(elem);
33     for fn_i in 0..self.hash_fn_count {
34         let index = self.get_index(hashes, fn_i as u64);
35         self.bits[index] = true;
36     }
37 }
38
39 // Data query
40 fn contains(&self, elem: &T) -> bool {
41     let hashes = self.make_hash(elem);
42     (0..self.hash_fn_count).all(|fn_i| {
43         let index = self.get_index(hashes, fn_i as u64);
44         self.bits[index]
45     })
46 }
47
48 // Calculate hash value
49 fn make_hash(&self, elem: &T) -> (u64, u64) {
50     let hasher1 = &mut self.hashers[0].clone();
51     let hasher2 = &mut self.hashers[1].clone();
52     elem.hash(hasher1);
53     elem.hash(hasher2);
54
55     (hasher1.finish(), hasher2.finish())
56 }
57
58 // Get the index of a certain bit in the bit bucket
59 fn get_index(&self, (h1,h2):(u64,u64),fn_i:u64) -> usize {
60     let ih2 = fn_i.wrapping_mul(h2);
61     let h1pih2 = h1.wrapping_add(ih2);
62     (h1pih2 % self.bits.len() as u64) as usize
63 }
64 }

```

```

1 fn main() {
2     let mut bf = BloomFilter::new(100, 0.08);
3     (0..20).for_each(|i| bf.insert(&i));
4     let res1 = bf.contains(&2);
5     let res2 = bf.contains(&200);
6     println!("2 in bf: {res1}, 200 in bf: {res2}");
7     // 2 in bf: true, 200 in bf: false
8 }

```

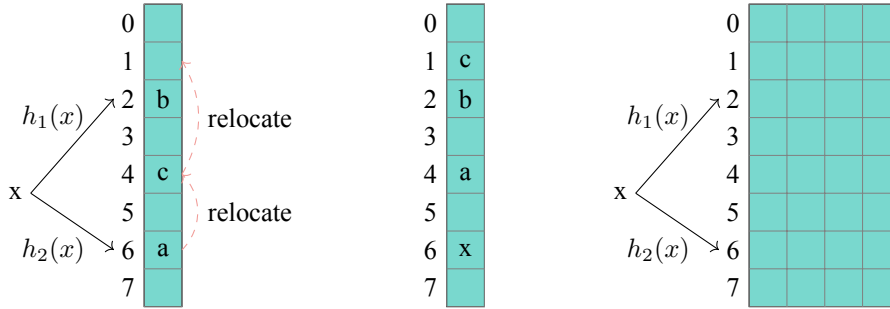
By analyzing the Bloom filter, we can conclude that its space complexity is $O(m)$, while the time complexity of insert and contains is $O(k)$. However, k is typically small, so we can consider the time complexity as $O(1)$.

10.4.2 The Cuckoo Filter

The Bloom filter, as discussed earlier, is relatively easy to implement, but it has a couple of disadvantages. Firstly, as more data is inserted, the error rate increases. Secondly, it does not support the deletion of data. Lastly, it uses random access memory, which does not perform well on CPUs with cache, as stated in CloudFlare’s article “When Bloom filters don’t bloom”. To overcome these issues, the Cuckoo filter^[18] was developed as an improved version of the Bloom filter. The Cuckoo filter uses paired hash functions to map data to two positions: one is the saved position and the other is the alternate position, used to handle collisions.

The Cuckoo filter uses paired hash functions to map data to two positions: one is the saved position, and the other is the alternate position, which is used to handle collisions. It is named after the cuckoo bird, which lays its eggs in other birds’ nests and lets them hatch and raise its offspring. The cuckoo chick hatches earlier than the other chicks and pushes the other eggs out of the nest to monopolize the food from its foster parents. Inspired by this biological phenomenon, the Cuckoo filter also kicks out the original element at the collided position and replaces it with the new element. If the alternate position is also occupied, it repeats this process until it reaches a predefined limit and confirms that the hash table is full.

Unlike the Bloom filter, the Cuckoo filter stores not only 0s and 1s but also a fixed number of bits of data, known as fingerprints. The length of the fingerprint is determined by the false positive rate ϵ , with smaller ϵ requiring longer fingerprints. The Cuckoo filter is based on the Cuckoo hash table, which is extended into a two-dimensional matrix to store multiple fingerprints in each bucket.



When inserting an element into the Cuckoo filter, if two buckets are already occupied, the filter randomly chooses one of the elements to be kicked out to an alternate position, and the previously kicked-out element takes the newly emptied spot. This filter supports insertion, deletion, and lookup operations.

In the standard Cuckoo hash table shown in the left diagram, inserting a new item into the existing hash table requires accessing the original item to determine its position and make room for the new item. However, since the Cuckoo filter only stores fingerprints, there is no way to rehash the original item to find its alternate position. To overcome this limitation, a technique called partial-key Cuckoo hashing can be used to obtain an item’s alternate position based on its fingerprint. For item x , the two candidate bucket indices are calculated using the same hashing function as follows:

$$\begin{aligned} h_1(x) &= \text{hash}(x) \\ h_2(x) &= h_1(x) \oplus \text{hash}(\text{fingureprint}(x)) \end{aligned} \quad (10.6)$$

The XOR operation \oplus ensures that $h_1(x)$ and $h_2(x)$ can be calculated using the same formula, so it is not necessary to know the value of x to determine the position of the alternate bucket using Equation (10.7).

$$j = i \oplus \text{hash}(\text{fingureprint}(x)) \quad (10.7)$$

The lookup method is straightforward: first use Equation (10.6) to calculate the fingerprint of the item to be searched and the positions of the two alternate buckets, then read the values in those buckets. If any of the buckets contains a value that matches the fingerprint of the item being searched, then it exists

in the filter. The deletion method is similarly simple: check the values in the two alternate buckets, and if there is a match, delete the copy of the fingerprint in that bucket. It is important to ensure that the item is inserted before attempting to delete it, otherwise it may accidentally delete other values with the same fingerprint.

After extensive experimentation and testing, a bucket size of 4 was found to provide excellent performance, even the best performance. The Cuckoo filter boasts four main advantages over Bloom filters:

- (1) Supports dynamic addition and deletion of items.
- (2) Higher lookup performance than Bloom filters, even when approaching full capacity.
- (3) Easier to implement than other Bloom filter alternatives, such as quotient filters.
- (4) In practical applications, it uses less space than Bloom filters if the false positive rate ϵ is less than 3%.

In addition to Bloom filters and Cuckoo filters, there are many other filters with various characteristics that interested readers can search for and refer to.

Table 10.1: Comparison of Various Filters

Filter Type	Space Usage	Hash Function	Deletion
Bloom Filter	1	k	no
Blocked Bloom Filter	1x	1	no
Counting Bloom Filter	3x-4x	k	yes
D-left Counting Bloom Filter	1.5x-2x	d	yes
Quotient Filter	1x-1.2x	≥ 1	yes
Cuckoo Filter	$\leq 1x$	2	yes

To implement the Cuckoo Filter, we need to extend the Bloom Filter to two dimensions. This requires us to create two new data structures: FingerPrint to store the fingerprints, and Bucket to store the FingerPrints. Since the implementation involves random access and hashing operations, we utilize libraries such as Rng and Serde in the code. We implement the Cuckoo Filter as a Rust library, where bucket.rs includes the definitions and operations of FingerPrint and Bucket, and util.rs includes the struct FaI for calculating fingerprints and bucket indexes. The code structure is as follows.

