

Computer Science



Lecture #1

? ? ~~COVID~~ EDITION
?

Professor: Carey Nachenberg

E-mail: climberkip@gmail.com

Class details: Franz 1178 - M/W 10am-12pm

Office hours: Eng VI, 372 (and via Zoom initially)

Mondays 12pm-1pm

Wednesdays 9am-10am

My Office: Eng VI, 297

The Question on Everyone's Mind!

What is the airspeed velocity of an unladen swallow?

When will we be
back in person?



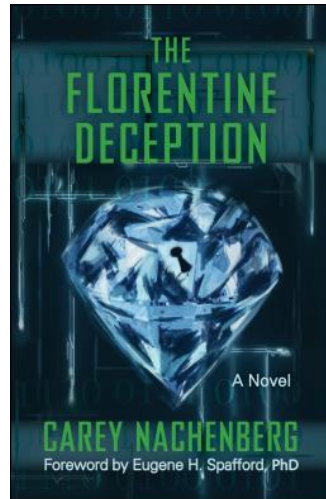
There's been no change - we're still supposed to be back
in-person after two weeks.

Don't worry - I'll tell you the moment I hear any news!

Who Am I?



299934



Carey Nachenberg

Age: 50

School: BS+MS in CS, UCLA '95

Work: Adjunct Prof at UCLA

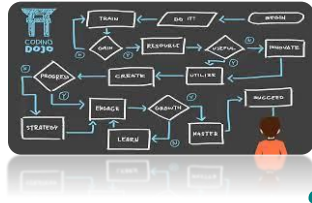
Hobbies: Rock climbing,
weight training,
teaching CS!

My goal: To make an impact on
your life (through
getting you excited
about programming)!

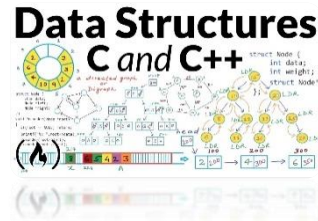
What You'll Learn in CS32



Advanced C++ Topics
Object Oriented Programming
and C++ language features



Algorithms
The most useful
algorithms (e.g.,
sorting, searching)



Data Structures
The most useful data
structures (e.g., lists,
trees, hash tables)



Building Big Programs
How to write large
programs (1000+ lines)

Basically, once you complete CS32,
you'll know 95% of what you need to
succeed in industry!



Official Class Websites

<http://www.cs.ucla.edu/classes/winter22/cs32/>

The class website has the **syllabus**, **grading policy**, **academic integrity agreement** and **assignments/deadlines**.

Bruin Learn Website

While we're virtual, I will post **recordings of all lectures** on Bruin Learn.

Warning: I will not always announce homework/projects so you have to track this on your own and be on top of the trash*!

* Being on top of the trash means being **responsible****.



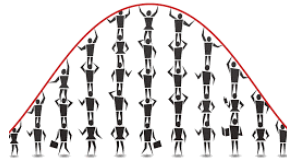
** If you're not responsible, you could get **rekt**.

The Syllabus...

I know you won't read it... but...



Academic
integrity
agreement



Curving and
grading policy



Late
submission
policy



TA Office
hours



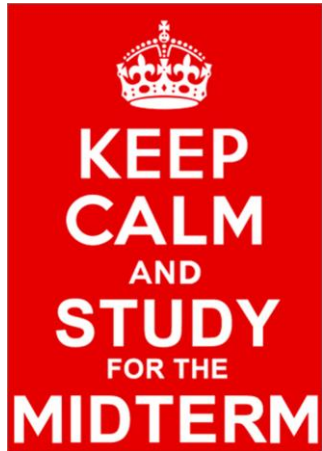
Testing your
code w/ 2
different
compilers



Deadlines for
all hw/projects

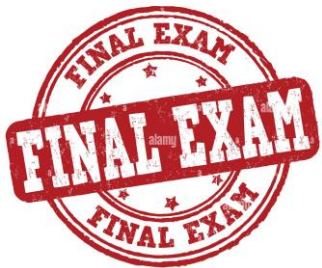
Important Dates

Project #1: Due Wed, Jan 12th (next Wed!)



Midterm #1: Tues, Feb 1st
6pm-7:30pm (not during class hours!!)

Midterm #2: Tues, Feb 22nd
6pm-7:30pm (not during class hours!!)



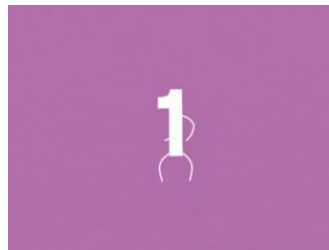
Final Exam: Sat, March 12th
11:30am-2:30pm

(This is the Saturday **BEFORE** Finals Week. Don't forget!)

Project #1: Due **NEXT WED!**

In P1, you have to add a few features to a **C++ program** that we provide.

It's worth exactly



% of your total grade!

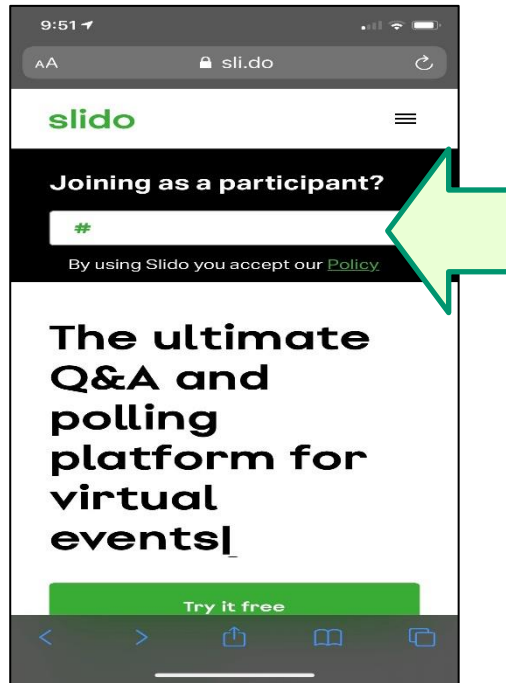
It's mostly for you to evaluate how ready you are for CS32.



If you find it difficult, I strongly encourage you to do a bit more prep and take the class in spring. 😊

Making Virtual Class Tolerable

(Until we're back in person!)



We'll be using **Slido** for questions!

On your laptop/phone, go to:

<https://sli.do>

Use event code: **UCLA-CS32**

You can ask and upvote questions,
all anonymously.

I'll also be doing surveys to make sure
everyone is following along.

To make things more
interesting, we'll have
5 min "detox" activities
from time to time...





Participate for Prizes!

(Who says bribery doesn't work)

I'll be giving one **\$20 Amazon Gift Card** out after each **virtual lecture** to reward participation.



If you **answer one of my questions** or **ask a relevant question**, you'll be entered in a raffle each day of class!

Carey's Thoughts on Teaching



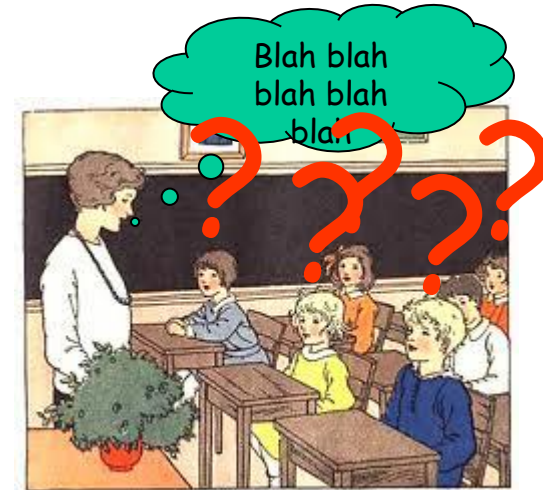
It's more important that **everyone understand** a topic than I **finish a lecture on time**.

Don't be shy!!!

If something **confuses you...**

it probably **confuses 5 other people** too.

So **always ask questions** if you're confused!



Always save more **advanced questions** for office hours or break.

I reserve the right to **wait until office hours** to answer advanced questions.



Questions? Bring Em On!

Alright, bring on those questions!



Obfuscated C Programming Contest: What Does it Do?

Answer: A maze solver! Given a file, it will find a path from the @ to the exclamation point!

