Programming Assignment #1

Sorting (on-line submission due: 13:00, 24 March 2024)

TA in charge of PA#1 is Yi-Chen Lin <u>r12943096@ntu.edu.tw</u>

Submission & Online Resources: on NTU COOL Introduction:

In this PA, you are required to implement various sorters that we learnt in class. You can download the *PA1.tar* file from NTU COOL website. Uncompress it using Linux command,

You can see the following directories after uncompressing it.

Name	Description			
bin/	Directory of binary file			
doc/	Directory of document			
inputs/	Directory of unsorted data			
lib/	Directory of library source code			
outputs/	Directory of sorted data			
src/	Directory of source code			
utility/	Directory of checker			

Input/output Files:

In the input file (*.in), the first two lines starting with '#' are just comments. Except comments, each line contains two numbers: index followed by the unsorted number. The range of unsorted number is between 0 and 1,000,000. Two numbers are separated by a space. For example, the file 5.case1.in contains five numbers

```
# 5 data points
# index number
0 16
1 13
2 0
3 6
4 7
```

The output file(*.out) is actually the same as the input file except that the numbers are sorted in *increasing* order. For example, 5.case1.out is like:

```
# 5 data points
# index number
0 0
1 6
2 7
3 13
```

PLOT:

You can visualize your unsorted/sorted numbers by using the gnuplot tool by the command gnuplot. After that, please key in the following

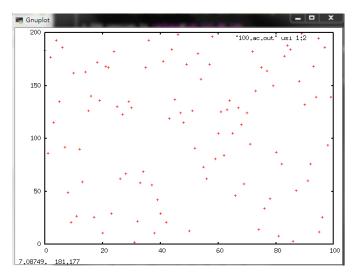
```
set xrange [0:5]
set yrange [0:20]
plot "5.case1.in" usi 1:2
plot "5.case1.out" usi 1:2

# if you want to save to png files
set terminal png
set output "5.case1.out.png"
replot
```

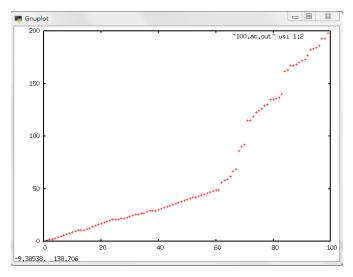
You need to allow X-window display to see the window if you are login remotely. For more gnuplot information, see

http://people.duke.edu/~hpgavin/gnuplot.html

There are two example "before" and "after" sort pictures with 100 numbers benchmark. Before sort:



After sort:



Command line parameters:

In the command line, you are required to follow this format

```
NTU_sort -[IS|MS|QS|HS] <input_file_name> <output_file_name>
```

where IS represents insertion sort, MS is merge sort, QS is quick sort and HS is heap sort. The square bracket with vertical bar '[IS|MS|QS|HS]' means that only one of the four versions is chosen.

The angle bracket <input_file_name> should be replaced by the name of the input file,
*.[case1|case2|case3].in, where case1 represents test case in random order,
case2 is test case in increasing order, and case3 is test case in reverse order. For the best
case, all the numbers are sorted in increasing order. For the worst case, all numbers are
sorted in descending order. For the average case, numbers are in random order.

The output file names are *. [case1|case2|case3].out. Please note that you do NOT need to add '[|]' or '<>' in your command line. For example, the following command sorts 1000.case1.in to 1000.case1.out using insertion sort.

```
./bin/NTU sort -IS inputs/1000.casel.in outputs/1000.casel.out
```

Source code files:

Please notice that all of the source code files have been already finished except sort_tool.cpp. You only need to complete the different sorting functions of class SortTool in sort_tool.cpp. You can still modify other source code files if you think it is necessary. The following will simply introduce the source code files.