```
shieber@Kew:cuckoofilter/ tree
/Cargo.toml
/src
|- bucket.rs
|- lib.rs
|- util.rs
```

The Cuckoo Filter code is quite lengthy, so we will only list the lib.rs file here. For the remaining code, please refer to the source code provided with the book.

```
1 // lib.rs
2 mod bucket;
3 mod util;
4
5 use std::fmt;
6 use std::cmp::max;
7 use std::iter::repeat;
8 use std::error::Error;
9 use std::hash::{Hash, Hasher};
10 use std::marker::PhantomData;
11 use std::collections::hash_map::DefaultHasher;
12
```



```

13 // Support serialization
14 use rand::Rng;
15 #[cfg(feature = "serde_support")]
16 use serde_derive::{Serialize, Deserialize};
17
18 use crate::util::FaI;
19 use crate::bucket::{Bucket, FingerPrint,
20                     BUCKET_SIZE, FINGERPRINT_SIZE};
21
22 const MAX_RELOCATION: u32 = 100;
23 const DEFAULT_CAPACITY: usize = (1 << 20) - 1;
24
25 // Error handling
26 #[derive(Debug)]
27 enum CuckooError {
28     NotEnoughSpace,
29 }
30
31 // add print function
32 impl fmt::Display for CuckooError {
33     fn fmt(&self, f: &mut fmt::Formatter) -> fmt::Result {
34         f.write_str("NotEnoughSpace")
35     }
36 }
37
38 impl Error for CuckooError {
39     fn description(&self) -> &str {
40         "Not enough space to save element, operation failed!"
41     }
42 }
43
44 // Definition of cuckoo filter
45 struct CuckooFilter<H> {
46     buckets: Box<[Bucket]>, // Bucket
47     len: usize,             // length
48     _phantom: PhantomData<H>,
49 }
50
51 // add default value
52 impl Default for CuckooFilter<DefaultHasher> {
53     fn default() -> Self {
54         Self::new()
55     }
56 }
57
58 impl CuckooFilter<DefaultHasher> {
59     fn new() -> Self {
60         Self::with_capacity(DEFAULT_CAPACITY)
61     }
62 }
63
64 impl<H: Hasher + Default> CuckooFilter<H> {

```

```

65     fn with_capacity(cap: usize) -> Self {
66         let capacity = max(1, cap.next_power_of_two()
67                             / BUCKET_SIZE);
68         Self {
69             // build a CuckooFilter with capacity buckets
70             buckets: repeat(Bucket::new())
71                           .take(capacity)
72                           .collect::<Vec<_>>(),
73             .into_boxed_slice(),
74             len: 0,
75             _phantom: PhantomData,
76         }
77     }
78
79     fn try_insert<T: ?Sized + Hash>(&mut self, elem: &T)
80     -> Result<bool, CuckooError> {
81         if self.contains(elem) {
82             Ok(false)
83         } else {
84             self.insert(elem).map(|_| true)
85         }
86     }
87
88     fn insert<T: ?Sized + Hash>(&mut self, elem: &T)
89     -> Result<(), CuckooError> {
90         let fai = FaI::from_data::<_, H>(elem)
91         if self.put(fai.fp, fai.i1)
92             || self.put(fai.fp, fai.i2) {
93             return Ok(());
94         }
95
96         // collision, relocate
97         let mut rng = rand::thread_rng();
98         let mut i = fai.random_index(&mut rng);
99         let mut fp = fai.fp;
100         for _ in 0..MAX_RELOCATION {
101             let other_fp;
102             {
103                 let loc = &mut self.buckets[i % self.len]
104                             .buffer[rng.gen_range(0,
105                                     BUCKET_SIZE)];
106                 other_fp = *loc;
107                 *loc = fp;
108                 i = FaI::get_alt_index::<H>(other_fp, i);
109             }
110             if self.put(other_fp, i) {
111                 return Ok(());
112             }
113             fp = other_fp;
114         }
115         Err(CuckooError::NotEnoughSpace)
116     }

```

