# YEAH A8

The Adventures of Links



## **Assignment overview**

This is the **penultimate** 106B assignment!

This assignment is due Friday 3/12, and the grace period expires the following Sunday.

You're welcome to work in pairs on this assignment.

#### Assignment overview

#### This assignment consists of two parts:

- 1. Labyrinth Using your debugger skills, can you escape a linked list labyrinth, made specifically for you?
- 2. DNA Now that you're comfortable inspecting linked lists, you'll need to implement a series of functions that operate on **nucleotides** represented by linked lists!

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To escape, you'll need to collect three magic items: a wand, a spellbook, and a potion.

This maze exists as a linked structure consisting of these elements:

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struct MazeCell {
    Item whatsHere; // Item present, if any.
    MazeCell* north; // The cell north of us, or nullptr if we can't go north.
    MazeCell* south;
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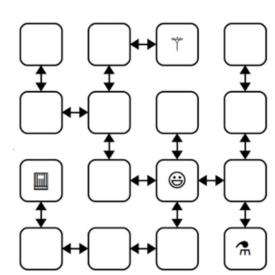
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};

    Remember enums?
    Wish you could forget them?:D

    NOTHING, SPELLBOOK, POTION, WAND
};
```

Here's what an example maze might look like!

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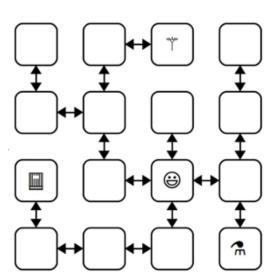
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Notice that not all have 4 valid arrow directions.

Directions that don't lead to other cells are **nullptr**.

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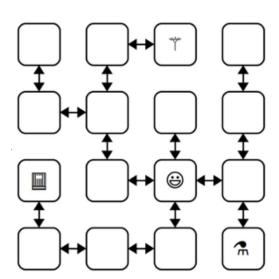
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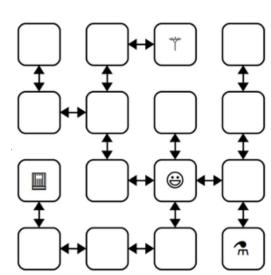
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As you can see, most **MazeCells** are empty, but some have the magical items in them!

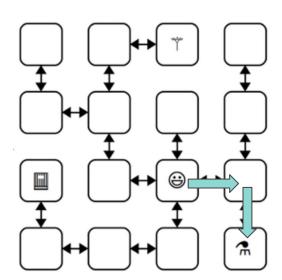
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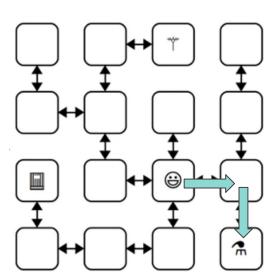
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In this path, we effectively went from cell start to start->south.

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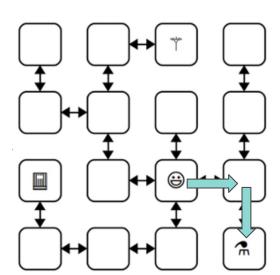


In this path, we effectively went from cell start to start->south.

This path was valid because we didn't go through any "walls" (i.e. cells that aren't connected), and we even found a magic item!

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    MazeCell* south;
    MazeCell* east;
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};
```



In this path, we effectively went from cell **start** to **start->east->south**.

This path was valid because we didn't go through any "walls" (i.e. cells that aren't connected), and we even found a magic item!

You could represent this subpath as the string "ES", where each character represents a direction you went from the starting location.

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A path leads to **freedom** if it: 1- never attempts to move into a direction that's **nullptr** (i.e. never goes through a "wall"), and 2 - picks up all 3 magical items.

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You should expect to iterate through the **entire** string, not stopping early unless you're asked to go through a wall (return false).

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Some notes about the problem:

- You can implement this iteratively or recursively, whichever you choose :)

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- One tricky edge case if you find all three items, **keep looping through the string.** If you happen to **also** move through a wall, you should **return false** anway.

Any Questions?

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Now that you've written a function that can determine whether a path gets you out of a labyrinth, it's your turn to find your way out of a **personalized** one!