main.cpp: main program for PA1

```
// *************************
         // File
[main.cpp]
         //
             Author
                            [Yu-Hao Ho]
                            [The main program of 2019 fall Algorithm PA1]
[2020/9/15 Mu-Ting Wu]
         //
              Synopsis
              Modify
         #include <cstring>
         #include <iostream>
         #include <fstream>
#include "../lib/tm_usage.h"
         #include "sort_tool.h"
         using namespace std;
         void help_message() {
   cout << "usage: NTU_sort -[IS|MS|QS|HS] <input_file> <output_file>" << endl;
   cout << "options:" << endl;
   cout << " IS - Insersion Sort" << endl;</pre>
                             MS - Merge Sort" << endl;
QS - Quick Sort" << endl;
               cout << "
              cout << "
              cout << "
                             HS - Heap Sort" << endl;
         int main(int argc, char* argv[])
               if(argc != 4) {
                   help_message();
                   return 0;
               CommonNs::TmUsage tmusg;
               CommonNs::TmStat stat;
               //////// read the input file /////////
               char buffer[200];
               fstream fin(argv[2]);
               fstream fout;
               fout.open(argv[3],ios::out);
               fin.getline(buffer,200);
fin.getline(buffer,200);
               int junk.num:
               vector<int> data;
               while (fin >> junk >> num)
                    data.push_back(num); // data[0] will be the first data.
                                                  // data[1] will be the second data and so on.
               //////// the sorting part ///////////
               tmusg.periodStart();
SortTool NTUSortTool;
               if(!strcmp(argv[1],"-QS")) {
                    NTUSortTool.QuickSort(data);
               else if(!strcmp(argv[1],"-IS")) {
                    NTUSortTool.InsertionSort(data);
               else if(!strcmp(argv[1],"-MS")) {
                    NTUSortTool.MergeSort(data);
               else if(!strcmp(argv[1],"-HS")) {
                    NTUSortTool.HeapSort(data);
               else {
                    help_message();
                    return 0;
               tmusg.getPeriodUsage(stat);
               cout <<"The total CPU time: " << (stat.uTime + stat.sTime) / 1000.0 << "ms" << endl;
               cout <<"memory: " << stat.vmPeak << "KB" << endl; // print peak memory
               ///////// write the output file ////////
fout << "# " << data.size() << " data points" <<endl;
fout << "# index number" << endl;
               for (int i = 0; i < data.size(); i++)
                    fout << i << " " <<data[i] << endl;
               fin.close();
               fout.close();
               return 0;
```

- Line 36-46: parse unsorted data from input file and push them into the vector.
- Line 52-67: call different function depending on given command.

Line 74-77: write the sorted data file.

sort tool.h: the header file for the SortTool Class

```
// *************************
2. 3. 4. 5. 6. 7. 8. 9. 10. 11. 12. 13. 14. 15. 16. 17. 18. 19. 20. 21. 22. 23. 24. 25. 26. 27. 28. 29. 30.
           File
        //
                        [sort_tool.h]
           Author
                         [Yu-Hao Ho]
                        [The header file for the SortTool Class]
       //
           Synopsis
                         [2020/9/15 Mu-Ting Wu]
           Modify
        #ifndef_SORT_TOOL_H
        #define SORT TOOL H
        #include<vector>
        using namespace std;
        class SortTool {
             public:
                                 SortTool(); // constructor
                  void
                                InsertionSort(vector<int>&); // sort data using insertion sort
                                MergeSort(vector<int>&); // sort data using merge sort
                  void
                  void
                                QuickSort(vector<int>&); // sort data using quick sort
                  void
                                HeapSort(vector<int>&); // sort data using heap sort
             private:
                  void
                                QuickSortSubVector(vector<int>&, int, int); // quick sort subvector
                  int
                                Partition(vector<int>&, int, int); // partition the subvector
                                MergeSortSubVector(vector<int>&, int, int); // merge sort subvector
                  void
                                Merge(vector<int>&, int, int, int, int); // merge two sorted subvectors
                  void
                  void
                                MaxHeapify(vector<int>&, int); // make tree with given root be a max-heap
                                                                          //if both right and left sub-tree are max-heap
                  void
                                BuildMaxHeap(vector<int>&); // make data become a max-heap
                                heapSize; // heap size used in heap sort
                  int
31.
        };
32.
        #endif
```