```

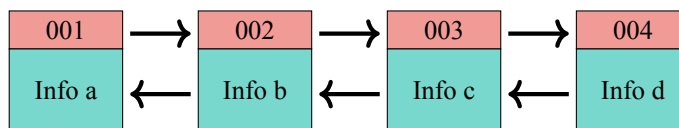
117
118 // add fingerprint
119 fn put(&mut self, fp: FingerPrint, i: usize) -> bool {
120     if self.buckets[i % self.len].insert(fp) {
121         self.len += 1;
122         true
123     } else {
124         false
125     }
126 }
127
128 fn remove(&mut self, fp: FingerPrint, i: usize) -> bool {
129     if self.buckets[i % self.len].delete(fp) {
130         self.len -= 1;
131         true
132     } else {
133         false
134     }
135 }
136
137 fn contains<T: ?Sized + Hash>(&self, elem: &T) -> bool {
138     let FaI { fp, i1, i2 } = FaI::from_data::<_, H>(elem);
139     self.buckets[i1 % self.len]
140         .get_fp_index(fp)
141         .or_else(|| {
142             self.buckets[i2 % self.len]
143                 .get_fp_index(fp)
144         })
145         .is_some()
146 }
147 }

```

As we can see from the code, the Cuckoo Filter supports insertion, deletion, and querying operations.

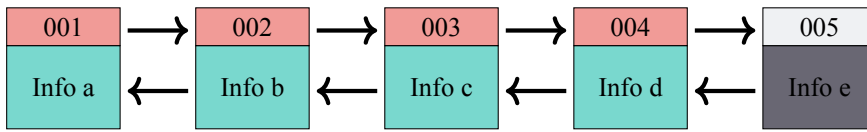
10.5 Least Recently Used(LRU) Algorithm

The cache eviction algorithm, or page replacement algorithm, is a commonly used memory management technique in virtual paging storage and data caching. Its principle is based on the assumption that if data has been recently accessed, it is more likely to be accessed again in the future. Therefore, data blocks that are in memory but not used will be removed based on which data is the least recently used, freeing up space and saving memory. LRU (Least Recently Used) is a popular eviction algorithm used to eliminate data based on the access record of data when storage is limited. Assuming a hash-linked list is used to cache user information with a capacity of 5, and currently caching 4 user information, as shown in the figure.

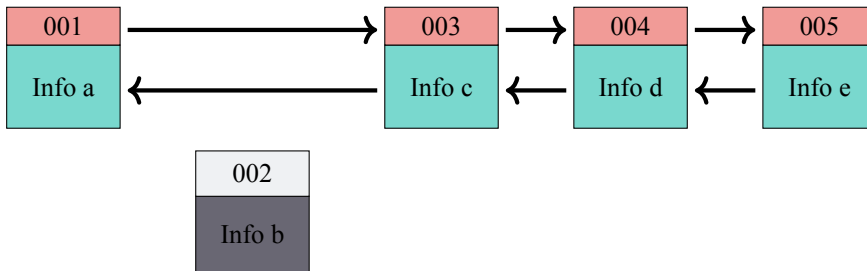


When a client queries user 5, the data is not in the hash-linked list, so it needs to be read from the database and inserted into the cache. At this point, the rightmost node of the linked list represents the

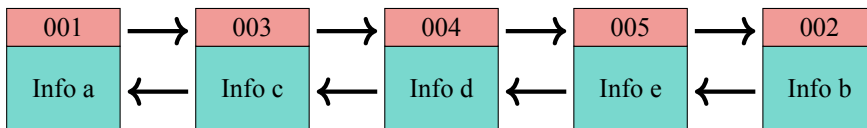
most recently accessed user 5, and the leftmost represents the least recently accessed user 1.



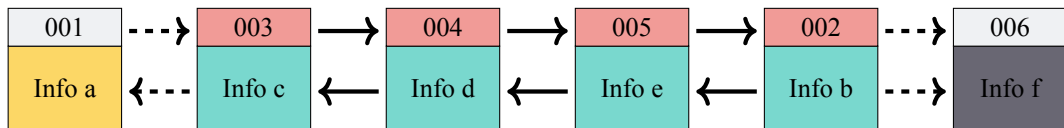
Next, when another client accesses user 2, the data is found in the hash-linked list, so user 2 is simply removed from its current position and reinserted into the rightmost of the linked list.



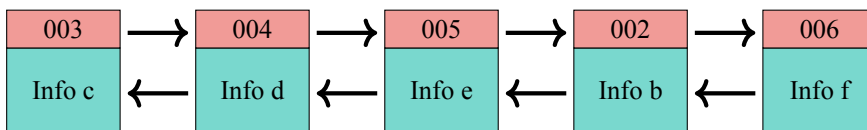
After these accesses, the hash-linked list looks like the figure shown.



Later, a client accesses user 6, but the data is not in the cache, so it also needs to be inserted.



However, since the cache has reached its limit, the least recently accessed data, user 1, needs to be deleted first. Then, user 6 can be inserted at the rightmost end of the hash-linked list.



The LRU algorithm is a common cache eviction algorithm that frees up memory by removing the least recently used data. To implement this algorithm, we need to define the data structure and operations abstracted from the graph. This includes managing keys, entries, and front and back pointers. The necessary operation functions include insert, remove, contains, and auxiliary functions.

```
1 // lru.rs
2 use std::collections::HashMap;
3
4 // Definition for entry of LRU
5 struct Entry<K, V> {
6     key: K,
7     val: Option<V>,
```

```

8     next: Option<usize>,
9     prev: Option<usize>,
10 }
11
12 struct LRUCache<K, V> {
13     cap: usize,
14     head: Option<usize>,
15     tail: Option<usize>,
16     map: HashMap<K, usize>,
17     entries: Vec<Entry<K, V>>,
18 }

```

To store the keys, we will use a `HashMap`, and a `Vec` to store the entries. The head and tail pointers can be simplified to the index of the `Vec`. To customize the cache capacity, we can implement a `with_capacity` function, with new setting the capacity to 100 by default.

```

1 // lru.rs
2 use std::hash::Hash;
3
4 const CACHE_SIZE: usize = 100;
5
6 impl<K: Clone + Hash + Eq, V> LRUCache<K, V> {
7     fn new() -> Self {
8         Self::with_capacity(CACHE_SIZE)
9     }
10
11     fn len(&self) -> usize {
12         self.map.len()
13     }
14
15     fn is_empty(&self) -> bool {
16         self.map.is_empty()
17     }
18
19     fn is_full(&self) -> bool {
20         self.map.len() == self.cap
21     }
22
23     fn with_capacity(cap: usize) -> Self {
24         LRUCache {
25             cap: cap,
26             head: None,
27             tail: None,
28             map: HashMap::with_capacity(cap),
29             entries: Vec::with_capacity(cap),
30         }
31     }
32 }

```

When inserting data, if the key already exists, the value will be updated directly and the original value will be returned. If the inserted value does not exist, the returned value should be `None`, hence the use of `Option` as the return type. The access method is used to remove the original value and update the value, and `ensure_room` is used to remove the least recently used data when the cache reaches capacity.

```

1 // lru.rs
2
3 impl<K: Clone + Hash + Eq, V> LRUCache<K, V> {
4     fn insert(&mut self, key: K, val: V) -> Option<V> {
5         if self.map.contains_key(&key) {
6             // Update if key exists
7             self.access(&key);
8             let entry = &mut self.entries[self.head.unwrap()];
9             let old_val = entry.val.take();
10            entry.val = Some(val);
11            old_val
12        } else {
13            // insert if key not exists
14            self.ensure_room();
15
16            // Update the original head pointer
17            let index = self.entries.len();
18            self.head.map(|e| {
19                self.entries[e].prev = Some(index);
20            });
21
22            // The new head node
23            self.entries.push(Entry {
24                key: key.clone(),
25                val: Some(val),
26                prev: None,
27                next: self.head,
28            });
29
30            self.head = Some(index);
31            self.tail = self.tail.or(self.head);
32            self.map.insert(key, index);
33
34            None
35        }
36    }
37
38    fn get(&mut self, key: &K) -> Option<&V> {
39        if self.contains(key) { self.access(key); }
40
41        let entries = &self.entries;
42        self.map.get(key).and_then(move |&i| {
43            entries[i].val.as_ref()
44        })
45    }
46
47    fn get_mut(&mut self, key: &K) -> Option<&mut V> {
48        if self.contains(key) { self.access(key); }
49
50        let entries = &mut self.entries;
51        self.map.get(key).and_then(move |&i| {

```