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In the file LabyrinthEscape.cpp, you'll need to put your name in the string myName:

```
Labyrinth.cpp

LabyrinthEscape.cpp

SplicingAndDicing.cpp

Other files

MARNING: Once you've set set this constant and started exploring your maze,

* WARNING: Once you've set set this constant and started exploring your maze,

* do NOT edit the value of MyName. Changing MyName will change which maze you

* get back, which might invalidate all your hard work!

11

*/

12

const string MyName = "put your name(s) here!";
```

Once you've put your name into the file, set a breakpoint on the first test in the file (for me it's line 33). Run your code in the debugger, and you'll be placed in a maze personally generated for your name!

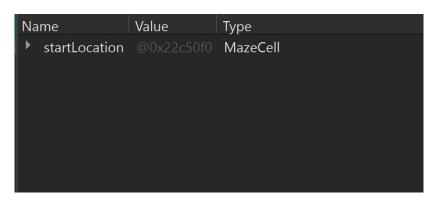
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This is what your debugger pane should look like when you hit the breakpoint:

Na	ıme	Value	Туре
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You've been given a startLocation! Open it up, and it should behave like the MazeCell's that you've seen before.

Here's what opening the startLocation yields me:

Name			Value	Туре
•	sta	artLocation	@0x22c50f0	MazeCell
		east	0x0	MazeCell *
		north	0x0	MazeCell *
		south	0x0	MazeCell *
	•	west	@0x2230850	MazeCell
		whatsHere	Item::NOTHING (0)	ltem

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Notice that there's nothing in the whatsHere field!

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Notice that there's nothing in the whatsHere field!

Also, the only non-wall place we can go is in the west direction! Let's go there:)

Here's what opening the west location yields me:

Name	Value	Туре
▼ startLocation		MazeCell
east	0x0	MazeCell *
north	0x0	MazeCell *
south	0x0	MazeCell *
▼ west	@0x2230850	MazeCell
east		MazeCell
north	0x0	MazeCell *
south		MazeCell
west	0x0	MazeCell *
whatsHere	Item::NOTHING (0)	ltem
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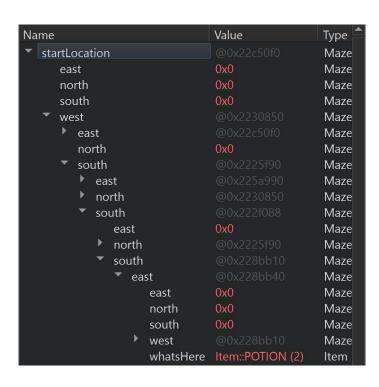
Rats! Nothing here too!

Here's what opening the west location yields me:

Name	Value	Туре
▼ startLocation		MazeCell
east	0x0	MazeCell *
north	0x0	MazeCell *
south	0x0	MazeCell *
▼ west	@0x2230850	MazeCell
east		MazeCell
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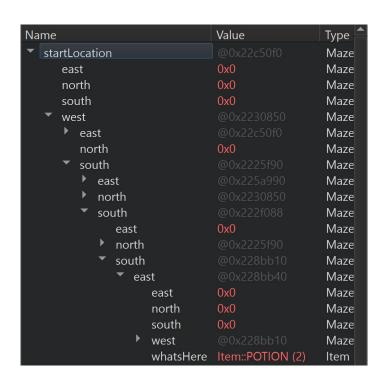
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Look inside the indented (west) struct. Notice that the west->east memory address is identical to the startLocation address. In this problem, all paths are bi-directional, so take a second to register why this must always be true.



Eventually, you'll find an item!

Quiz question: What is the string path starting at **startLocation** that find the POTION?



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Answer: "WSSSE"

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## Part 1.5: Escape from the Labyrinth

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Once you have a good idea of where your items are, write out the string path that hits all 3 of your items (i.e. NNSSEWEWWWESNEENESNWS). Put that path in the variable **ThePathOutOfMyMaze** variable and run the tests -- if the test passes, then you've successfully escaped!

## Part 1.5: Escape from the Labyrinth

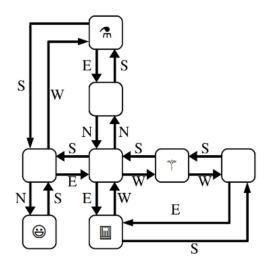
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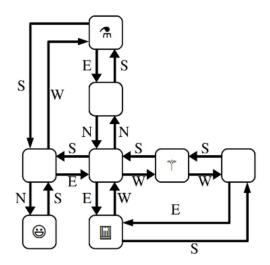
In the last portion of part 1, your job is to escape from a labyrinth like the one from the last part.... but this one's a bit different.



