

Tries and Applications

ITP 435 Week 4, Lecture 2





Tries



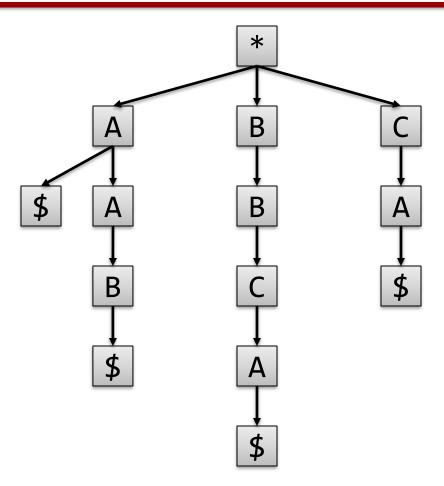
What's a trie?



Usually pronounced "try"

 A tree used (typically) to store strings and allows for efficient matching of patterns and implementing things like autocomplete

 Example on the right (ignore the * and \$ for now)



What to store for each node?



We're going to "rule of zero" this, so no raw pointers!

- Array of unique_ptrs to children number of elements is alphabet size + 1 (we'll talk about why the + 1 later)
- char for the letter stored at the node

How will we declare the trie?



template <size_t AlphabetSize, typename LetterToIdxFunc>
class Trie

- AlphabetSize How many different letters there are in the alphabet the trie needs to support (for space efficiency)
- LetterToldxFunc A function to map a letter to a specific index in the array of children we store at each node

Sample LetterToldxFunc



Suppose we have an alphabet size of 3 and the three letters are A,
 B, and C:

```
struct BasicTextIdx {
    size_t operator()(char c) const {
        switch (c) {
            case 'A': return 0;
            case 'B': return 1;
            case 'C': return 2;
            default: return 0;
        }
    }
}
```

What member data does the trie have?



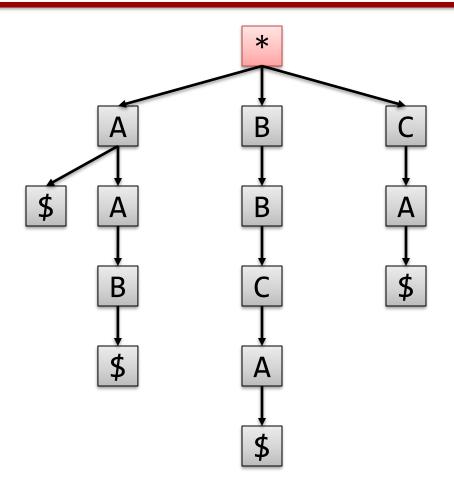
Pointer to root node

Instance of LetterToldxFunc object

What letter do we store at the root node?



- We'll just use *
- (This assumes * doesn't exist in the alphabet)



Handling Common Prefixes

 Suppose we need to store "BANANA" and "BANANAS"

 The proposed trie on the right does not encode enough information

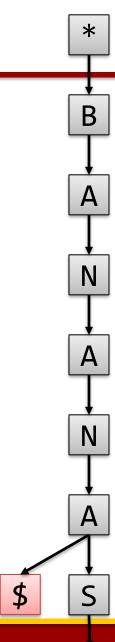




Handling Common Prefixes

 Suppose we need to store "BANANA" and "BANANAS"

- Use a special character \$ to denote the end of a string
- (This assumes \$ is not a valid letter in the alphabet)





Accounting for \$ and the LetterToldxFunc



- Just assume that \$ is always index 0
- Add 1 to LetterToldxFunc result to get the correct final location in the array:

```
struct BasicTextIdx {
    size_t operator()(char c) const {
        switch (c) {
            case 'A': return 0;
            case 'B': return 1;
            case 'C': return 2;
            default: return 0;
        }
    }
}
```

Even though BasicTextIdx says 'A' is at 0, we'll actually store it at 1



Insertion



Insertion



 The general idea is you get a string to insert and when you insert you either use existing nodes when appropriate or create new nodes if they don't already exist

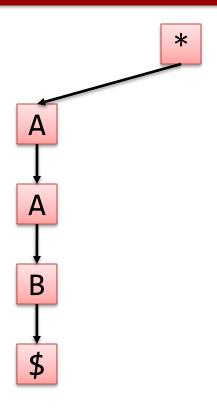
void Insert(std::string_view word)



```
Trie<3, BasicTextIdx> trie;
trie.Insert("AAB");
trie.Insert("A");
trie.Insert("BBCA");
trie.Insert("CA");
```