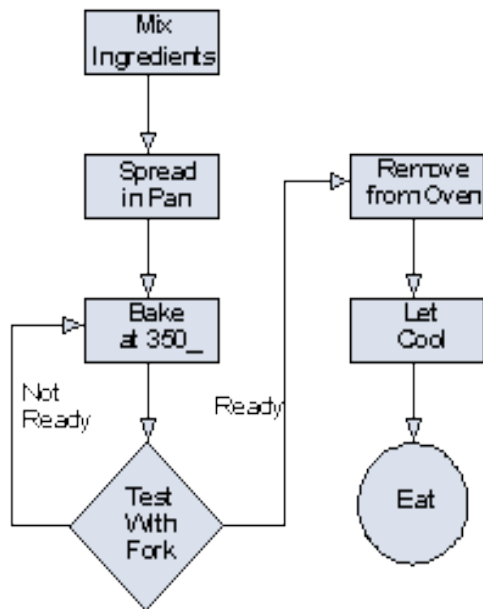
[illegible]

Credit: Joshua Karns

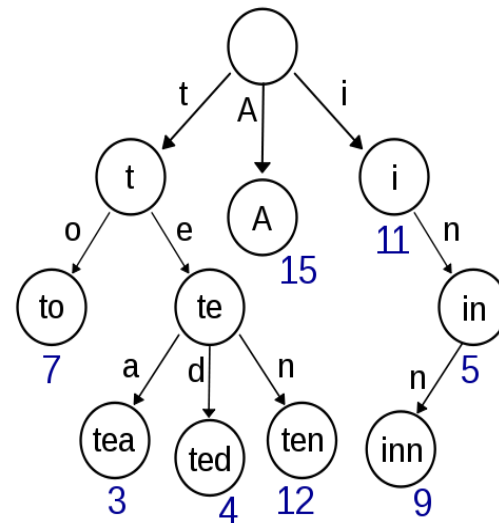
Alright... Enough administration!

Let's learn about...

Algorithms



Data Structures



What is an Algorithm



An **algorithm** is a set of **instructions/steps** that solves a particular problem.

Each algorithm operates on **input data**.



42

Each algorithm produces an **output result**.

Each algorithm can be classified by **how long it takes to run** on a particular input.



Each algorithm can be classified by **the quality of its results**.

Algorithm Comparison

"Guess My Word"

Let's say you're building a word guessing game.

The user secretly
picks a random word
from the dictionary.



Our program then must
discover the word the user
picked as quickly as possible.

Let's consider **two** different algorithms...

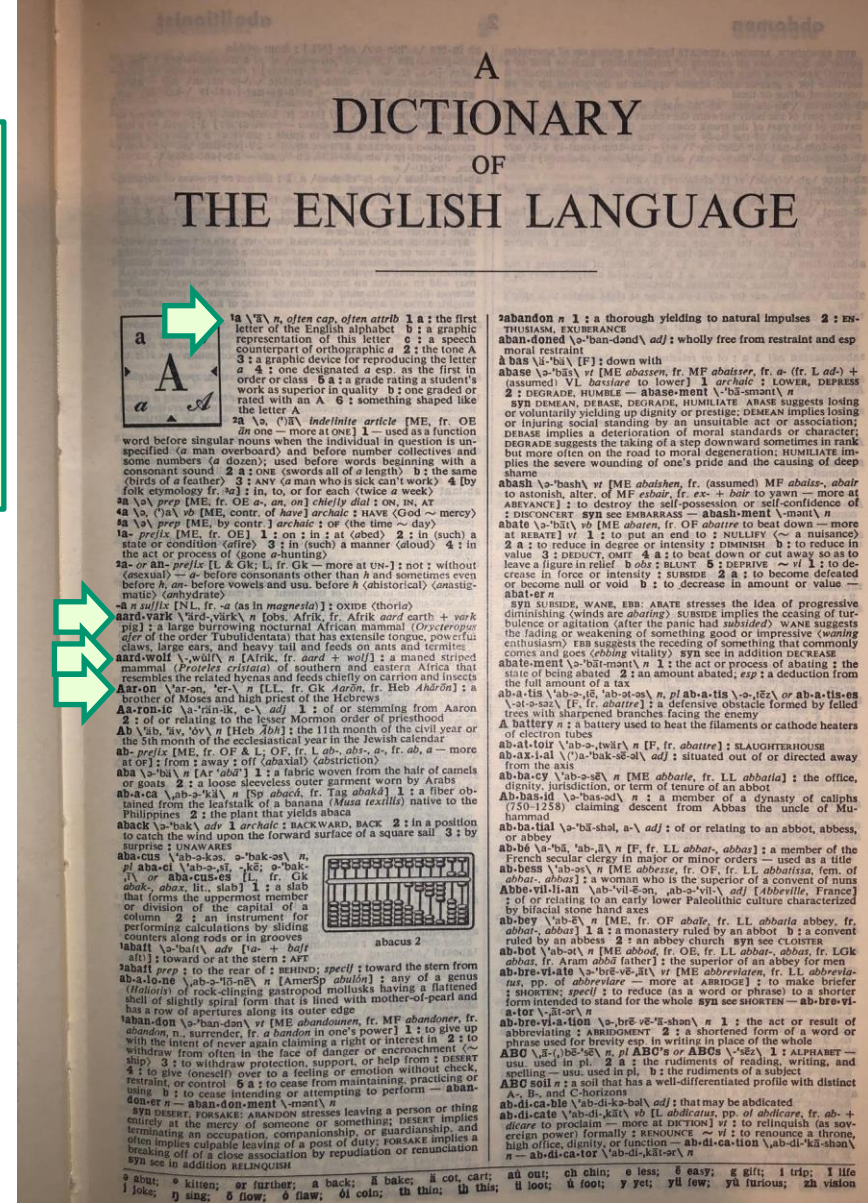
Algorithm #1: Linear Search

Let's try a simple algorithm...

1. Ask the user to think of a random word
2. Start at the first word in the dictionary
3. While we haven't found the user's word:
 - a. Ask: "Is that your word?"
 - b. If not, advance to the next word

Question: If there are 100,000 total words in our dictionary, on average, how many guesses will our algorithm require?

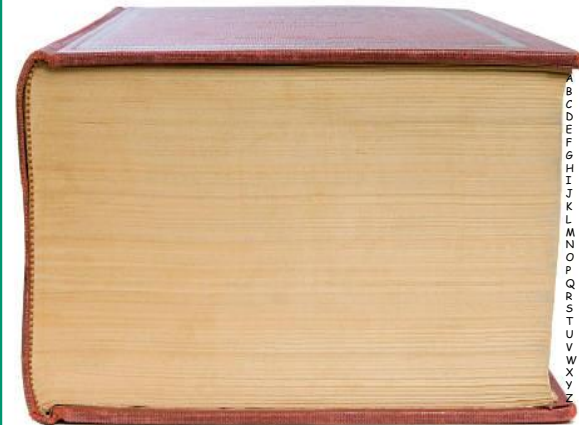
Ok, let's try it.



Algorithm #2: Binary Search

Alright, for our second strategy let's try a more intelligent approach called **binary search**:

1. Print: "Think of a random word"
2. While we haven't found the word:
 - a. Pick the middle word in the dictionary
 - b. Ask: "Does your word come before or after that word?"
 - c. If their word comes before, throw away the 2nd half of the dictionary
 - d. If their word comes after, throw away the 1st half of the dictionary



See how it works? If the user says their word comes before the middle word in the dictionary, we can immediately rule out the second half of the dictionary! We know the user's word must be in the first half of the dictionary. Then we can repeat this process for the remaining first half of the dictionary, and so on. Eventually we'll arrive at a single word, which **MUST** be the user's word!

Question: If there are **100,000 total words** in our dictionary, on average, how many guesses will our Binary Search algorithm require? [see next slide]

Binary Search: How Many Guesses?

We keep on dividing our search area in half until we finally arrive at our word.

In the worst case, we must keep halving our search area until it contains just a single word – our word!



If our dictionary had 16 words, how many times would we need to halve it until we have just one word left?

16 8 4 2 1

It would take 4 steps

Ok, what if our dictionary had 131,072 words?

131072 65536 32768 16384 8192 4096 2048 1024 512 256 128 64 32 16 8 4 2 1

It would take just 17 steps!!! WOW!

Wow! That's Significant!