sort tool.h

- Line 17-20: sort function which will be called in *main.cpp*.
- Line 22: This function will be used in quick sort. It will sort sub vector with given lower and upper bound. This function should be implemented to partition the sub vector and recursively call itself.
- Line 23: This function will be used in quick sort and should be implemented to partition the sub vector.
- Line 24: This function will be used in merge sort. It will sort sub vector with given lower and upper bound. This function should be implemented to call itself for splitting and merging the sub vector.
- Line 25: This function will be used in merge sort and should be implemented to merge two sorted sub vectors.
- Line 26: This function will be used in heap sort and should be implemented to make the tree with given root be a max-heap if both of its right subtree and left subtree are max-heap.
- Line 28: This function will be used in heap sort and should be implemented to make input data be a max-heap.
- sort_tool.cpp: the implementation of the SortTool Class

```
// File
[sort\_tool.cpp]
        //
           Author
                        [Yu-Hao Ho]
        //
                        [The implementation of the SortTool Class]
           Synopsis
                        [2020/9/15 Mu-Ting Wu]
           Modify
        #include "sort_tool.h"
        #include<iostream>
        // Constructor
        SortTool::SortTool() {}
        // Insertsion sort method
        void SortTool::InsertionSort(vector<int>& data) {
            // Function : Insertion sort
            // TODO : Please complete insertion sort code here
        // Quick sort method
        void SortTool::QuickSort(vector<int>& data){
             QuickSortSubVector(data, 0, data.size() - 1);
        // Sort subvector (Quick sort)
        void SortTool::QuickSortSubVector(vector<int>& data, int low, int high) {
             // Function : Quick sort subvector
            // TODO : Please complete QuickSortSubVector code here
            // Hint : recursively call itself
                       Partition function is needed
        int SortTool::Partition(vector<int>& data, int low, int high) {
            // Function : Partition the vector
            // TODO : Please complete the function
            // Hint : Textbook page 171
        // Merge sort method
        void SortTool::MergeSort(vector<int>& data){
             MergeSortSubVector(data, 0, data.size() - 1);
        // Sort subvector (Merge sort)
        void SortTool::MergeSortSubVector(vector<int>& data, int low, int high) {
            // Function : Merge sort subvector
             // TODO : Please complete MergeSortSubVector code here
            // Hint : recursively call itself
                       Merge function is needed
        // Merge
        void SortTool::Merge(vector<int>& data, int low, int middle1, int middle2, int high) {
            // Function : Merge two sorted subvector
             // TODO : Please complete the function
        // Heap sort method
        void SortTool::HeapSort(vector<int>& data) {
            // Build Max-Heap
             BuildMaxHeap(data);
            // 1. Swap data[0] which is max value and data[i] so that the max value will be in correct location
             // 2. Do max-heapify for data[0]
             for (int i = data.size() - 1; i >= 1; i--) {
                 swap(data[0], data[i]);\\
                 heapSize--
                 MaxHeapify(data,0);
        }
        void SortTool::MaxHeapify(vector<int>& data, int root) {
            // Function : Make tree with given root be a max-heap if both right and left sub-tree are max-heap
            // TODO : Please complete max-heapify code here
        //Build max heap
        void SortTool::BuildMaxHeap(vector<int>& data) {
             heapSize = data.size(); // initialize heap size
             // Function : Make input data become a max-heap
            // TODO : Please complete BuildMaxHeap code here
```

- Line 15-18: please complete the function of insertion sort here.
- Line 21-23: the function of quick sort will call function of Sorting sub-vector and give initial lower/upper bound.
- Line 25-30: please complete the function of sorting sub-vector using quick sort algorithm here.
- Line 32-36: please complete the function of partition here.
- Line 39-41: the function of merge sort will call function of Sorting sub-vector and give initial lower/upper bound.
- Line 44-49: please complete the function of sorting sub-vector using merge sort algorithm here.
- Line 52-55: please complete the function of merging two sorted sub-vector here.
- Line 58-68: the function of heap sort will build max-heap first. And then, exchange data iteratively.
- Line 71-74: please complete the function of max-heapify which makes the tree with given root be a max-heap if its right and left sub-tree are both max-heap.
- Line 77-81: please complete the function of building max-heap with given input data.

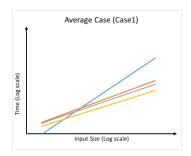
Requirements:

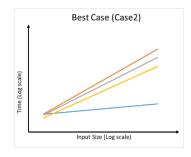
- 1. Please check the source code files under the src directory. You may need to complete the functions of class SortTool in *sort_tool.cpp*. You can also modify *main.cpp* and *sort_tool.h* if you think it is necessary.
- 2. Your source code must be written in C or C^{++} . The code must be executable on EDA union lab machines.
- 3. In your report, compare the running time of four versions of different input sizes. Please fill in the following table. Please use –O2 optimization and turn off all debugging message. You should specify where you run the data, on EDA union lab machines or your local terminal.

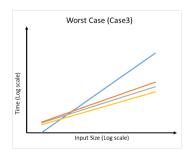
Input size	IS		MS		QS		HS	
	CPU time	Memory						
	(s)	(KB)	(s)	(KB)	(s)	(KB)	(s)	(KB)
4000.case2								
4000.case3								
4000.case1								
16000.case2								
16000.case3								
16000.case1								
32000.case2								
32000.case3								
32000.case1								
1000000.case2								
1000000.case3								
1000000.case1								

4. In your report, plot the trendline of four sorting algorithms to show the growth of run time as a function of input size, and try to analyze the slopes of the curves as well as their relation (as the following example, where each curve represents an algorithm.) Please note that you should transfer the run time and input size to log scale first, then draw the figures. If your trendline is different from following example, you should give reasonable explanation. For example: You may find out Mergesort and Quicksort have same time tendency as insertion sort in worst case, why? How to solve this?