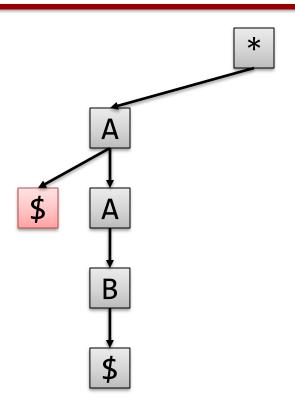


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trie.Insert("AAB");
trie.Insert("A");
trie.Insert("BBCA");
trie.Insert("CA");
```



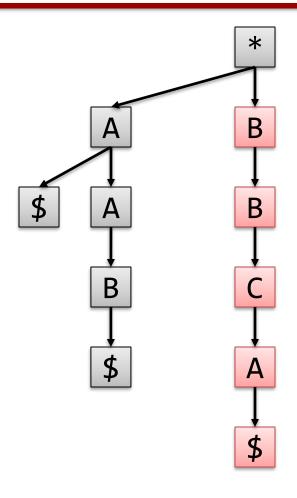


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Trie<3, BasicTextIdx> trie;
trie.Insert("AAB");
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trie.Insert("CA");
```



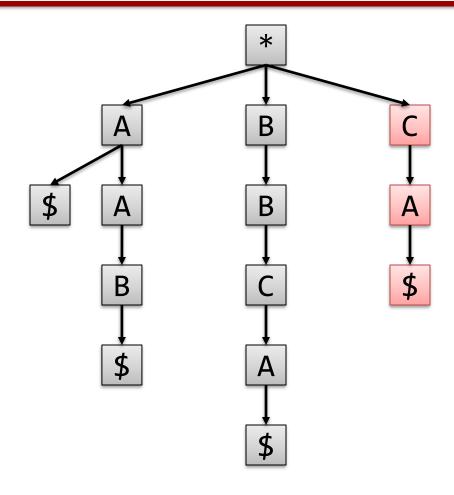


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trie.Insert("AAB");
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trie.Insert("CA");
```



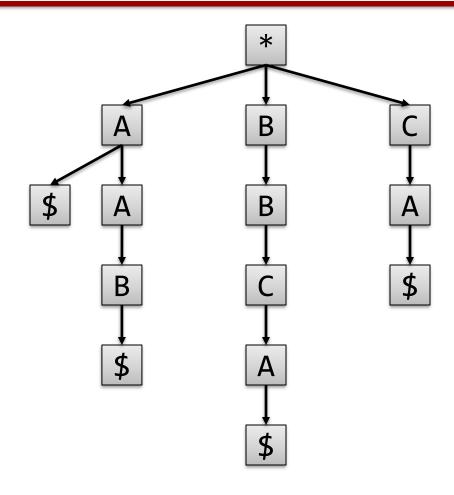


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trie.Insert("AAB");
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```





```
Trie<3, BasicTextIdx> trie;
trie.Insert("AAB");
trie.Insert("A");
trie.Insert("BBCA");
trie.Insert("CA");
```





Breadth-First-Search (BFS)



Breadth-First-Search Traversal



- This is not a particularly useful operation on a trie, but it is helpful for writing test cases to ensure the trie is constructed to spec!
- Will call visitFunc on each node in the tree in a BFS manner, starting at root

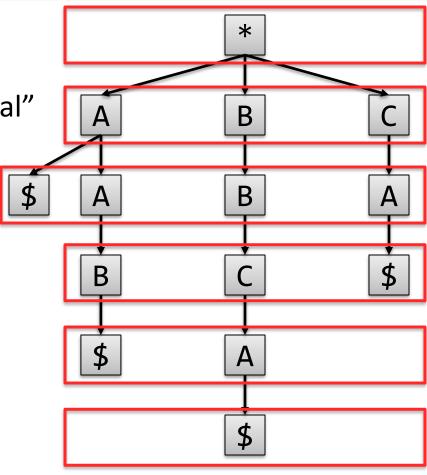
```
void BFS(std::function<void(char)> visitFunc)
```

Breadth-First-Search



 Can just implement using a std::queue

Equivalent to "level-order traversal"



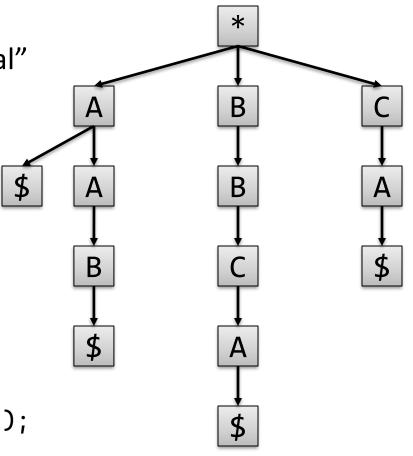
Breadth-First-Search Example



Can just implement with a queue

Equivalent to "level-order traversal"