```

52         entries[i].val.as_mut()
53     })
54 }
55
56 fn contains(&mut self, key: &K) -> bool {
57     self.map.contains_key(key)
58 }
59
60 // Ensure there is enough capacity, remove the
61 // least recently used item if full
62 fn ensure_room(&mut self) {
63     if self.cap == self.len() {
64         self.remove_tail();
65     }
66 }
67
68 fn remove_tail(&mut self) {
69     if let Some(index) = self.tail {
70         self.remove_from_list(index);
71         let key = &self.entries[index].key;
72         self.map.remove(key);
73     }
74     if self.tail.is_none() {
75         self.head = None;
76     }
77 }
78
79 // Get the value of a key, remove the
80 // old position and add to the head
81 fn access(&mut self, key: &K) {
82     let i = *self.map.get(key).unwrap();
83     self.remove_from_list(i);
84     self.head = Some(i);
85 }
86
87 fn remove(&mut self, key: &K) -> Option<V> {
88     self.map.remove(&key).map(|index| {
89         self.remove_from_list(index);
90         self.entries[index].val.take().unwrap()
91     })
92 }
93
94 fn remove_from_list(&mut self, i: usize) {
95     let (prev, next) = {
96         let entry = self.entries.get_mut(i).unwrap();
97         (entry.prev, entry.next)
98     };
99
100     match (prev, next) {
101         // The data item is in the middle of the cache
102         (Some(j), Some(k)) => {
103             let head = &mut self.entries[j];

```

```

104         head.next = next;
105         let next = &mut self.entries[k];
106         next.prev = prev;
107     },
108     // The data item is at the end of the cache
109     (Some(j), None) => {
110         let head = &mut self.entries[j];
111         head.next = None;
112         self.tail = prev;
113     },
114     // The data item is at the head of the cache
115     - => {
116         if self.len() > 1 {
117             let head = &mut self.entries[0];
118             head.next = None;
119             let next = &mut self.entries[1];
120             next.prev = None;
121         }
122     },
123 }
124 }
125 }

```

The following are use example of LRU.

```

1 // lru.rs
2
3 fn main() {
4     let mut cache = LRUCache::with_capacity(2);
5     cache.insert("foo", 1);
6     cache.insert("bar", 2);
7     cache.insert("baz", 3);
8     cache.insert("tik", 4);
9     cache.insert("tok", 5);
10
11     assert!(!cache.contains(&"foo"));
12     assert!(!cache.contains(&"bar"));
13     assert!(cache.contains(&"baz"));
14     assert!(cache.contains(&"tik"));
15
16     cache.insert("qux", 6);
17     assert!(cache.contains(&"qux"));
18 }

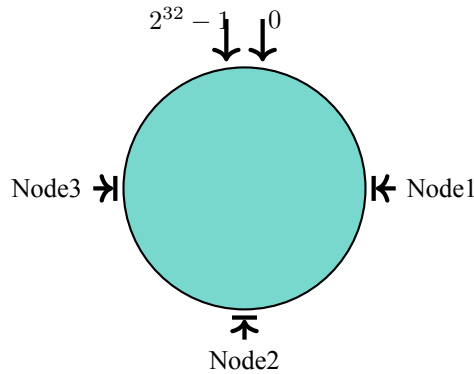
```

10.6 Consistent Hashing

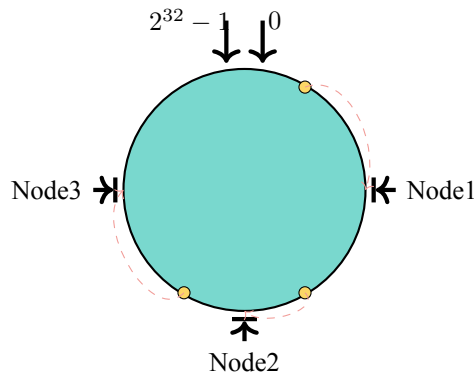
The Consistent Hashing algorithm was initially proposed by Karger et al. of MIT to address hot-spot issues in distributed caching. However, after years of development, it has become a widely applied algorithm.

To understand its application, let's consider caching images in Redis. A single Redis instance can handle small data volumes and low access rates, and a master-slave configuration may be sufficient.

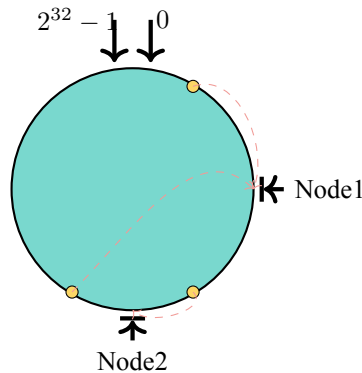
However, when data volumes grow, and access rates increase, a cluster must be created to distribute data across multiple machines. For example, if there are 5 machines, the position index of an image would be $\text{index} = \text{hash}(\text{key}) \% 5$, where key is an index related to the image. However, adding or removing machines changes N , and previously calculated indices will become invalid. This is where consistent hashing comes in. It creates a circle that ranges from 0 to $2^{32} - 1$, mapping data to a position on the circle.



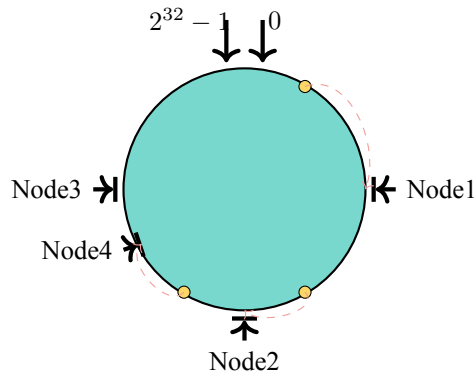
To achieve caching, data must be added to a specific segment of the circle based on its hash value. The corresponding node located just clockwise from the segment will store the data.



Suppose Node3 fails, causing data between Node2 and Node3 to be transferred to Node1.



When a new machine, Node4, is added, the data originally stored on Node3 will now be stored on Node4.



The Consistent Hashing algorithm is fault-tolerant and scalable since it only relocates a small portion of the data when nodes are added or removed. This feature enables the algorithm to achieve consistency. To implement Consistent Hashing, a Ring is needed to store nodes representing machines.

```

1 // conshash.rs
2
3 use std::fmt::Debug;
4 use std::string::ToString;
5 use std::hash::{Hash, Hasher};
6 use std::collections::{BTreeMap, hash_map::DefaultHasher};
7
8 // Ring node definition, used for storing the host, ip and port
9 #[derive(Clone, Debug)]
10 struct Node {
11     host: &'static str,
12     ip: &'static str,
13     port: u16,
14 }
15
16 // implement a to_string function
17 impl ToString for Node {
18     fn to_string(&self) -> String {
19         self.ip.to_string() + &self.port.to_string()
20     }
21 }
22
23 // Definition for the Ring
24 struct Ring<T: Clone + ToString + Debug> {
25     replicas: usize,           // partition number
26     ring: BTreeMap<u64, T>, // store data
27 }

```

Replicas in the Ring are utilized to avoid node clustering and ensure even data distribution across nodes. Multiple virtual nodes can be created for each node to solve the node clustering problem.

The default hash calculator provided by the standard library, with a default of 10 nodes, can be used for hash calculation, or custom nodes can be created. For the Consistent Hashing algorithm, at least node insertion, node deletion, and query functions must be supported. Additionally, batch versions of insertion and deletion can be implemented for batch processing.