Your figures should be clear and easy to distinguish the curves.







You can skip the test case if the run time is more than 10 minutes.

5. Notice: You are not allowed to include the header <algorithm> or <queue> in STL!

Compile

We expect your code can compile and run in this way.

Type the following commands under <student id> pal directory,

```
make
cd bin
./NTU_sort -[IS|MS|QS|HS] <input_file_name> <output_file_name>
```

We provide the sample makefile, please modify into yours if needed.

Control the stack size

To prevent stack overflow cause by the recursion function calls, please set the stack size to 256MB using the following Linux comment:

ulimit -s 262144

```
# CC and CFLAGS are varilables
1. 2. 3. 4. 5. 6. 7. 8. 9. 10. 11. 12. 13. 14. 15. 16. 17. 18. 19. 22. 22. 22. 22. 22. 23. 33. 34. 35. 36. 37. 38. 34. 41. 42.
       CFLAGS = -c
       AR = ar
       ARFLAGS = rcv
       # -c option ask g++ to compile the source files, but do not link.
       # -g option is for debugging version
       # -O2 option is for optimized version
       DBGFLAGS = -g - D_DEBUG_ON_
       OPTFLAGS = -O2
       # make all
       all: bin/NTU sort
             @echo -n ""
       # optimized version
       bin/NTU_sort: sort_tool_opt.o main_opt.o lib
            $(CC) $(OPTFLAGS) sort_tool_opt.o main_opt.o -ltm_usage -Llib -o bin/NTU_sort
       main opt.o: src/main.cpp lib/tm usage.h
            $(CC) $(CFLAGS) $< -Ilib -o $@
       sort_tool_opt.o: src/sort_tool.cpp src/sort_tool.h
            $(CC) $(CFLAGS) $(OPTFLAGS) $< -0 $@
       # DEBUG Version
       dbg: bin/NTU_sort_dbg
             @echo -n ""
       bin/NTU sort dbg: sort tool dbg.o main dbg.o lib
            $(CC) $(DBGFLAGS) sort_tool_dbg.o main_dbg.o -ltm_usage -Llib -o bin/NTU_sort_dbg
       main_dbg.o: src/main.cpp lib/tm_usage.h
            $(CC) $(CFLAGS) $< -Ilib -o $@
       sort_tool_dbg.o: src/sort_tool.cpp src/sort_tool.h
            $(CC) $(CFLAGS) $(DBGFLAGS) $< -0 $@
       lib: lib/libtm_usage.a
       lib/libtm usage.a: tm usage.o
            $(AR) $(ARFLAGS) $@ $<
       tm_usage.o: lib/tm_usage.cpp lib/tm_usage.h
            $(CC) $(CFLAGS) $<
       # clean all the .o and executable files
            rm -rf *.o lib/*.a bin/*
```

makefile

- Line 38-39: compile the object file tm usage.o from tm usage.cpp and tm usage.h
- Line 36-37: archive *tm_usage.o* into a static library file *libtm_usage.a*. Please note that library must start with *lib* and ends with *.a*.
- Line 37: this small library has only one objet file. In a big library, more than one objective files can be archived into a single *lib*.a* file like this

```
ar rcv libx.a file1.o [file2.o ...]
```

- Lines 12-21: When we type 'make' without any option the makefile will do the first command (line.12 in this sample). Thus, we can compile the optimization version when we type 'make'. This version invokes options '-O2' for speed improvement. Also 'DEBUG ON' is not defined to disable the printing of arrays in *sort tool.cpp*.
- Lines 23-32: Compile the debug version when we type 'make dbg'. This version invokes options '-g' (for DDD debugger) and also '-D_DEBUG_ON_' to enable the printing of arrays in *sort tool.cpp*.
- Lines 13,25: @echo -n "" will print out the message in "". In this sample we print

nothing.

```
Notice: $< represent the first dependency.
```

\$@ represent the target itself.

```
Example: a.o : b.cpp b.h  \$ (CC) \$ (CFLAGS) \$ (DBGFLAGS) \$ < -o \$ \emptyset   \$ <= b.cpp \$ (a) = a.o
```

You can find some useful information here.

Makefile Tutorial By Example

Validation:

You can verify your answer very easily by comparing your output with case2 which is the sorted input. Or you can see the gnuplot and see if there is any dot that is not sorted in order.

Also, you can use our result checker which is under utility directory to check whether your result is correct or not. To use this checker, simply type

```
./PA1_Result_Checker <input_file> <your output file>
```

Please notice that it will not check whether the format of result file is correct or not. You have to check the format by yourself if you modify the part of writing output file in *main.cpp*.