```
std::string result;
trie.BFS([&result](char c) {
   result += c;
});
REQUIRE(result == "*ABC$ABABC$$A$");
```





FindPrefix



FindPrefix



 Given a string, finds the longest prefix of that string which exists in the trie (or an empty string if none exists)

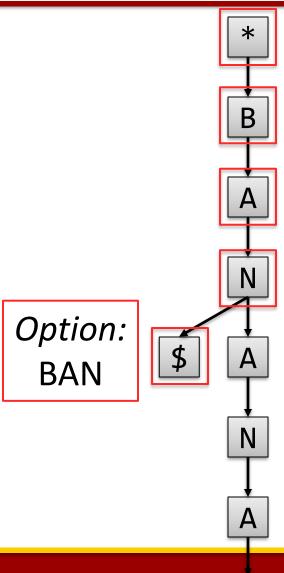
std::string FindPrefix(std::string_view word)

FindPrefix Examples



```
Trie<26, EnglishTextIdx> t;
t.Insert("BAN");
t.Insert("BANANA");

t.FindPrefix("BAN")
??
"BAN"
```



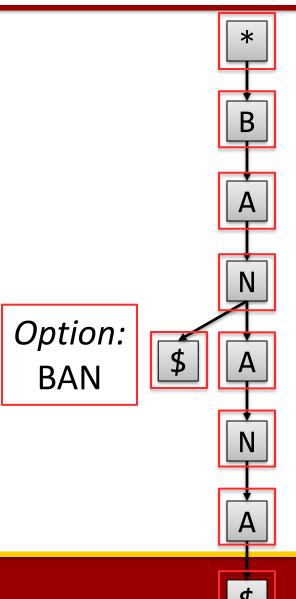
FindPrefix Examples



```
Trie<26, EnglishTextIdx> t;
t.Insert("BAN");
t.Insert("BANANA");