This time, the NSEW pointers may point in any direction! (for example, in one MazeCell, east may actually point south!)

More simply, put, the names of the 4 pointers in the MazeCell have no directional meaning.

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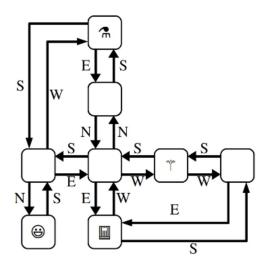


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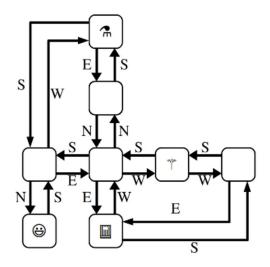
Luckily, you can still find a way out of the maze! For example, the path SW finds one of the magical items!

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When solving this maze, we've guaranteed that the directions wont change on you (i.e. the path SW will always find the top item in this maze), but you're going to have to spend more time mapping this maze out.

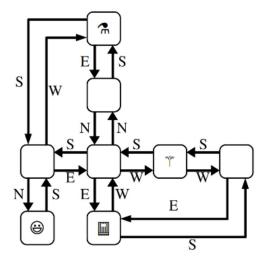
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We've also guaranteed again that all paths are bi-directional, but as you can see, you're going to need to be careful to figure out which path is the way back.

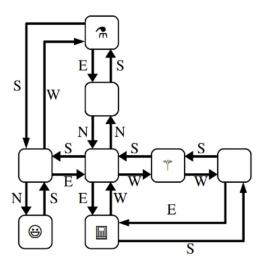
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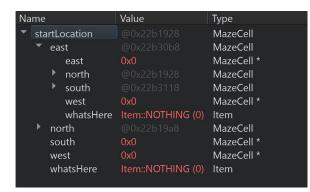
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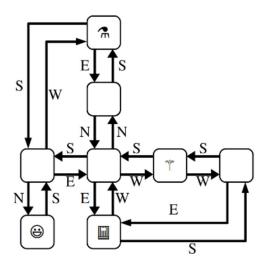


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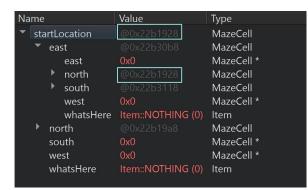


If I move east, which direction moves me back to the start location?

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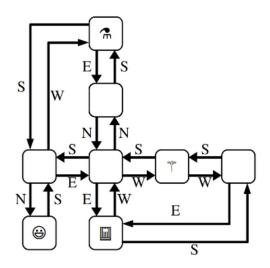


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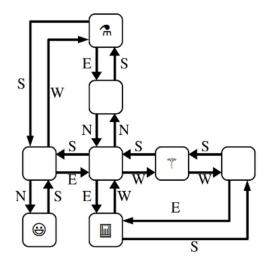
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In this way, you will (slowly and carefully) figure out which links bring you to new maze locations and which ones turn you in loops. Pay attention to the memory addresses at all times because you might end up looping back to a cell that you had visited long ago!

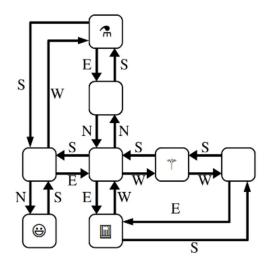
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Once you have the string path to get you out, fill it in at **ThePathOutOfMyTwistyMaze** and you'll be good to go!

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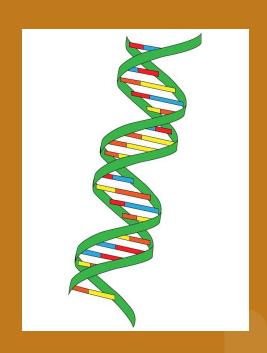


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One last point -- if you re-run your program, the memory addresses will become different, so try and do this in one run!

Part 2: DNA



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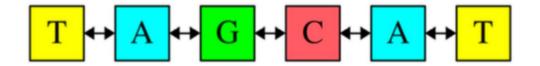
```
struct Nucleotide {
    char value;
    Nucleotide* next;
    Nucleotide* prev;

/* This custom macro assists with memory leak detection. */
    TRACK_ALLOCATIONS_OF(Nucleotide);
};
```

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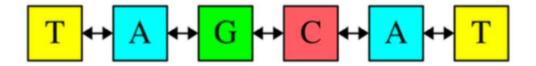
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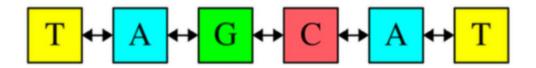


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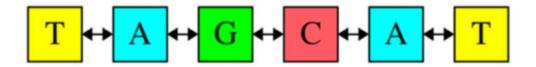
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An example of what a doubly linked list looks like:

Note that the first **T** has a **nullptr** as its **prev** pointer, and the last **T** has a **nullptr** as its **next** pointer.