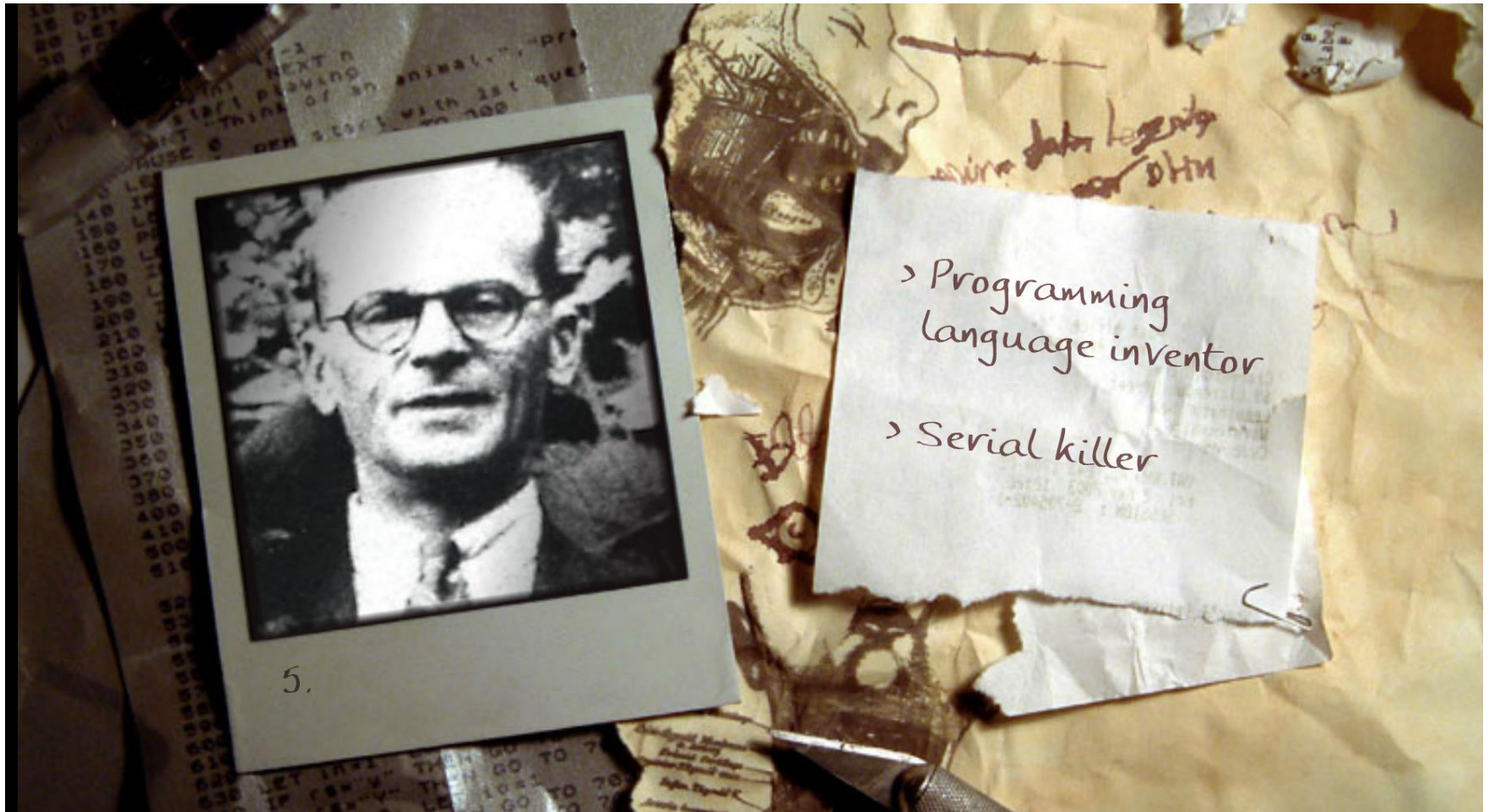
Linear search requires ~50,000 steps to guess a word from a 100k word dictionary.

But our binary search algorithm requires just 17 steps, on average, to guess the user's secret word!



In CS32, you'll learn:
all the major algorithms,
how to analyze them, and
how to pick the best one


And now for a fun game!



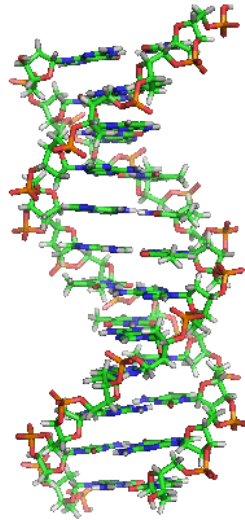
Is this guy a **programming language inventor**... or a **SERIAL KILLER?!?!?**

Killed 8 women in London, was arrested after new tenants tracing an unpleasant odor peeled off the kitchen wallpaper to reveal a corpse.

Data Structures

 A **data structure** is the **set of variable(s)** that an algorithm uses to solve a problem.

Let's consider a data structure to efficiently store **millions of DNA sequences** so they can be searched.



DNA is made up of four bases:
Adenine, **Cytosine**, **Guanine** and **Thymine**

A Data Structure For DNA

If I wanted to store millions of bacterial DNA sequences, each with 10 bases:

```
atggacatct  
acattaacga  
caccctcacc  
accgtagaat  
caccgccagct  
caccgaaatt
```

...

What **simple data structure** might I use?

A Data Structure For DNA

Right! You could use a **sorted array** of strings!

```
string dna_seq[100000000];
```

```
dna_seq
```

```
[0] acattaacga
```

```
[1] accgtagaat
```

```
[2] atggacatct
```

```
[3] caccgccagct
```

```
[4] caccctcacc
```

```
[5] caccgaaatt
```


```
... ..
```

Not bad. Given a new sequence, I could use **binary search** to determine if its is held in our data structure.

A Data Structure For DNA

Are there any **drawbacks** of the array data structure?

```
dna_seq
[0] acattaacga
[1] accgtagaat
[2] atggacatct
[3] caccgccagct
[4] caccctcacc
[5] caccgaaatt
... ..
```



atgaacatct

Right! If we wanted to add **even one new sequence**, we'd have to **move millions of items**, and that'd be slow.

A Data Structure For DNA

Are there any other **drawbacks** of the array data structure?

```
dna_seq
[0] acattaacga
[1] accgtagaat
[2] atggacatct
[3] caccgccagct
[4] caccctcacc
[5] caccgaaatt
... ..
```

Right! **Arrays** have a **fixed capacity**, so we're limited in how many DNA sequences we can add!

A Better Data Structure for DNA

Instead of an array, let's use a **Trie** data structure.

A Trie is just like a **balanced mobile**.



Let's see!

A Trie for DNA

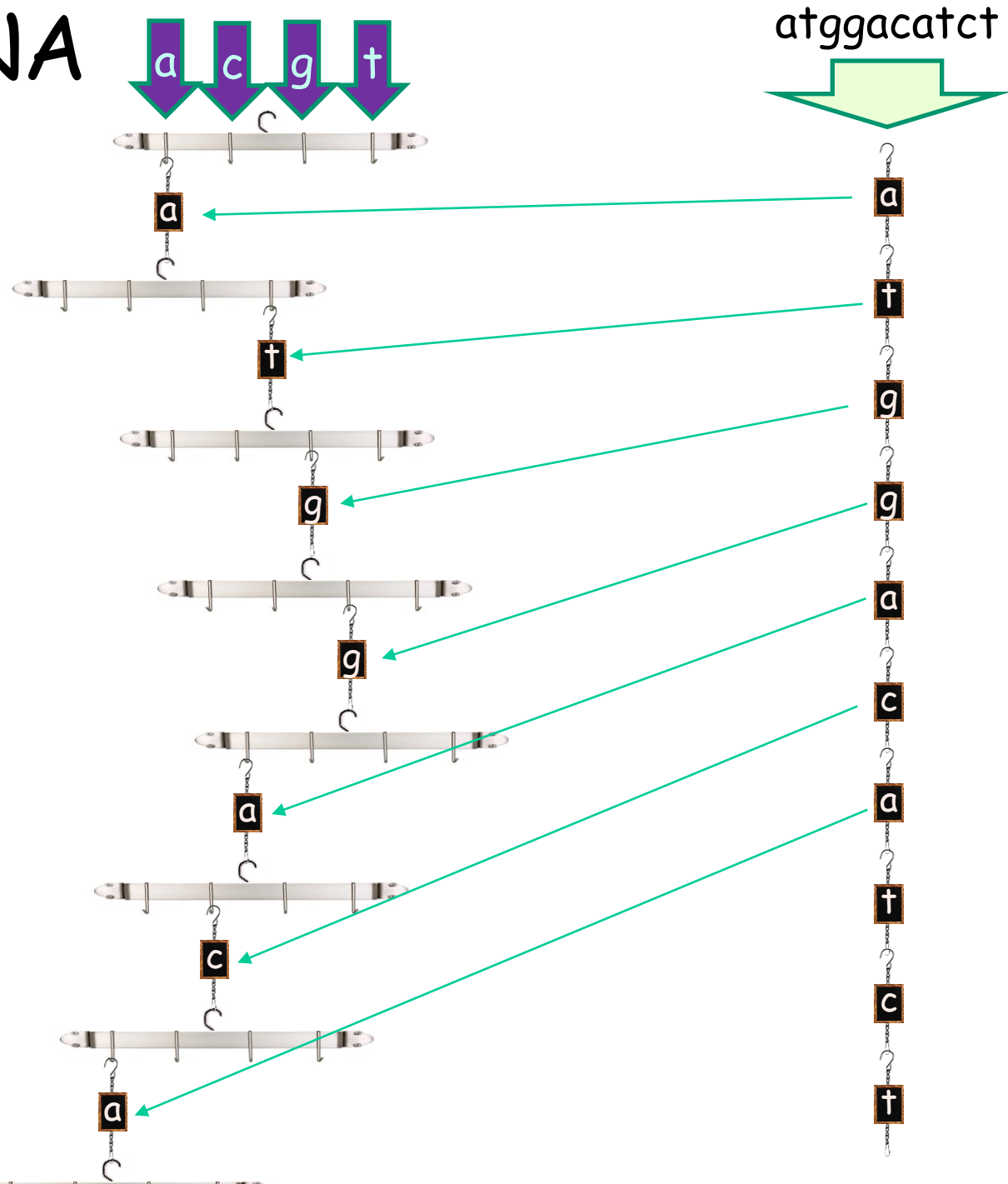
Imagine each DNA sequence dangling from ten linked chains.