```

1 // conshash.rs
2
3 const DEFAULT_REPLICAS: usize = 10;
4
5 // Hash calculation function
6 fn hash<T: Hash>(val: &T) -> u64 {
7     let mut hasher = DefaultHasher::new();
8     val.hash(&mut hasher);
9
10    hasher.finish()
11 }
12
13 impl<T> Ring<T> where T: Clone + ToString + Debug {
14     fn new() -> Self {
15         Self::with_capacity(DEFAULT_REPLICAS)
16     }
17
18     // new with a replicas parameter
19     fn with_capacity(replicas: usize) -> Self {
20         Ring {
21             replicas: replicas,
22             ring: BTreeMap::new()
23         }
24     }
25
26     // Batch insertion of nodes
27     fn add_multi(&mut self, nodes: &[T]) {
28         if !nodes.is_empty() {
29             for node in nodes.iter() {
30                 self.add(node);
31             }
32         }
33     }
34
35     fn add(&mut self, node: &T) {
36         for i in 0..self.replicas {
37             let key = hash(&(node.to_string()
38                             + &i.to_string()));
39             self.ring.insert(key, node.clone());
40         }
41     }
42
43     // Batch deletion of nodes
44     fn remove_multi(&mut self, nodes: &[T]) {
45         if !nodes.is_empty() {
46             for node in nodes.iter() {
47                 self.remove(node);
48             }
49         }
50     }
51

```

```

52     fn remove(&mut self, node: &T) {
53         // assertion to ensure that the node exists
54         assert!(!self.ring.is_empty());
55
56         for i in 0..self.replicas {
57             let key = hash(&(node.to_string()
58                             + &i.to_string()));
59             self.ring.remove(&key);
60         }
61     }
62
63     // Get nodes
64     fn get(&self, key: u64) -> Option<&T> {
65         if self.ring.is_empty() {
66             return None;
67         }
68
69         let mut keys = self.ring.keys();
70         keys.find(|&k| k >= &key)
71             .and_then(|k| self.ring.get(k))
72             .or(keys.nth(0).and_then(|x| self.ring.get(x)))
73     }
74 }
75
76 fn main() {
77     let replica = 3;
78     let mut ring = Ring::with_capacity(replica);
79     let node = Node{
80         host:"localhost",
81         ip:"127.0.0.1",
82         port:23
83     };
84     ring.add(&node);
85
86     for i in 0..replica {
87         let key = hash(&(node.to_string() + &i.to_string()));
88         let res = ring.get(key);
89         assert_eq!(node.host, res.unwrap().host);
90     }
91
92     println!("{:?}", &node);
93     ring.remove(&node);
94     // Node { host: "localhost", ip: "127.0.0.1", port: 23 }
95 }

```

10.7 Base58 Encode and Decode

Base58 is an encoding algorithm that only uses certain characters from the Base64 table, removing the characters highlighted in red. It was originally used to represent Bitcoin wallet addresses and was invented by Satoshi Nakamoto.

Table 10.2: Base 64 encoding characters

ASCII	Char	ASCII	Char	ASCII	Char	ASCII	Char	ASCII	Char
0	0	13	D	26	Q	39	d	52	q
1	1	14	E	27	R	40	e	53	r
2	2	15	F	28	S	41	f	54	s
3	3	16	G	29	T	42	g	55	t
4	4	17	H	30	U	43	h	56	u
5	5	18	I	31	V	44	i	57	v
6	6	19	J	32	W	45	j	58	w
7	7	20	K	33	X	46	k	59	x
8	8	21	L	34	Y	47	l	60	y
9	9	22	M	35	Z	48	m	61	z
10	A	23	N	36	a	49	n	62	+
11	B	24	O	37	b	50	o	63	/
12	C	25	P	38	c	51	p		

To avoid ambiguous characters, Base58 removes 0 (zero), O (capital O), I (capital i), l (lowercase L), plus sign (+), and slash (/) from the Base64 table. The remaining 58 characters are used as encoding characters. These characters are easily distinguishable and also prevent problems with line breaks when copying the code.

Base58 encoding is essentially a conversion from a large number base. First, the character is converted to ASCII and then to decimal, followed by 58 base conversion. Finally, the corresponding character is selected from the encoding table to form the Base58 encoding string. However, due to the need for base conversion, the efficiency of Base58 encoding is relatively low. The encoding algorithm (10.1) illustrates the encoding principle.

Algorithm 10.1: Base58 Encoding Algorithm

Data: Original string s

Result: Encoded string $b58_str$

1 Initialize an empty string $b58_str$ to store the result.

2 **for** $c \in s$ **do**

3 Convert byte c in s to ASCII value (256 base).

4 Convert 256 base number to decimal.

5 Convert decimal number to 58 base number.

6 Convert 58 base number to corresponding character according to the Base58 character table.

7 Append the resulting character to $b58_str$.

8 **end**

9 Return the encoded string $b58_str$.

Decoding is the reverse process of encoding and involves converting from a large number base. The characters in the Base58 string are first converted to ASCII values, then to 10 decimal values, followed by conversion to 256 decimal values, and finally to ASCII characters. The decoding process is described in detail in Algorithm (10.2).

Algorithm 10.2: Base58 Decoding Algorithm**Data:** Encoded string b58**Result:** Decoded string new_str

```

1 Initialize an empty string new_str to store the result
2 for  $c \in b58$  do
3   Convert byte c in b58 to ASCII value (58 base)
4   Convert 58 base number to 10 decimal value
5   Convert 10 decimal value to 256 decimal value
6   Convert 256 decimal value to corresponding ASCII character
7   Append the resulting character to new_str
8 end
9 Return the decoded string new_str

```

Encoding and decoding are essentially two-way string transformations between different spaces, similar to a mapping in the encoding space. To implement a Base58 encoder and decoder, we first prepare the encoding characters ALPHABET and the encoding conversion table DIGITS_MAP. Additionally, we define the maximum conversion base 58 and substitute 1 for the leading 0 as constants. Using constants for operations is preferable to using magic numbers directly, as it improves code readability.

```

1 // base58.rs
2
3 // Conversion base 58
4 const BIG_RADIX: u32 = 58;
5
6 // Substitute 1 for the leading 0
7 const ALPHABET_INDEX_0: char = '1';
8
9 // Base58 encoding characters
10 const ALPHABET: &[u8;58] = b"123456789ABCDEFGHJKLMNPQRSTUVWXYZ
11 abcdefghijklmnopqrstuvwxyz";
12
13 // Mapping relationship between the bases
14 const DIGITS_MAP: &'static [u8] = &[
15     255,255,255,255,255,255,255,255,255,255,255,255,255,255,255,255,
16     255,255,255,255,255,255,255,255,255,255,255,255,255,255,255,255,
17     255,255,255,255,255,255,255,255,255,255,255,255,255,255,255,255,
18     255, 0, 1, 2, 3, 4, 5, 6, 7,
19     8,255,255,255,255,255,255,
20     255, 9,10,11,12,13,14,15,16,255,17,18,19,20,
21     21,255,
22     22,23,24,25,26,27,28,29,30,31,
23     32,255,255,255,255,255,
24     255,33,34,35,36,37,38,39,40,41,42,43,255,44,45,
25     46,
26     47,48,49,50,51,52,53,54,55,56,
27     57,255,255,255,255,255,
28 ];

```

To handle potential errors during encoding and decoding, we have implemented a custom error type for Base58 encoding to manage illegal characters, length errors, and other cases. Two traits, `Encoder` and `Decoder`, have been created for encoding and decoding, respectively. `Encoder` and `Decoder` contain the methods `encode_to_base58` and `decode_from_base58`, respectively.

