Project title: Suffix Tree provided in Sample project 2 (Bioinformatics)

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Accomplished:

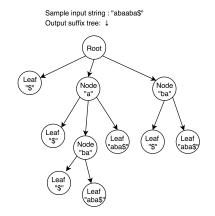
- Everything proposed in the project proposal except MUMs
- Two approaches to build a suffix trie
 - Create suffix trie directly using tuple and map
 - Generate a new suffix trie by inserting a new suffix string into the old one (start with Root [])
- Convert suffix trie into suffix tree
- Indexed suffix tree
- Find whether a given string s0 is a substring of s1

Not completed: MUMs

Code accounting: 229(SfxTree.hs) + 37(Main.hs) = 266

Sample inputs and outputs:

Run main and main2 functions to check the sample outputs:



Also, we can type "build <input string>" to check the result:

```
[*Main> build "gacttg"
-----Testing-----Testing-----
Input String: gacttg
Parse: gacttg$
Index: [6,5,4,3,2,1,0]
```

In addition, "sfxtrie1 \$ parse < input string>" or "sfxtrie2 \$ parse2 < input string>" can generate a suffix trie of the input string; "sfxtree1 < input string>" or "sfxtree2 < input string>" can generate a suffix tree of the input string; "indexTree < input string>" can generate a indexed suffix tree of the input string; "insertTrie < new suffix> < old suffixTrie>" can generate a new suffix trie after suffix insertion; "trieToTree < suffixTrie> < suffixTree>" can convert a suffix trie into a suffix tree; "isSubstr < string A> < string B>" can find whether string A is a substring of string B.

Instructions for running the code:

Only Need: **ghc**, emacs (ghc is enough actually since we can run the code via terminal or cmd) Details: just go to the directory of the source fils and type following commands:

```
[jk:production dihuang$ ghci
GHCi, version 8.2.2: http://www.haskell.org/ghc/ :? for help
[Prelude> :1 Main.hs
[1 of 2] Compiling SfxTree (SfxTree.hs, interpreted)
[2 of 2] Compiling Main (Main.hs, interpreted)
Ok, two modules loaded.
*Main>
```