Now imagine a hanging bar with **four hooks**, one hook for **A**, one for **C**, one for **G** and one for **T**.

Starting with the first base of the sequence, **hang its chain** from the proper hook...

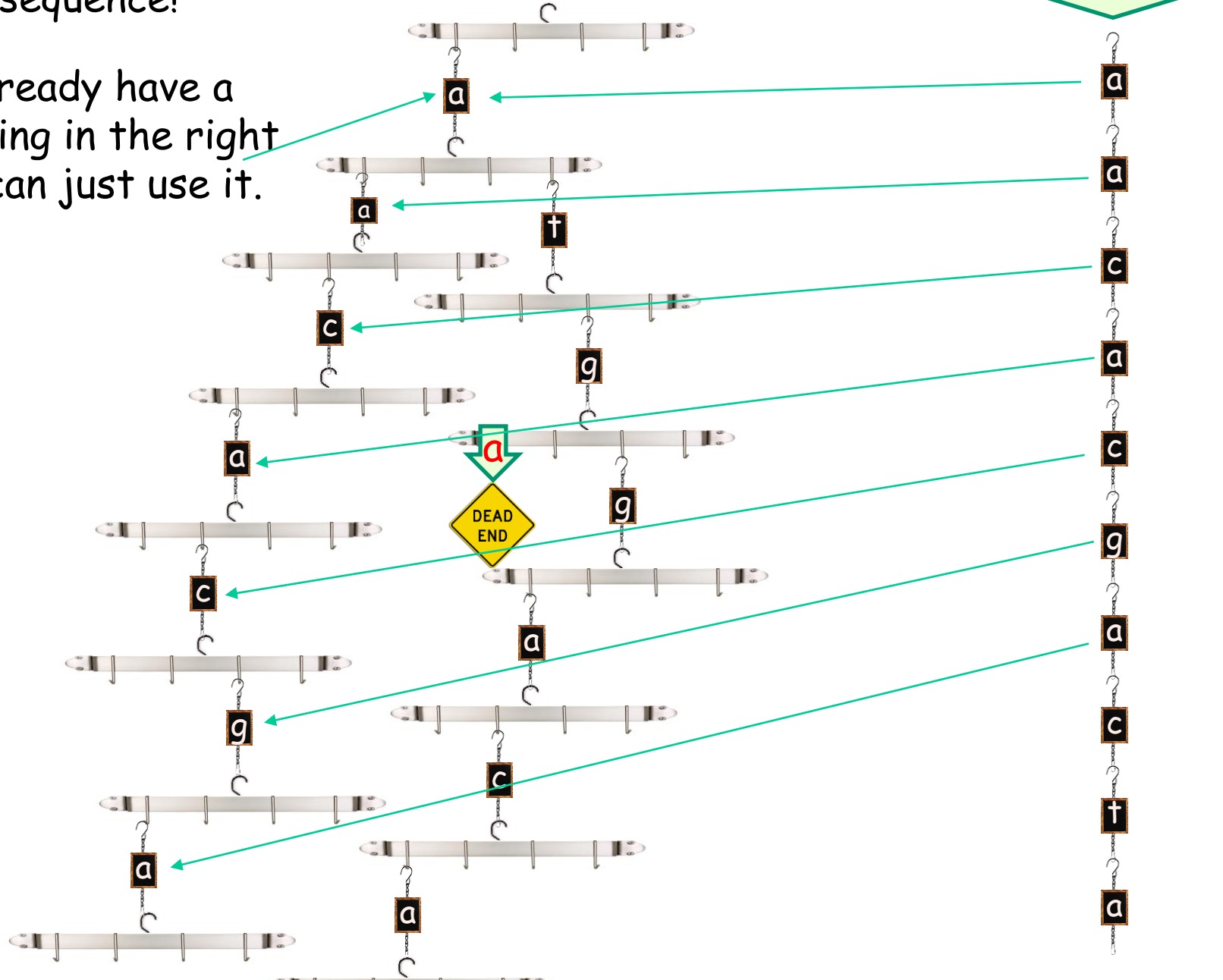
Add a new bar under the chain.

Repeat for all bases.



Ok, let's add a second DNA sequence!

If we already have a chain hanging in the right spot, we can just use it.



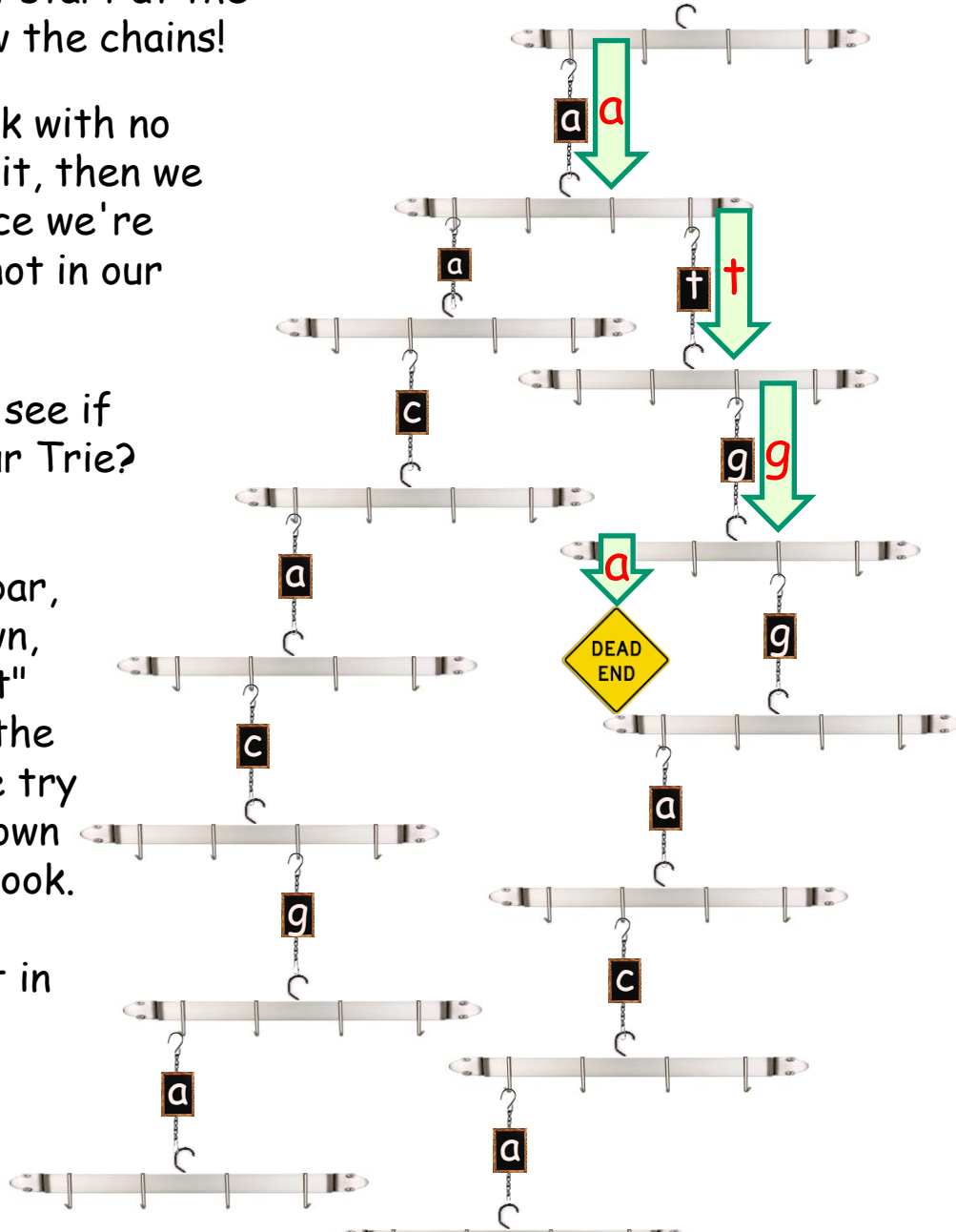
Now, to search if a sequence is in our Trie, we just start at the top bar and follow the chains!