```

1 // base58.rs
2
3 // Decoding error type
4 #[derive(Debug, PartialEq)]
5 pub enum DecodeError {
6     Invalid,
7     InvalidLength,
8     InvalidCharacter(char, usize),
9 }
10
11 // Encoding and decoding traits
12 pub trait Encoder {
13     // Encoding method
14     fn encode_to_base58(&self) -> String;
15 }
16
17 pub trait Decoder {
18     // Decoding method
19     fn decode_from_base58(&self) -> Result<String, DecodeError>;
20 }

```

In the next step, we will implement the methods that correspond to the two traits mentioned earlier. Although these traits are implemented for strings, it is better to use `u8` for internal calculations since the characters in the string may contain multiple `u8` values.

```

1 // base58.rs
2
3 // Implement Base58 encoding
4 impl Encoder for str {
5     fn encode_to_base58(&self) -> String {
6         // Convert to bytes for processing
7         let str_u8 = self.as_bytes();
8         // Count the number of leading zeros
9         let zero_count = str_u8.iter()
10             .take_while(|&&x| x == 0)
11             .count();
12         // Space required after conversion is log(256)/log(58),
13         // which is approximately 1.38 times the original data.
14         // We don't need leading zeros
15         let size = (str_u8.len() - zero_count) * 138 / 100 + 1;
16         // Convert characters from one base to another
17         let mut i = zero_count;
18         let mut high = size - 1;
19         let mut buffer = vec![0u8; size];
20         while i < str_u8.len() {
21             // j is the decreasing index, corresponding to
22             // counting from the back
23             let mut j = size - 1;

```

```

24
25         // carry is the character read from the front
26         let mut carry = str_u8[i] as u32;
27
28         // Store the converted data from back to front
29         // in turn
30         while j > high || carry != 0 {
31             carry += 256 * buffer[j] as u32;
32             buffer[j] = (carry % BIG_RADIX) as u8;
33             carry /= BIG_RADIX;
34
35             if j > 0 {
36                 j -= 1;
37             }
38         }
39         i += 1;
40         high = j;
41     }
42
43     // Handle multiple leading zeros
44     let mut b58_str = String::new();
45     for _ in 0..zero_count {
46         b58_str.push(ALPHABET_INDEX_0);
47     }
48
49     // Get the encoded characters and concatenate
50     // them into a string
51     let mut j = buffer.iter()
52         .take_while(|&x| x == 0)
53         .count();
54     while j < size {
55         b58_str.push(ALPHABET[buffer[j] as usize] as char);
56         j += 1;
57     }
58
59     // Return the encoded string
60     b58_str
61 }
62 }

```

Decoding is the process of converting Base58 encoded data back to its original form, which is essentially a conversion between numeral systems. The specific implementation of the decoding method is as follows:

```

1 // base58.rs
2
3 // Implement Base58 decoding
4 impl Decoder for str {
5     fn decode_from_base58(&self) -> Result<String, DecodeError>
6     {
7         // Store conversion characters
8         let mut bin = [0u8; 132];
9         let mut out = [0u32; (132 + 3) / 4];

```



```

10
11     // Number of remaining bits after processing data
12     // in units of 4
13     let bytes_left = (bin.len() % 4) as u8;
14     let zero_mask = match bytes_left {
15         0 => 0u32,
16         _ => 0xffffffff << (bytes_left * 8),
17     };
18
19     // Count leading zeros
20     let zero_count = self.chars()
21         .take_while(|&x| x == ALPHABET_INDEX_0)
22         .count();
23
24     let mut i = zero_count;
25     let b58: Vec<u8> = self.bytes().collect();
26     while i < self.len() {
27         // Invalid characters
28         if (b58[i] & 0x80) != 0 {
29             return Err(DecodeError::InvalidCharacter(
30                 b58[i] as char, i));
31         }
32         if DIGITS_MAP[b58[i] as usize] == 255 {
33             return Err(DecodeError::InvalidCharacter(
34                 b58[i] as char, i));
35         }
36
37         // Number system conversion
38         let mut j = out.len();
39         let mut c = DIGITS_MAP[b58[i] as usize] as u64;
40         while j != 0 {
41             j -= 1;
42             let t = out[j] as u64 * (BIG_RADIX as u64) + c;
43             c = (t & 0x3f00000000) >> 32;
44             out[j] = (t & 0xffffffff) as u32;
45         }
46
47         // Data is too long
48         if c != 0 {
49             return Err(DecodeError::InvalidLength);
50         }
51
52         if (out[0] & zero_mask) != 0 {
53             return Err(DecodeError::InvalidLength);
54         }
55
56         i += 1;
57     }
58
59     // Handle remaining bits
60     let mut i = 1;
61     let mut j = 0;

```

```

62     bin[0] = match bytes_left {
63         3 => ((out[0] & 0xff0000) >> 16) as u8,
64         2 => ((out[0] & 0xff00) >> 8) as u8,
65         1 => {
66             j = 1;
67             (out[0] & 0xff) as u8
68         },
69         _ => {
70             i = 0;
71             bin[0]
72         }
73     };
74
75     // Process data in units of 4 and
76     // use bit shifting for division
77     while j < out.len() {
78         bin[i] = ((out[j] >> 0x18) & 0xff) as u8;
79         bin[i + 1] = ((out[j] >> 0x10) & 0xff) as u8;
80         bin[i + 2] = ((out[j] >> 8) & 0xff) as u8;
81         bin[i + 3] = ((out[j] >> 0) & 0xff) as u8;
82         i += 4;
83         j += 1;
84     }
85
86     // Get number of leading zeros
87     let leading_zeros = bin.iter()
88         .take_while(|&x| x == 0)
89         .count();
90
91     // Get decoded string
92     let new_str = String::from_utf8(
93         bin[leading_zeros - zero_count..]
94         .to_vec());
95
96     // Return valid result
97     match new_str {
98         Ok(res) => Ok(res),
99         Err(_) => Err(DecodeError::Invalid),
100     }
101 }
102 }
103
104 fn main() {
105     println!("{:?}", "abc".encode_to_base58());
106     println!("{:?}", "ZiCa".decode_from_base58().unwrap());
107
108     println!("{:?}", "loverust".encode_to_base58());
109     println!("{:?}", "K8xdoM2VJtK".decode_from_base58());
110
111     println!("{:?}", "iloveyou".encode_to_base58());
112     println!("{:?}", "Jdjxs3pmuK2".decode_from_base58());
113 }

```

The following are the results after Base58 encoding and decoding.