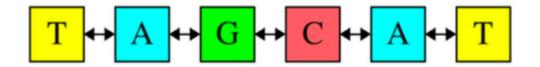
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Function 1: Given a pointer to the beginning of a DNA strand, free all of the associate memory.

void deleteNucleotides(Nucleotide\* dna);



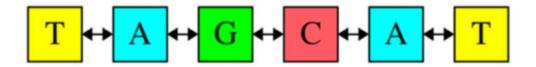
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Function 1: Given a pointer to the beginning of a DNA strand, free all of the associate memory.

Function 2: Given a pointer to the beginning of a DNA strand, return a string representing the content of the strand.

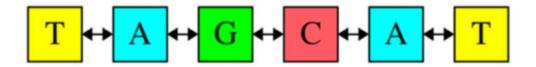
Using this function on the above strand would return **TAGCAT**.

string fromDNA(Nucleotide\* dna);



Function 3: Given a string representing a DNA strand, build up a corresponding doubly linked list and return a pointer to the beginning of the strand.

Nucleotide\* toStrand(const string& dna);



Function 3: Given a string representing a DNA strand, build up a corresponding doubly linked list and return a pointer to the beginning of the strand.

Calling this function give TAGCAT would construct the strand above.

# Nucleotide\* toStrand(const string& dna);

- If you are having trouble with doubly linked list:
  - o End of lecture 21.
  - o Problems 6 to 9 in Section 7.

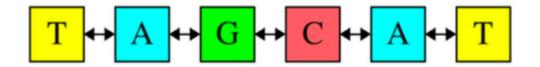
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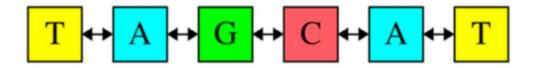
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- Draw pictures! Even Keith does that :)
- Test your code before moving on. These functions are the foundation moving forward.



Function 4: Given a pointer to the beginning of a DNA strand **dna**, look for a target sequence **target**.

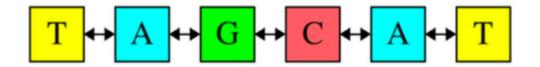
Nucleotide\* findFirst(Nucleotide\* dna, Nucleotide\* target);



Function 4: Given a pointer to the beginning of a DNA strand **dna**, look for a target sequence **target**.

If found, return a pointer to the beginning of that sequence.

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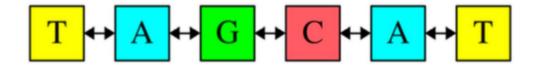


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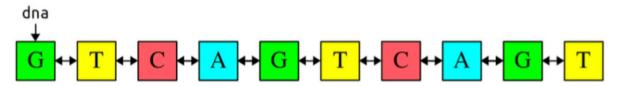
If found, return a pointer to the beginning of that sequence.

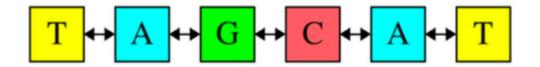
Otherwise, return a **nullptr**.

Nucleotide\* findFirst(Nucleotide\* dna, Nucleotide\* target);

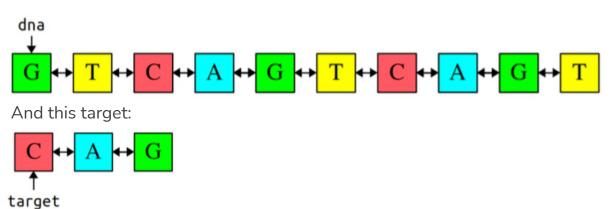


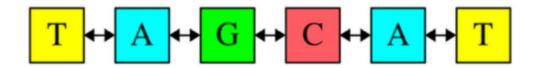
#### Given this dna:



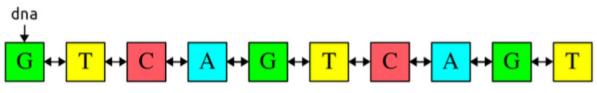


#### Given this dna:

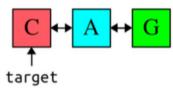




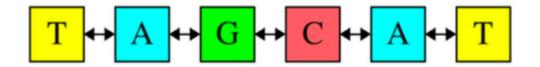
#### Given this dna:



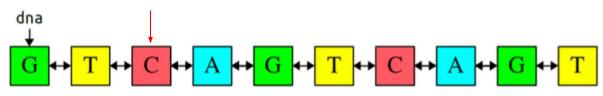
And this target:



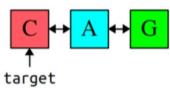
Where should the pointer we are returning be pointing to?