If we reach a hook with no chain hanging from it, then we know the sequence we're searching for is not in our Trie!

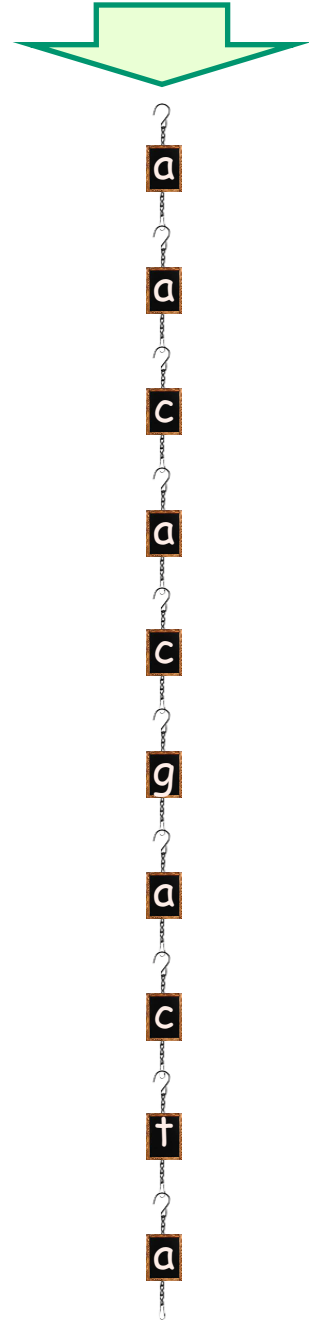
For instance, let's see if "atgattacgt" is in our Trie?

We start the top bar,
follow the "a" down,
then follow the "t"
down, then follow the
"g" down. Finally, we try
to follow the "a" down
and find an empty hook.

The sequence
"atgattacgt" is not in
our Trie!



aacacgacta

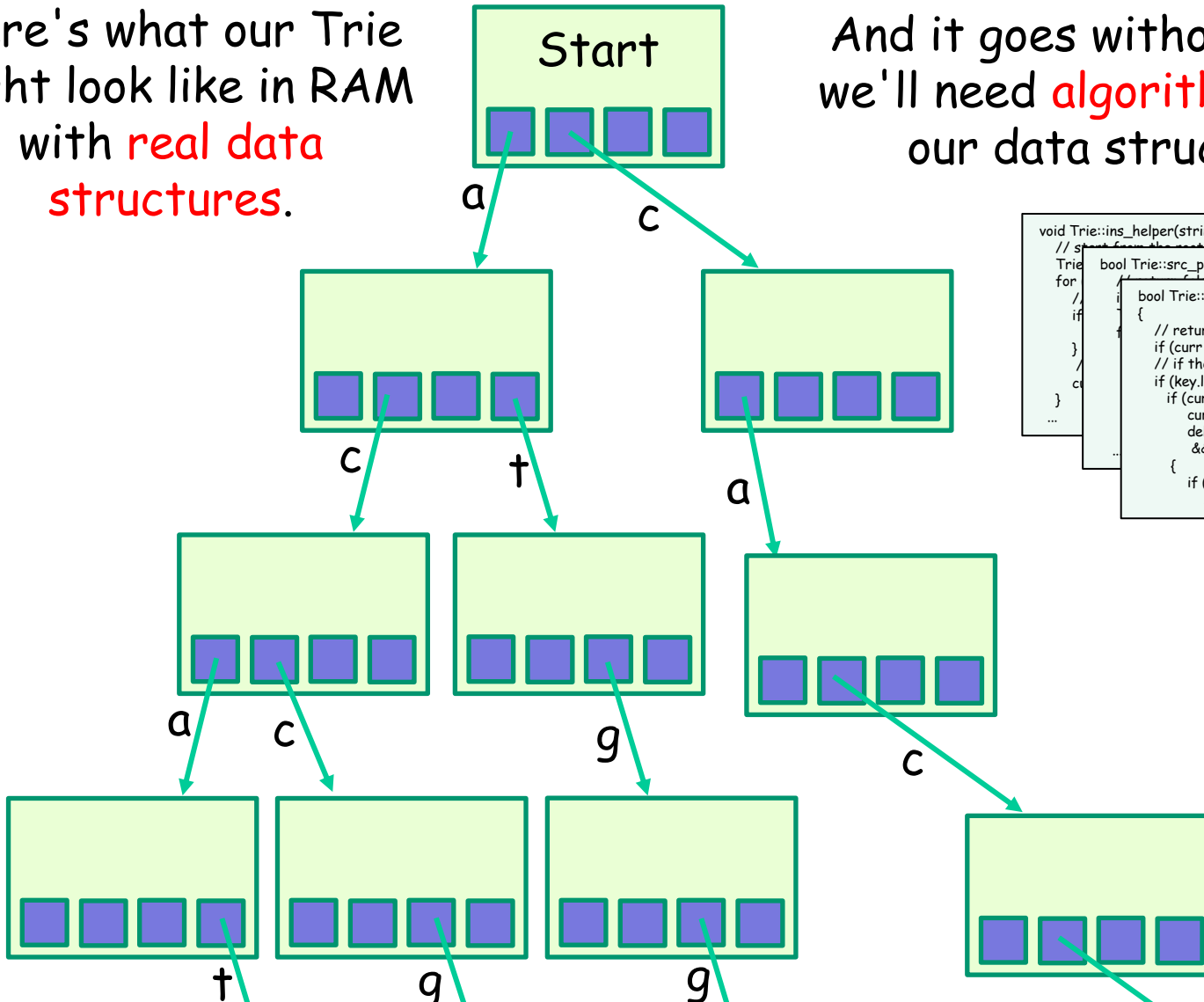


A Trie for DNA

Enough analogies with chains!

Here's what our Trie
might look like in RAM
with **real data**
structures.

And it goes without saying that we'll need **algorithms** to process our data structure too...



```
void Trie::ins_helper(string key) {
    // ...
    bool Trie::src_primitive(string key) {
        // ...
        bool Trie::del_nodes(Trie*& curr, string key)
        {
            // return if Trie is empty
            if (curr == nullptr) { return false; }
            // if the end of the key is not reached
            if (key.length() > 0) {
                if (curr != nullptr &&
                    curr->character[key[0]] != nullptr &&
                    deletion(curr->character[key[0]], key.substr(1))
                    && curr->isLeaf == false)
                {
                    if (!haveChildren(curr))
                        ...
                }
            }
        }
    }
}
// ...
}
```