```
"ZiCa"  
"abc"  
"K8xdoM2VJtK"  
Ok(  
    "loverust",  
)  
"Jdjsx3pmuK2"  
Ok(  
    "iloveyou",  
)
```

With the completion of the entire Base58 algorithm, similar methods can be used to implement encoding and decoding algorithms for other encoding schemes of interest, such as Base32, Base36, Base62, Base64, Base85, Base92, and so on. In Chapter 1, we used Base64 to create a password generator, but it can also be replaced with Base58. I have already made the replacement and included the specific code in the corresponding repository for this chapter.

10.8 Blockchain

Blockchain is a digital technology that has gained significant attention in recent years. This attention has been due, in part, to the rise in the price of Bitcoin, which is closely tied to blockchain technology, as well as other related concepts such as Ethereum, virtual currencies, and the digital economy. The endorsement of virtual currencies by influential figures such as Tesla CEO Elon Musk has further fueled the development of the field. Additionally, governments around the world have been formulating policies related to blockchain technology, adding to the momentum of the industry. Ultimately, blockchain technology is a tool closely tied to economic development and the trade of commodities.

Online transactions rely heavily on financial institutions as third-party intermediaries to process payment information. However, this credit-based model has inherent weaknesses that limit the feasibility of small daily payment transactions, increase the cost of transactions, and prevent completely irreversible transactions due to the involvement of financial intermediaries to coordinate disputes. This limits the potential of internet trade as both parties in a transaction must trust each other, and merchants must guard against potential fraud. In contrast, physical cash transactions do not require third-party intermediaries and thus do not face such limitations.

To address these issues, we need an electronic payment system based on cryptographic principles that enables two parties to make payments directly without intermediaries. This eliminates the possibility of rolling back payment transactions, protecting sellers from fraud. We propose a method for generating electronic transaction proofs using a peer-to-peer distributed timestamp server that records transactions in chronological order. The system is secure as long as the total computing power controlled by honest nodes exceeds that of cooperating attackers.

The passage above is the introduction to the Bitcoin whitepaper (Bitcoin: A Peer-to-Peer Electronic Cash System^[19]), written by Satoshi Nakamoto, the inventor of Bitcoin. It outlines the reason for the invention of Bitcoin, which was to address the practical problem of the global financial crisis of 2008, inflation, and its impact on various countries. Nakamoto was dissatisfied with the existing financial environment and used their professional knowledge to invent the blockchain technology to address this problem.

10.8.1 The Principles of Blockchain and Bitcoin

Blockchain and Bitcoin are often mentioned together in media reports, but their relationship is not always clear. Blockchain technology is a new decentralized infrastructure and computing paradigm that

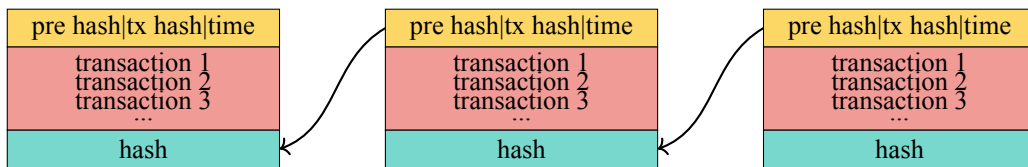
uses a chain data structure to store and verify data, distributed node consensus algorithms to generate and update data, cryptography to ensure data security, and smart contracts to program and operate data. In contrast, Bitcoin is a type of digital currency that was invented using blockchain technology.

To put it simply, blockchain is a decentralized distributed ledger, where data is not only stored on each node but replicated and shared across the entire network. Blockchain is a ledger that records transactions and consumes a lot of resources to do so. As transactions are recorded, rewards and transaction fees are generated in the form of digital currency, which is used to maintain the system's operation. Bitcoin is the first digital currency created using blockchain technology and serves as a guarantee of transactions and a type of incentive.

In summary, blockchain is a distributed transaction medium, and Bitcoin is a type of digital currency that serves as a guarantee of transactions and an incentive. Blockchain itself consists of components such as blocks, blockchains, transactions, accounts, miners, transaction fees, and rewards. To implement a blockchain system, one must start by implementing these basic components.

10.8.2 A Primary Blockchain

The blockchain structure consists of a block header, block body, and block hash. The header contains the hash of the previous block (pre_hash), the current block transaction hash value (tx_hash), and the block packaging time (time). The block body includes all transaction data (transactions), and the block hash (hash) is calculated from the block header and block body. The block and blockchain structures are shown in the following figure.



Hash values play a vital role in the blockchain, as evident from the structure discussed above. Thus, the initial step is to compute the hash value, and serialization of the block structure is a more efficient approach.

The basic blockchain's first function is to perform serialization and hash value computation, demonstrated in the following code. To handle blocks with varying transaction numbers, we employ the "Sized" trait. The serialization process utilizes the "bincode" library, while the "crypto's Sha3" library computes the hash. For better readability, we convert all hashes to strings. The serialized data is of type "&[u8]", and "hash_str" retrieves the data and returns it as a string.

```

1 // serializer.rs
2
3 use bincode;
4 use serde::Serialize;
5 use crypto::digest::Digest;
6 use crypto::sha3::Sha3;
7
8 // Data serialization
9 pub fn serialize<T: ?Sized>(value: &T) -> Vec<u8>
10     where T: Serialize
11 {
12     bincode::serialize(value).unwrap()
13 }
14
15 // calculate the hash value and return it as a string
16 pub fn hash_str(value: &[u8]) -> String {
  
```

```

17     let mut hasher = Sha3::sha3_256();
18     hasher.input(value);
19
20     hasher.result_str()
21 }

```

The hash computation function can calculate the hash values for the block's hash, `pre_hash`, and `tx_hash`, while the time can be obtained by capturing the block's generation time. Initially, we simplify the problem by using strings to represent transactions and place them in a `Vec` to simulate multiple transactions. In Rust, we can utilize the `struct` to represent blocks and block headers.

```

1 // block.rs
2
3 // Block structure
4 pub struct Block {
5     pub header: BlockHeader,
6     pub tranxs: String,
7     pub hash: String,
8 }
9
10 // Block header structure
11 pub struct BlockHeader {
12     pub time: i64,
13     pub pre_hash: String,
14     pub txs_hash: String,
15 }

```

Since hash values are crucial in the blockchain, the first step is to calculate the hash. To achieve this, we serialize the block structure before computing the hash value for efficiency purposes.

