

## CIS241 Lab 3: play with pointers

### String Processing

A DNA can be represented as an array of N characters, where N is very large (on the order of tens of thousands). A problem in genetic research is to discover whether any pattern of length LEN is ever repeated in the array, where LEN is a fixed and small constant, say 4.

Use the command below to download the source code **dna.c** from the instructor's web page to your current directory.

wget <http://cis.gvsu.edu/~wangx/teaching/cis241/dna.c>

Program **dna.c** is intended to find if a substring of length 4 in the given string is repeated. For the string that is stored in array **line**, this program would show that substring "book" is repeated, where its first occurrence is pointed to by pointer p1 and its second occurrence by p2. Note that the out loop in this program allows p1 to point to a substring in the given string and the inner loop allows p2 to point to a substring after p1. As such, these two nested while-loops make it possible to compare every possible pair of substrings in a given string. If the substring of LEN characters that p1 points to is identical to the one that p2 points to, the results are displayed on the screen and the program terminates. If no matches are located, the program display such a message and terminates.

In this lab, you will write C code to replace each line of pseudo-code in this program. Note that the given pseudo-code is intended to show the main logic of this program. You may add additional statements to address what you find necessary.

Also take the following steps to familiarize yourself with the GNU debugger **gdb**, which would be helpful when you test and debug your program.

- Type **gcc -g dna.c -o dna** to prepare an executable for **gdb**
- Type **gdb dna** to run **gdb** on the executable
- Use **list** to show the first ten lines of code in **dna**
- Use **list** again to show the next ten lines of code in **dna**
- Use **break lineNo** to set one or more breakpoints; for example, one at the first line of the out loop and one at the first line of the inner loop
- Use **run** to run **dna** until the first breakpoint is reached
- Use **print variable** to check the current value of variable
- Use **step** or **step 2** to run one or two lines of code from where this program stopped
- Use **cont** to make this program continue from where it is to the next breakpoint or the end of this program if no more breakpoints are on the way
- Use **quit** to leave **gdb** when the program terminates

At the **gdb** prompt, you may use **help command** to find a description about the named command. Refer the PDF document "Unix Programming Tools" (Section 3) that is available from the following link:

<http://cslibrary.stanford.edu/107/UnixProgrammingTools.pdf>

When your program runs correctly, change the value of LEN from 4 to 5 and then run your program again. It is a case in which no match can be found. Show your program to the instructor after you have done all the above successfully.