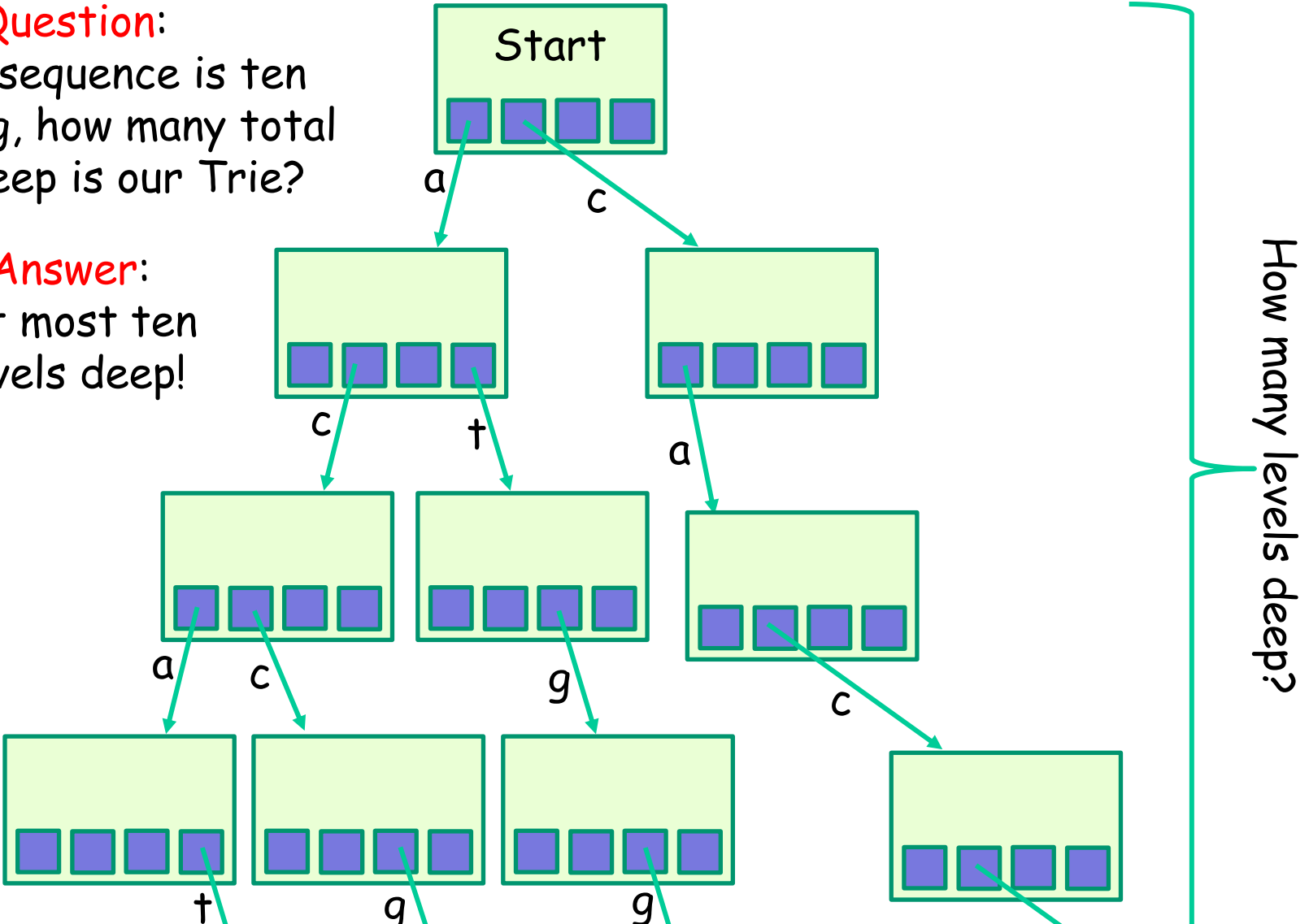
A Trie for DNA

Question:

If each sequence is ten bases long, how many total levels deep is our Trie?

Answer:

At most ten levels deep!



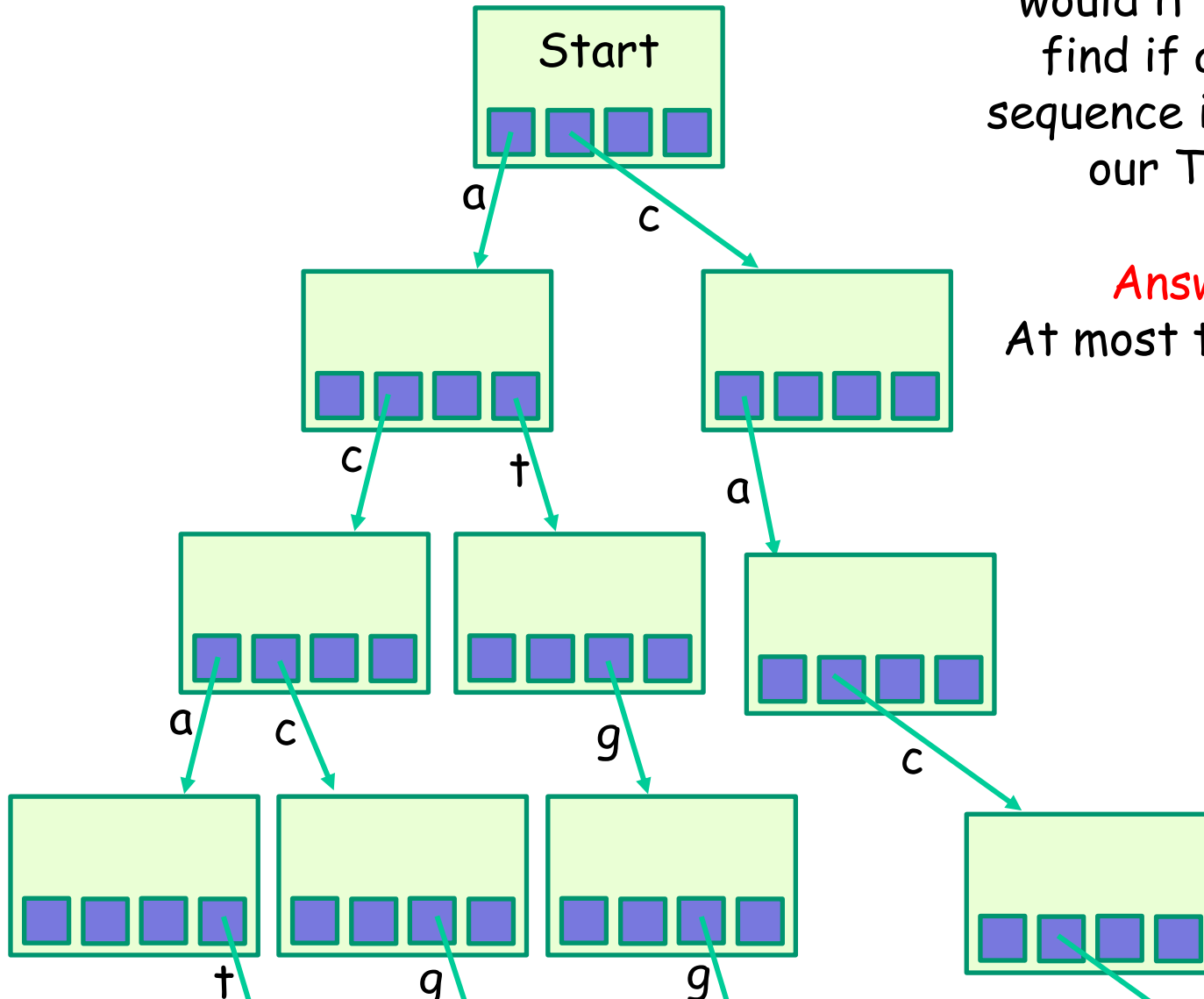
A Trie for DNA

Question:

How many steps
would it take to
find if a DNA
sequence is held in
our Trie?

Answer:

At most ten steps!



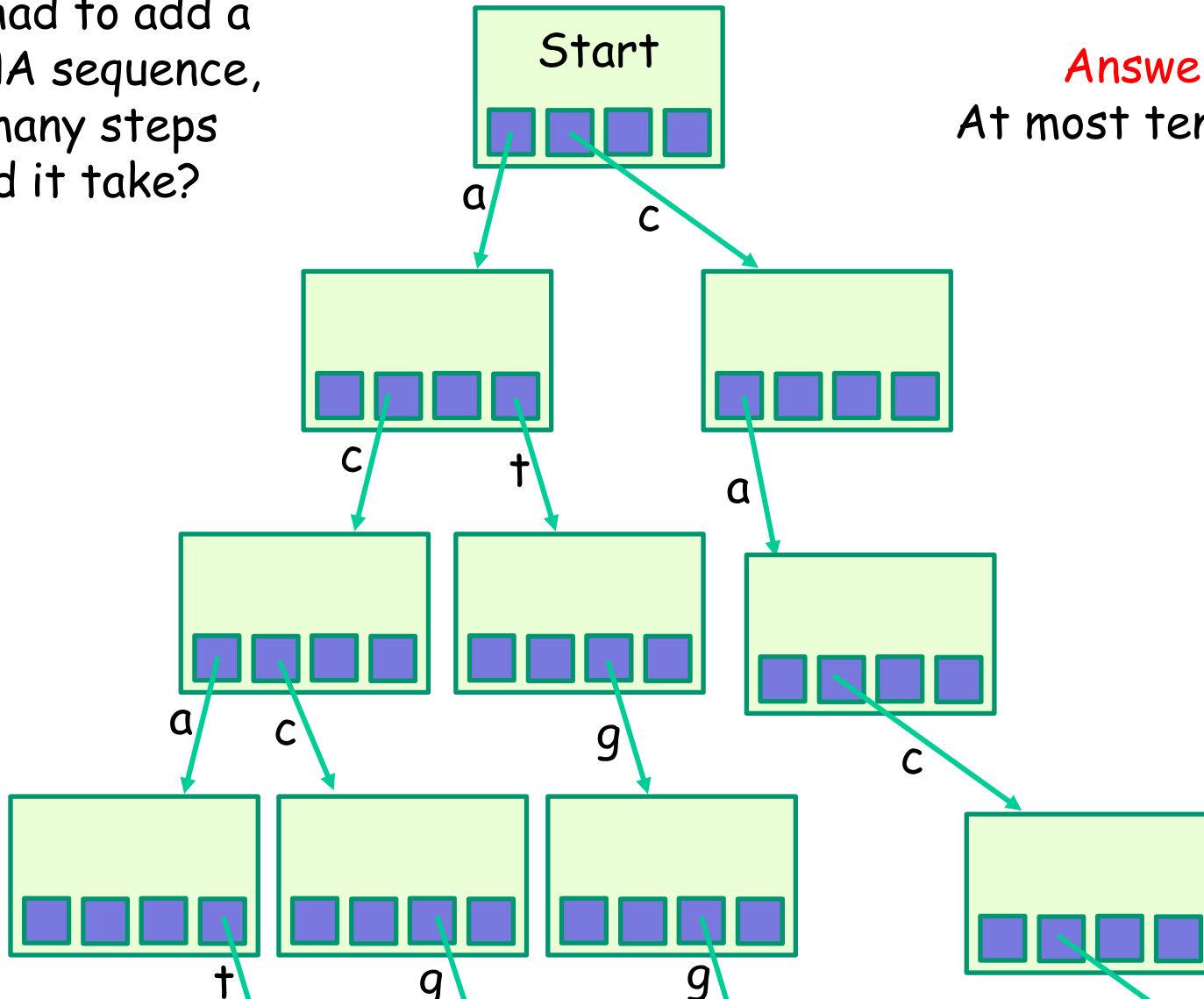
A Trie for DNA

Question:

If we had to add a new DNA sequence, how many steps would it take?