```

1 // block.rs
2
3 use std::thread;
4 use std::time::Duration;
5 use chrono::prelude::*;
6 use utils::serializer::{serialize, hash_str};
7 use serde::Serialize;
8
9 // Definition of block header
10 #[derive(Serialize, Debug, PartialEq, Eq)]
11 pub struct BlockHeader {
12     pub time: i64,
13     pub pre_hash: String,
14     pub txs_hash: String,
15 }
16
17 // Definition of block
18 #[derive(Debug)]
19 pub struct Block {
20     pub header: BlockHeader,
21     pub tranxs: String,
22     pub hash: String,
23 }

```

```

24
25 impl Block {
26     pub fn new(txs: String, pre_hash: String) -> Self {
27         // Introduce a 3-second delay for mining
28         println!("Start mining .... ");
29         thread::sleep(Duration::from_secs(3));
30
31         // Prepare timestamp and calculate transaction hash
32         let time = Utc::now().timestamp();
33         let txs_ser = serialize(&txs);
34         let txs_hash = hash_str(&txs_ser);
35
36         let mut block = Block {
37             header: BlockHeader {
38                 time: time,
39                 txs_hash: txs_hash,
40                 pre_hash: pre_hash,
41             },
42             tranxs: txs,
43             hash: "".to_string(),
44         };
45
46         block.set_hash();
47         println!("Produce a new block!\n");
48
49         block
50     }
51
52     // Calculate and set block hash value
53     fn set_hash(&mut self) {
54         let header = serialize(&(self.header));
55         self.hash = hash_str(&header);
56     }
57 }

```

To form a blockchain, we must link the created blocks together, which can be accomplished using a `Vec` to store multiple blocks. The system should enable the generation of the first block (genesis block) and adding new blocks to the chain. Since the genesis block does not have a `pre_hash`, it needs to be set manually. We will utilize the base64 value of "Bitcoin hit \$60000" as the `pre_hash` for the genesis block. Lastly, we will require a function to create a new block and update its hash value.

```

1 // blockchain.rs
2 use crate::block::Block;
3
4 // Genesis block pre_hash
5 const PRE_HASH: &str = "22caaf24ef0aea3522c13d133912d2b7
6                        22caaf24ef0aea3522c13d133912d2b7";
7 pub struct Blockchain {
8     pub blocks: Vec<Block>,
9 }
10
11 impl Blockchain {
12     pub fn new() -> Self {

```

```

13         Blockchain { blocks: vec![Self::genesis_block()] }
14     }
15
16     // Generate the genesis block
17     fn genesis_block() -> Block {
18         Block::new("genesis block".to_string(), PRE_HASH.
19             to_string())
20     }
21
22     // Add blocks to form a blockchain
23     pub fn add_block(&mut self, data: String) {
24         // Get the hash value of the previous block
25         let pre_block = &self.blocks[self.blocks.len() - 1];
26         let pre_hash = pre_block.hash.clone();
27         // Build a new block and add it to the chain
28         let new_block = Block::new(data, pre_hash);
29         self.blocks.push(new_block);
30     }
31
32     // Print block information
33     pub fn block_info(&self) {
34         for b in self.blocks.iter() { println!("{:?}", b); }
35     }

```

To execute a blockchain system, we must generate transactions to create blocks. In the code below, the string "tx" represents a transaction, and information is printed after the transaction has been packed and finalized.

```

1 // main.rs
2 use core::blockchain::Blockchain as BC;
3
4 fn main() {
5     println!("-----Mine Info-----");
6
7     let mut bc = BC::new();
8     let tx = "0xabcd -> 0xabce: 5 btc".to_string();
9     bc.add_block(tx);
10    let tx = "0xabcd -> 0xabcf: 2.5 btc".to_string();
11    bc.add_block(String::from(tx));
12
13    println!("-----Block Info-----");
14    bc.block_info();
15 }

```

To ensure that these codes work coherently, we need to arrange them logically. Since the hash calculation is not directly related to the blockchain, it should be included in the "utils" directory as a tool. On the other hand, the "block" and "blockchain" should be stored in the "core" directory. The main function should call the "core" directory to execute the creation and addition of blocks. To generate the blockchain project, we can use Cargo, as demonstrated in the [blockchain1](#) repository on GitHub.

The code above has facilitated the implementation of a basic blockchain project that can create and add blocks. The program's output, including mining information and block information, is depicted on next page.

```

-----Mine Info-----
Start mining ... Produce a new block!

Start mining ... Produce a new block!

Start mining ... Produce a new block!

-----Block Info-----
Block {
  header: BlockHeader {
    time: 1679965388,
    txs_hash: "f056286693df436978f73e333ab580220e05782b2eadedea8e84dc0c4d4af408" ,
    pre_hash: "22caaf24ef0aea3522c13d133912d2b722caaf24ef0aea3522c13d133912d2b7" ,
  },
  tranxs: "genesis block" ,
  hash: "55d057cbcc55dd468b093004fb94f51581ed3d6b315c50cf545c9d28bce256d5" ,
}
Block {
  header: BlockHeader {
    time: 1679965391,
    txs_hash: "84eeeb7be34240b4a5c45534fc0951ec8f375d93cd3a77d2f154fbdfdde080c1" ,
    pre_hash: "55d057cbcc55dd468b093004fb94f51581ed3d6b315c50cf545c9d28bce256d5" ,
  },
  tranxs: "0xabcd -> 0xabce: 5 btc" ,
  hash: "0c1c5383bb5844fa17f59d55a287def80ae0f887c5940acf1ad93948fe4010cb" ,
}
Block {
  header: BlockHeader {
    time: 1679965394,
    txs_hash: "ca51ee57941a2af26ae2fa02d05f6276f6c2c050535314d6393501b797b13438" ,
    pre_hash: "0c1c5383bb5844fa17f59d55a287def80ae0f887c5940acf1ad93948fe4010cb" ,
  },
  tranxs: "0xabcd -> 0xabcf: 2.5 btc" ,
  hash: "ed9ca0f1474f7e9d18aa3e4fc586b605350d6dc5b9c5d37b6285d828d72f6e8f" ,
}

```

Our implementation is a simplified version of a blockchain. A comprehensive blockchain involves various components such as proof of work, transactions, accounts, hashing, miners, mining, bitcoin rewards, block storage, and many others. Readers can gradually incorporate these features on top of the fundamental blockchain implementation.

10.9 Summary

In this chapter, we covered a plethora of useful data structures. We began by exploring the implementation of trie, Bloom filter, and cuckoo filter. Next, we delved into Hamming and edit distance, followed by an understanding of LRU cache eviction algorithm and consistent hashing algorithm. Lastly, we learned about the principles of blockchain and implemented a rudimentary blockchain.

Throughout this book, we have covered a wide range of data structures and implemented a considerable amount of Rust code. Although some of the codes may not be optimal, they can still be useful references for readers. Any feedback from readers via Github to further improve the content is greatly appreciated.

Finally, I hope that this book proves helpful to readers and serves as a valuable resource to explore rust and its applications in data structures.

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