t.FindPrefix("BANANAS")
??
"BANANA"
```

To implement this efficiently, update the best option along the way during the search



*Option:*BANANA

In-class Week 4 Lecture 2



FindPrefix practice





CompleteFromPrefix

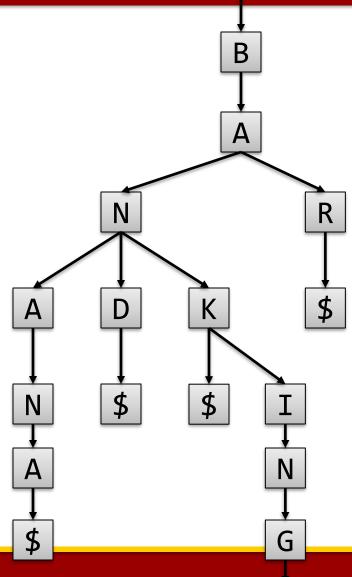


CompleteFromPrefix



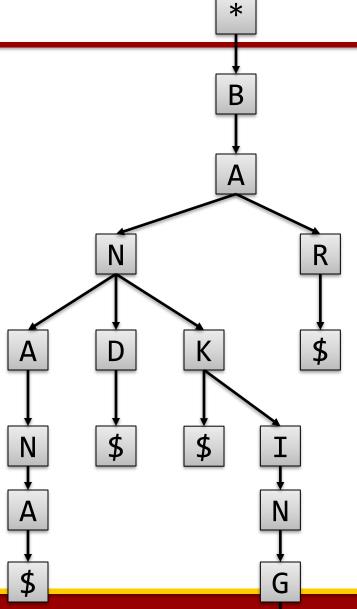
 Given a string, finds the X shortest words in the trie that match the prefix

Given this trie



*

t.CompleteFromPrefix("BAN");



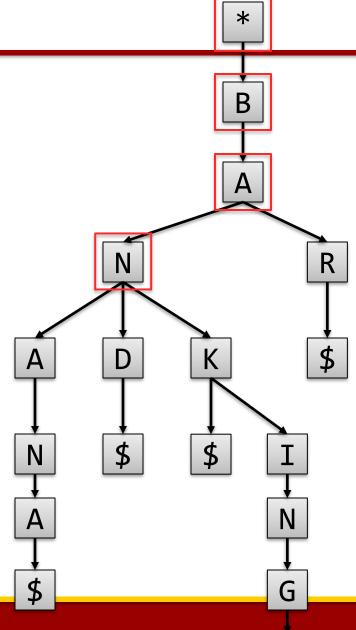




t.CompleteFromPrefix("BAN");

First, match "BAN"

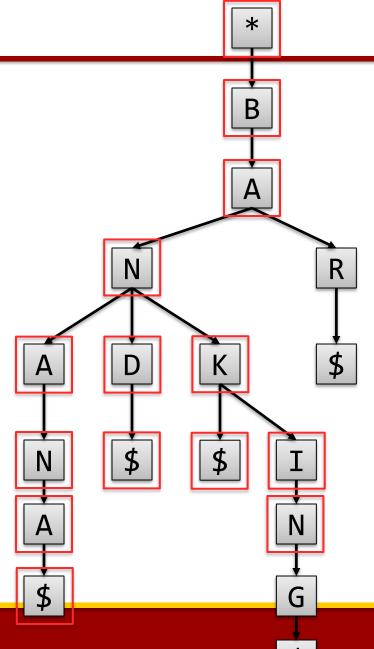
(If there is no match in the trie then there are no results)



t.CompleteFromPrefix("BAN");

Next, do a BFS from that node stopping when you hit count complete words (or stopping if the BFS runs out)

BAND BANK BANANA

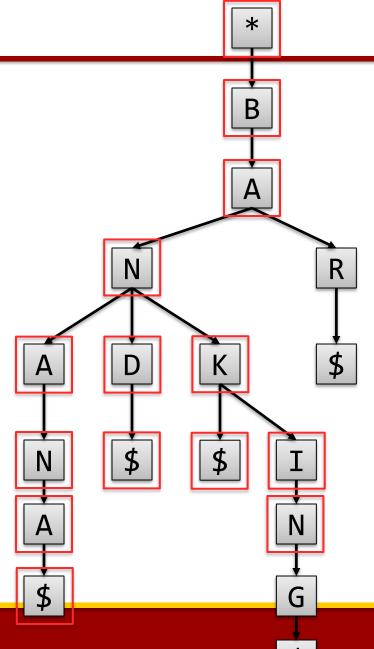


Southern California

How to reconstruct the words from the trie?

Option 1: During the BFS also save the partially-completed word in the queue, so you can just add the additional letter as it continues (fastest)

Option 2: Once you hit a \$, traverse the parents back to root and reverse the word



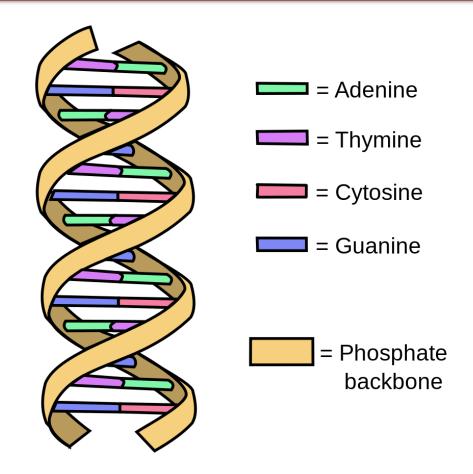


DNA Pattern Matching



DNA





A, T, C, and G are called nucleotides

FASTA File Format



- A simple, text-based file format used to describe (among other things) DNA nucleotide sequences
- First line has a comment/description of the file, and subsequent lines have a sequence of nucloetides
- Example:

>gi|319999821:c124527448-124526573 Pan troglodytes isolate Yerkes chimp ATGATACCCATCCAACTCACTGTCTTCTTCATGATCATCTATGTGCTTGAGTCCTTGACAATTATTGTGCAG AGCAGCCTAATTGTTGCAGTGCTGGGCAGAGAATGGCTGCAAGTCAGAAGGCTGATGCCTGTGGACATGATT CTCATCAGCCTGGGCATCTCTCTCTCTCTCTACAGTGGGCATCAATGCTGAACAATTTTTTGCTCCTATTTT AATTTGAATTATGTACTTTGCAACTTAACAATCACCTGGGAATTTTTTTAATATCCTTACATTCT

DNA Pattern Matching



- Given a long DNA sequence, find the largest prefix in the trie that matches
- Start at letter 0 of the sequence, FindPrefix, then go on to letter 1, FindPrefix, etc
- Will use to find some transcription factor proteins



Autocomplete



Autocomplete



 Essentially just have a large dictionary you load into the trie, and given the prefix use CompleteFromPrefix



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