Answer:

At most ten steps!



Sorted Array vs. Trie

Sorted Array	Trie
Limited capacity	Infinitely expandable
Need to move millions of items every time we add an item	Just ten steps to add a new item!
Searchable in dozens of steps	Searchable in ten steps every time
Easy to implement	More complex to implement

Important point: Always choose the simplest data structure possible that meets your project's requirements. It's never a good idea to show off with a more complex data structure if you can use a simpler, easier-to-code, easier-to-understand one!

Data Structures

As we can see, the right data structure can make your algorithms far more efficient!

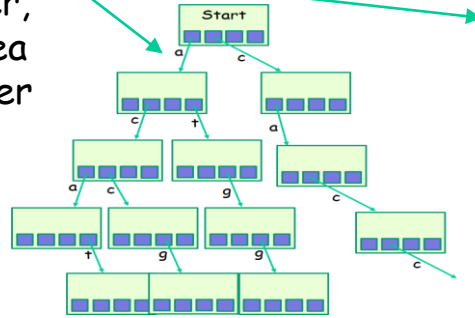
better

So in CS32, we'll also learn all of the most efficient data structures!

Data Structures + Algorithms = Confusion!

As we've seen, your data structures and algorithms can get quite complex.

If you gave your Trie code to another programmer, they would have no idea how to use it! It's super complex!



So it always helps to also **create a few simple functions** that ~~hide the gory details...~~

Such a collection of simple functions is called an "interface."

An **interface** lets any programmer **use your code** without having to dive into your complex data structures or logic.

```

void Trie::ins_helper(string key) {
    // start from the root node
    Trie* curr = root;
    for (int i = 0; i < key.length(); i++) {
        // return if Trie is empty
        if (curr == nullptr) { return false; }
        // if the end of the key is not reached
        if (key[i] != '\0') {
            // if curr->next[key[i]] is nullptr, create a new node
            if (curr->next[key[i]] == nullptr) {
                curr->next[key[i]] = new Trie(key.substr(i));
            }
            // go to the next node
            curr = curr->next[key[i]];
        }
        // if it is the end of the key, mark the node as leaf
        else {
            curr->isLeaf = true;
        }
    }
}

bool Trie::src_primitive(string key) {
    Trie* curr = root;
    for (int i = 0; i < key.length(); i++) {
        // return if Trie is empty
        if (curr == nullptr) { return false; }
        // if the end of the key is not reached
        if (key[i] != '\0') {
            // if curr->next[key[i]] is nullptr, return false
            if (curr->next[key[i]] == nullptr) { return false; }
            // go to the next node
            curr = curr->next[key[i]];
        }
        // if it is the end of the key, check if the node is leaf
        else {
            if (curr->isLeaf) { return true; }
            else { return false; }
        }
    }
}

bool Trie::del_nodes(Trie*& curr, string key) {
    // return if Trie is empty
    if (curr == nullptr) { return false; }
    // if the end of the key is not reached
    if (key.length() > 0) {
        // if curr->next[key[0]] is nullptr, return false
        if (curr->next[key[0]] == nullptr) { return false; }
        // deletion of the node
        deletion(curr->next[key[0]], key.substr(1));
        // curr->isLeaf == false
        curr->isLeaf = false;
    }
    // if (I have children(curr))
    if (IhaveChildren(curr)) {
        // ...
    }
}

```

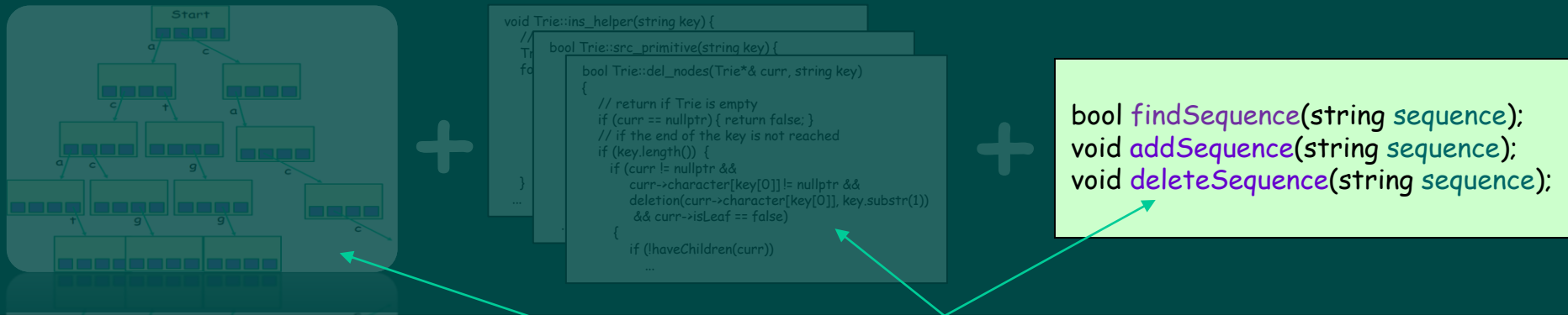
```
bool findSequence(string dna_sequence);  
void addSequence(string dna_sequence);  
void deleteSequence(string dna_sequence);
```

```
int main()
{
    addSequence("tgaccagact");
    if (findSequence("gcgttaacac") == true)
        cout << "This is a known DNA snippet!\n";
}
```

The Abstract Data Type (ADT)

In CS, we call a coordinated group of data structures, algorithms and interface functions an Abstract Data Type.

Abstract Data Type (for DNA searching)



In an ADT, the ~~data structures~~ and ~~algorithms~~ are ~~secret~~.

The ADT provides an **interface** (a simple set of functions) to enable the rest of the program to use it.

Typically, we build programs from a collection of ADTs, each of which solves a different sub-problem.

ADTs in C++

In C++, we use **classes** to define ADTs in our programs!

Each C++ class holds **data structures**, **algorithms** and **interface functions**!

Once we've defined our class, the rest of our program can use it trivially. All our program needs to do is call the functions in our class's **public interface**! The rest of the program can ignore the details of how our class works and just use its features!

```
int main()
{
    DNADatabase d;

    d.addSequence("gagagtcaca");
    d.addSequence("tcaggacata");
    ...

    string dna_seq;

    cout << "Enter a 10-base sequence: ";
    cin >> dna_seq;
    if (d.findSequence(dna_seq) == true)
        cout << "This bacteria is known!";
}
```

```
// A C++ DNA sequencer class...
// (this is really an ADT!)

class DNADatabase
{
    public:
        // our interface functions go here
        void findSequence(...);
        void addSequence(...);

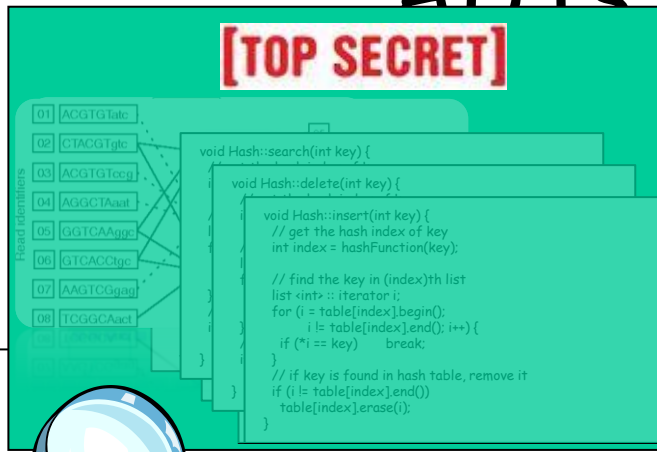
        ...

    private:
        // secret algorithms go here
        ...

        // secret data structures go here
        ...

};
```


ADTs in C++



```
int main()  
{
```

```
    DNADatabase d;
```

```
    d.addSequence("gagagtcaca");
```

```
    d.addSequence("tcaggacata");
```

```
    ...
```

```
    string dna_seq;
```

```
    cout << "Enter a 10-base sequence: ";
```

```
    cin >> dna_seq;
```

```
    if (d.findSequence(dna_seq) == true)
```

```
        cout << "This bacteria is known!";
```

```
}
```

Now what if I wanted to **improve** my class's implementation?

