```
Mon Apr 04 19:40:07 2022
ps5-readme.txt
   1: /*******************
   2: * readme
   3: * DNA Sequence Alignment
   5 •
   6: Name: Matthew Lorette Anaya
   7:
   8: Hours to complete assignment: 5
   9:
  10: /********************
      * Explain which approach you decided to use when implementing
       * (either recursive with memoization, recursive without memoization,
       * dynamic programming or Hirschberg\222s algorithm). Also describe why
  14:
       * you chose this approach and what its pros and cons are.
       *************************************
  15:
  16:
  17: Implementation of this program was done with the use of dynamic programmi
  18: and a matrix. I used the algorithm on the Princeton site in order to fill
  19: said matrix. In-order to find the alignment I used backtracking top-left
  20: bottom right, moving from the current matrix index to the next-lowest mat
  21: index. There was a certain case where if the diagonal was 1 higher than t
  22: downwards or rightwards option, diagonal was still the taken rout. In any
  23: case, depending on which direction I went, I either added a gap, or both
  24: letters, and incremented i and j counters to traverse back to the bottom
  25: right of the matrix.
  26:
  27:
  28:
  29:
  30: /*****************
      * Does your code work correctly with the endgaps7.txt test file?
  32:
      * This example should require you to insert a gap at the beginning
  33:
  34:
      * of the Y string and the end of the X string.
  36:
  37: Kinda confused here on what this question really is. The pdf is using exa
mple10.txt
  38: And this is asking for endgaps7.txt. Seems like there is a mix up of pdfs
between different years of this . So I'm going to use the what the HW pdf says
as there really isn't an example to compare to
  39: otherwise. Though it also says to put this all into a folder named ps3, w
hich is definitely incorrect.
  40:
  41: Input:
  42:
             Ê./EDistance < ./sequence/example10.txt</pre>
  43:
  44: Expected output:
  45:
  46:
             Edit distance = 7
  47:
             AT1
  48:
             AA0
  49:
             C-2
  50:
             AA0
  51:
             GG0
  52:
             TG1
  53:
             TT0
  54:
             A-2
  55:
             CC0
             CA1
  56:
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  57:
  58: What happened:
  59:
  60:
            Edit distance = 7
            A T 1
  61:
            A A 0
  62:
  63:
            C - 2
  64:
            A A 0
  65:
            G G 0
            T G 1
  66:
  67:
            Т Т О
  68:
            A - 2
            C C 0
  69:
  70:
            C A 1
  71:
            Execution time is 0.00094 seconds.
  72:
  73:
  74:
  75:
  76:
  77: /*******************
  78: * Look at your computer\222s specs in the settings.
  79: * How much RAM does your computer have and explain what this means?
  81:
  82: My Mac has 16gb of RAM. Random access memory gives applications a place t
  83: store and access data on a short-term basis. It stores the information yo
ur
  84: computer is actively using so that it can be accessed quickly.
  85:
  86:
  87: /*****************
      * For this question assume M=N. Look at your code and determine
         approximately how much memory it uses in bytes, as a function of
  90:
        N. Give an answer of the form a * N^b for some constants a
  91:
         and b, where b is an integer. Note chars are 2 bytes long, and
  92: *
        ints are 4 bytes long.
  93:
  94:
        Provide a brief explanation.
  95:
  96:
      ^{\star} What is the largest N that your program can handle if it is
  97: * limited to 8GB (billion bytes) of memory?
  100: N^2 is the area of the matrix, the number of integer slots that need to b
 101: filled in. 4 is the size of an integer in bytes.
 102:
 103: a = 4
 104: b = 2
 105: largest N = ^44,721
 106:
 107: Explanation:
            4 * 44,721^2 = 7,999,871,364 just shy of 8gb.
 108:
 109:
 111: * Run valgrind if you can and attach the output file to your submission.
 112: * If you cannot run it, explain why, and list all errors you\222re seein
a.
 113:
      * If you can run it successfully, does the memory usage nearly match tha
      * found in the question above?
 114:
 115: * Explain why or why not.
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 116: /******************************
 118: -----
 119: n time(i) total(B) useful-heap(B) extra-heap(B) sta
cks(B)
 120: -
 121: 67 6,808,807,498 3,085,122,584 3,084,577,402 545,182
  Ω
 122: 68 6,870,407,882 3,146,681,624 3,146,125,562
  0
 123: 69 30,570,148,197 3,201,904,240 3,201,338,395 565,845
  0
 124:
 125: It does not, its actually quite different and I'm not entirely sure as to
why. Not
 126: Sure if I'm reading valgrind output wrong or my equation is.
 127:
 128:
 130: * For each data file, fill in the edit distance computed by your
 131: \star program and the amount of time it takes to compute it.
 133: \star If you get segmentation fault when allocating memory for the last
 134: * two test cases (N=20000 and N=28284), note this, and skip filling
 135: * out the last rows of the table.
 137:
 138: data file
                    distance
                             time (seconds)
 139: -----
 140: ecoli2500.txt 118
141: ecoli5000.txt 160
                                         0.125216
                                         0.334861
                     194
 142: ecoli7000.txt
                                         0.521017
 143: ecoli10000.txt 223
144: ecoli20000.txt 3135
145: ecoli28284.txt 8394
                                   1.4272
                                 74.6052
                     8394
                                 177.645
 145: ecoli28284.txt
 146:
 148: * Here are sample outputs from a run on a different machine for
 149: * comparison.
 151:
              distance
 152: data file
                               time (seconds)
 154: ecoli2500.txt 118
155: ecoli5000.txt 160
                             0.171
                                   0.529
 156: ecoli7000.txt
                      194
                                   0.990
                    223
3135
 157: ecoli10000.txt
                                   1.972
 158: ecoli20000.txt
                                   7.730
 159:
 160:
 161:
 162: /***********************************
      * For this question assume M=N (which is true for the sample files
 163:
 164:
        above). By applying the doubling method to the data points that you
 165:
        obtained, estimate the running time of your program in seconds as a
      * polynomial function a * N^b of N, where b is an integer.
 166:
 167:
        (If your data seems not to work, describe what went wrong and use
        the sample data instead.)
 168:
 169:
 170: * Provide a brief justification/explanation of how you applied the
 171: * doubling method.
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172:
173: * What is the largest N your program can handle if it is limited to 1 \,
174: * day of computation? Assume you have as much main memory as you need.
176: a =
177: b =
178: largest N =
179:
180: /
182: * Did you use the lambda expression in your assignment? If yes, where
183: * (describe a method or provide a lines numbers)
185: No
186:
187:
188:
189: *******************
190: * List whatever help (if any) you received from the course TAs,
191: * instructor, classmates, or anyone else.
193:
194: N/a
195:
197: * Describe any serious problems you encountered.
199:
200:
201:
202: /*******************
203: * List any other comments here.
204:
  *******************
205:
```