#### Given this dna:



And this target:



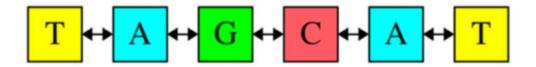
Where should the pointer we are returning be pointing to?

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- If the target sequence is empty, you should return a pointer to the beginning of the DNA strand.

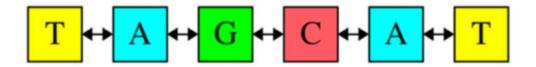
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- If the target sequence is empty, you should return a pointer to the beginning of the DNA strand.
- You may assume that target and dna are pointing to completely different Nucleotide objects
- When in doubt, pictures and well-constructed tests are your best friends!



Function 5: Given a pointer to the beginning of a DNA strand **dna** by reference, look for a target sequence **target**.

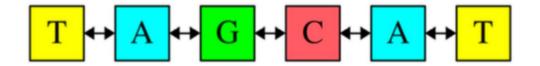
bool spliceFirst(Nucleotide\*& dna, Nucleotide\* target);



Function 5: Given a pointer to the beginning of a DNA strand **dna** by reference, look for a target sequence **target**.

If found, remove the sequence from that strand and return true.

bool spliceFirst(Nucleotide\*& dna, Nucleotide\* target);

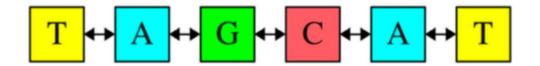


Function 5: Given a pointer to the beginning of a DNA strand **dna** by reference, look for a target sequence **target**.

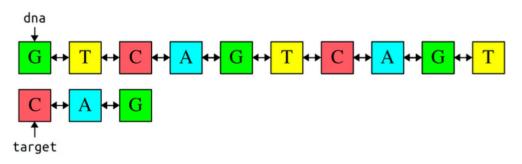
If found, remove the sequence from that strand and return true.

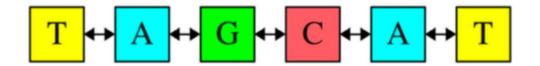
If not found, leave the strand as it is and return false.

bool spliceFirst(Nucleotide\*& dna, Nucleotide\* target);

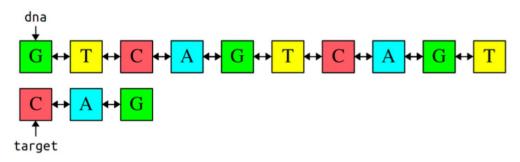


Only the first instance of the sequence should be removed:



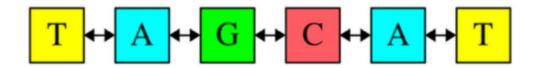


Only the first instance of the sequence should be removed:

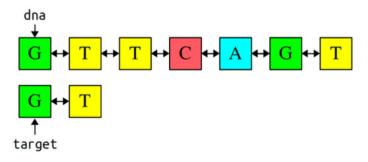


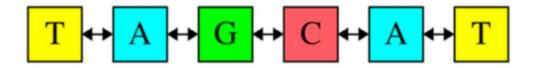
#### Result:

$$\begin{array}{c}
d_{\text{na}} \\
\downarrow \\
G \\
\downarrow \\
T \\
\downarrow \\
T \\
\downarrow \\
T \\
\downarrow \\
C \\
\downarrow \\
A \\
\downarrow \\
G \\
\downarrow \\
T
\end{array}$$

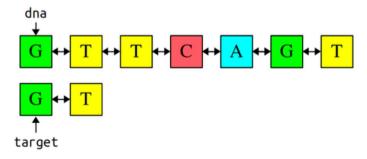


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\text{dna} \\
\downarrow \\
T \\
\downarrow \\
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A \\
\downarrow \\
G \\
\downarrow \\
T
\end{array}$$

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- Free the sub-strand you sliced out.

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- Every note from before holds.