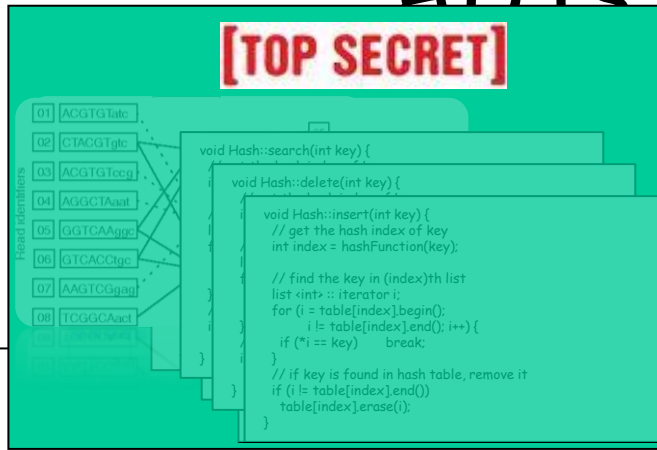
Let's say I made a **radical change** to my **data structures** & **algorithms**...

Would the **user** need to **change** any part of their program?

No! Because these details are hidden from the rest of our code!

The code that uses our class knows nothing about its private data structures and algorithms. All it knows how to use is the public interface.

ADTs in C++



```
int main()  
{
```

```
    DNADatabase d;
```

```
    d.addSequence("gagagtcaca");
```

```
    d.addSequence("tcaggacata");
```

```
    ...
```

```
    string dna_seq;
```

```
    cout << "Enter a 10-base sequence: ";
```

```
    cin >> dna_seq;
```

```
    if (d.findSequence(dna_seq) == true)
```

```
        cout << "This bacteria is known!";
```

```
}
```

This is a **huge benefit** of
Abstract Data Types!

We can **break up our programs** into
small, **self-contained ADTs**...

And **combine these smaller parts**
together to **solve bigger problems**.

What is Object Oriented Programming?



Object Oriented Programming (OOP) is simply a programming model based on the Abstract Data Type (ADT) concept we just learned!

In OOP, programs are constructed from multiple self-contained **classes**.

Each class holds a set of **data structures** and **algorithms** - we then access the class using a **set of interface functions**!

Classes **talk to each other** only by using **public interface functions** - each class knows nothing about how the others work inside.

Intermission Meme

Where Great Ideas Come From



Scientists



Musicians



Chefs



Programmers

@codezinga

C++ Class Review

As we've seen, a "class" is a self-contained problem solver that contains:

- Data structures
- Algorithms
- Interface functions

Since you've probably forgotten everything about classes...

Let's do a quick review of classes by defining our own
Nerd class!



Defining a New Class

```
class Nerd
{
    public:
        Nerd(int stink, int IQ) {
            myStinkiness = stink;
            myIQ = IQ;
        }
        void study(int hours) {
            myStinkiness += 3*hours;
            myIQ *= 1.01;
        }
        int getStinkyLevel() {
            int total_stink = myIQ * 10 +
                myStinkiness;
            return total_stink;
        }
    private:
        int myStinkiness, myIQ;
};
```


First, we write the outer shell of our **class** and give it a **name**.

Then we define our class's **public interface** functions...

Then we define our class's **private** variables and functions...

Our class defines an entirely new **data type**, like **string**, that we can now use in our program.

Alert: **Nerd** is **not a variable**!
It's a new C++ **data type**!



Don't forget the semicolon!

nerd.h

```
class Nerd
{
public:
    Nerd(int stink, int IQ) {
        myStinkiness = stink;
        myIQ = IQ;
    }
    void study(int hours) {
        myStinkiness += 3*hours;
        myIQ *= 1.01;
    }
    ...
private:
    int myStinkiness, myIQ;
};
```

ucla.cpp

```
#include "nerd.h" // #1
int main()
{
    int num_nerds = 1;
    Nerd david(30, 150); // #2
    david.study(10); // #3
}
```

Using a New Class

Once we define a new **class**, like **Nerd**, we can use it to define variables like any traditional data type.

- The Nerd class defines a new data type like int, float, or string
- You typically define each new class in its own **.h file** ("header file") and put the file in the same folder as your **.cpp files**.
- A header file is similar to a .cpp file except you typically only put class declarations and constants in it (you typically put the actual class function {bodies} in your cpp file).
- To use your new class, simply include its **header file** using "quotation marks" (#1)
- You can then define variables with it throughout your program (#2).
- On line #2, **david** is a **Nerd** variable with an initial stinkiness of 30 and an IQ of 150.
- Once you've defined your variable (#3) you can call its member functions, like study.

nerd.h

```
class Nerd
{
public:
    Nerd(int stink, int IQ) {
        myStinkiness = stink;
        myIQ = IQ;
    }
    void study(int hours) {
        myStinkiness += 3*hours;
        myIQ *= 1.01;
    }
    ...
private:
    int myStinkiness, myIQ;
};
```

ucla.cpp

```
#include "nerd.h"
int main()
{
    int num_nerds = 1;
    Nerd david(30, 150); // #1
    david.study(10);
}
```

Using a New Class

Alright, let's see our class in action!

num_nerds

david

```
Nerd(int stink, int IQ) {
    myStinkiness = stink; // #2
    myIQ = IQ;
}
void study(int hours) {
    myStinkiness += 3*hours; // #3
    myIQ *= 1.01;
}
... myStinkiness    myIQ
    -79342          12338
```

When you define your **david** variable, it gets its own copy of all of the **functions** and **member variables** defined in your class! As soon as the david variable is created (#1), C++ calls the **constructor** function inside the variable to initialize its state. Our constructor (#2) and other member functions (#3) have access to **david's** private member variables (like myIQ)! Note: A class's **primitive** member variables (e.g. int's, doubles like myStinkiness) all start out with random values and NOT zero! Your constructor must initialize them.

nerd.h

```
class Nerd
{
public:
    Nerd(int stink, int IQ) {
        myStinkiness = stink;
        myIQ = IQ;
    }
    int getStinkyLevel() {
        int total_stink = myIQ
            * 10 + myStinkiness;
        return total_stink;
    }
private:
    int myStinkiness, myIQ;
};
```

ucla.cpp

```
#include "nerd.h"

int main()
{
    int num_nerds = 1;
    Nerd david(30, 150);

    david.study(10);
}
```

Other Details

You typically only use member variables to store permanent attributes of your class.

- **Stinkiness** and **IQ** are inherent attributes of every **Nerd**, so we make these member variables.
- If you're using a variable for a temporary computation (like `total_stink` to the left), then just use a local variable for that.
- Never use member variables for temporary computation. Only use member variables to store values that you expect your object to retain over time (like the IQ of a nerd, their phone number, or address).

nerd.h

```
class Nerd
{
public:
    Nerd(int stink, int IQ) {
        myStinkiness = stink;
        myIQ = IQ;
    }
    void study(int hours) {
        myStinkiness += 3*hours;
        myIQ *= 1.01;
    }
    ...
private:
    int myStinkiness, myIQ;
};
```

ucla.cpp

```
#include "nerd.h"

int main()
{
    int num_nerds = 1;
    Nerd david(30, 150);

    david.study(10); // #1
}
```

Other Details

- All functions in the **public** section of your class (like the constructor or study()) can be seen/called by all parts of your program.
- All functions and data defined in the private section of your class (like myStinkiness) are hidden from the rest of your program. They may only be used by other functions defined in the Nerd class.
- Notice how the main function calls david.study(10) on line #1. This is legal because the study() method is in Nerd's public section.
- Also notice that your Nerd constructor and study method may access/change the Nerd's private member variables (like myIQ).
- However, other functions outside of your class, like main(), are forbidden from accessing these private member variables. If your main() function were to try to cout << david.myIQ; this would result in an error.
- Hiding the internal implementation details of a class is called "**encapsulation**."
- Encapsulation makes your program simpler, since each class works without knowledge of how the other classes work internally.
- So we can change how one class works internally (in CS lingo, we "refactor" it), and the other classes will continue to work as-is without any changes!