Submission:

You need to create a directory named **<student_id>_pa1**/ (e.g. b09901000_pa1/) (**student id should start with a lowercase letter**) which must contain the following materials:

- A directory named src/ contains your source codes: only *.h, *.hpp, *.c,
 *.cpp are allowed in src/, and no directories are allowed in src/;
- 2. A directory named bin/ containing your executable binary named NTU sort;
- 3. A directory named **doc**/ containing your report;
- 4. A makefile named **makefile** that produces an executable binary from your source codes by simply typing "make": the binary should be generated under the directory <student_id>_pa1/bin/;
- 5. A text readme file named **README** describing how to compile and run your program;
- 6. A report named **report.pdf** on the data structures used in your program and your findings in this programming assignment.

We will use our own test cases, so do NOT include the input files.

In summary, you should at least have the following items in your *.tgz file.

```
src/<all your source code>
lib/<library file>
bin/NTU_sort
doc/report.pdf
makefile
README
```

The submission filename should be compressed in a single file <student_id>_pa1.tgz. (e.g. b09901000_pa1.tgz). You can use the following command to compress a whole directory:

```
tar -zcvf <filename>.tgz <dir>
```

For example, go to the same level as PA1 directory, and type

```
tar -zcvf b09901000 pa1.tgz b09901000 pa1/
```

Please submit a single *.tgz file to NTU COOL system before 3/26(Sun.) 13:00.

You are required to run the checksubmitPA1 script to check if your .tgz submission file is correct. Suppose you are in the same level as PA1 directory

```
bash ./PA1/utility/checkSubmitPA1.sh b09901000 pa1.tgz
```

Please note the path must be correct. If you are located in the \sim / directory, then './PA1/utility/checkSubmitPA1.sh' means the path \sim /PA1/utility/checkSubmitPA1.sh and b09901000 pa.tgz means the path \sim /b99901000 pa1.tgz

If you see "Permission denied", you should use following command to fix the problem.

```
chmod +x <checker name>
```

Your program will be graded by automatic grading script. Any mistake in the submission will cost at least 20% penalty of your score. Please be very careful in your submission.

Grading:

70% correctness (including submission correctness and implementation correctness) 20% file format and location 10% report

Frequently Asked Questions:

Q: Should we take care of the error handling issue of the testbench?

A: The testbench for this PA is correct. However, it is always better to have your error handling protection.

Q: Where is my EDA Union account and password?

A: NTU COOL > 成績 > Account & Password > 評語.

Handout #1 26 February 2024 Shao-Hua Sun

Q: How to log in to the server?

A: Please see the file "Week 1 Slides" in NTU COOL > 課程內容 > Recitation.

Someone cannot log in by

"ssh alg24sXXX@edaunion.ee.ntu.edu.tw -p <port>"

can try another command

"ssh -p <port> alg24sXXX@edaunion.ee.ntu.edu.tw".

Your EDA Union account (a.k.a Web ID) is "alg24sXXX" starts with the lower letter "a".

Q: What is the runtime limit for each case?

A: The runtime limit for each case is 10 minutes.

Q: Can I use C++11? Can I use Mac OS? Can I use ...?

A: Yes you can. However, we will use and only use the EDA Union server for evaluation. So whatever platform/setting you use, test on the EDA Union server before you submit your final version. The submission that fails to compile and/or run on the EDA Union server will be severely penalized.

Q: Runtime warning message: cannot get memory usage

A: Our code does not support checking memory usage on Mac OS. To check memory usage, compile and run the code on the EDA Union server.

Q: Quicksort runtime error message: segmentation fault

A: If the code is correct, this might be caused by stack overflow. Also, other memory-related issues might be reported in the error message. Enlarging the stack size would help. See section "Control the stack size" in the pdf.

Q: Do the submission files include the lib/ folder?

A: Yes. You are also allowed to modify the codes in lib/, so it's recommended that you attach them in your submission.

Q: Permission denied when using PA1 checker.

A: Type "chmod u+x ./utility/PA1 result checker" and run again.

Q: If there is any compile error (e.g. "Error: expecting string instruction after 'rep'")

A: "Google" it first or try to compile it on another server.

Handout #1 26 February 2024 Shao-Hua Sun

NOTE:

- 1. TA will check your source code carefully. Copying other source code can result in zero grade for all students involved.
- 2. Implementation correctness means to follow the guideline on the handout to write the codes. Wrong implementation will result in penalty even if the output is correct.
- 3. If you have any problem, please either contact TA, or ask question in Algorithms @ NTU on FaceBook. If your question is not answered, you should send email to TA, or attend TA hours.