Intro to Algorithms: Homework #9

Due on April 22, 2021

Prof. Zaki

Jared Gridley

```
0 5 20 -10 10 5 45 55 00
Max Sum (5):
    L[0] - 6
 O(1) \times O(n) = O(n)
```

Figure 1: Page 1

Lab Results:

Results:

cox1-protein.fasta
edit distance = 230

alignment:

M-VQRWLYSTNAKDIAVLYFMLAIFSGMAGTAMSLIIRLELAAPGSQYLHGNSQLFNVLVVGHAVLMIFFLVMPALIGGFGNYLLPLMIGATDTA FPRINNIAFWVLPMGLVCLVTSTLVESGAGTGWTVYPPLSSIQAHSGPSVDLAIFALHLTSISSLLGAINFIVTTLNMRTNGMTMHKLPLFVWSI FITAFLLLLSLPVLSAGITMLLLDRNFNTSFFEVSGGGDPILYEHLFWFFGHPEVYILIIPGFGIISHVVSTY-S-KKPVFGEISMVYAMASIGL LGFLVWSHHMYIVGLDADTRAYFTSATMIIAIPTGIKIFSWLATIHGGSIRLATPMLYAIAFLFLFTMGGLTGVALANASLDVAFHDTYYVVGHF HYVLSMGAIFSLFAGYYYWSPQILGLNYNEKLAQIQFWLIFIGANVIFFPMHFLGINGMPRRIPDYPDAFAGWNYVASIGSFIATLSLFLFIYIL YDQLVNGLNNKVNNKSVIYNKAPDFVESNTIFNLNTVKSSSIEFLLTSPPAVHSFNTP-AVQS

MFADRWLFSTNHKDIGTLYLLFGAWAGVLGTALSLLIRAELGQPG--NLLGNDHIYNVIVTAHAFVMIFFMVMPIMIGGFGNWLVPLMIGAPDMA
FPRMNNMSFWLLPPSLLLLLASAMVEAGAGTGWTVYPPLAGNYSHPGASVDLTIFSLHLAGVSSILGAINFITTIINMKPPAMTQYQTPLFVWSV
LITAVLLLLSLPVLAAGITMLLTDRNLNTTFFDPAGGGDPILYQHLFWFFGHPEVYILILPGFGMISHIV-TYYSGKKEPFGYMGMVWAMMSIGF
LGFIVWAHHMFTVGMDVDTRAYFTSATMIIAIPTGVKVFSWLATLHGSNMKWSAAVLWALGFIFLFTVGGLTGIVLANSSLDIVLHDTYYVVAHF
HYVLSMGAVFAIMGGFIHWFPLFSGYTLDQTYAKIHFTIMFIGVNLTFFPQHFLGLSGMPRRYSDYPDAYTTWNILSSVGSFI---SL-------TAV-ML-----MIFMI-WEA--F-ASKRKVLMVEEPSMNLEWLYGCPPPYHTFEEPVYMKS

cox1-dna.fasta
edit distance = 634

alignment:

TACAA--C-GTTATCGTCACAGCCCATGC-ATTTGTAATAATCTTCTTCATAGTAATACCCATCATAATCGGAGGCTTTGGCAACTGACTAGTTC $\tt CCCTAATAATCGGTGCCCCCGATATGGCGTTTCCCCGCATAAACAACATAAGCTTCTGACTCTTACCTCCTCTCTCCTACTCCTGCTCGCAT-C$ TGCTATAGTGGAGGCCGG-AGCAGGAACAGGTTGAACAGTCTACCCTCCCTTAGCAGGGAACTA-CTCCCACCCTGGAGCCTCCGTAGACCTAAC CATCTTCTCCTT-ACACCTAGCAGGTGT-CTCCTCTATCTTAGGGGCCCATCAATTTCATCACAACAATTATCAATATAAAACCCCCTGCCATAAC $\tt CCAATACCAAACGCCCCTCTTCGTCTGATCCGTCCTAATCACAGCAGTCCT-ACTTCTCCTATCTCCCCAGTCCTAGCTGCC-ATCACTAT$ ACTACTAACAGACCGCAACCTCAACACCACCTTCTTCGA-CCCCG-CCGGAGGAGGAGACCCCATTCTATACCAACACCTATTCTGATTTTTCGGTCACCCTGAAGTTTATATTCTTATCCTACCAGGCTT-CGGAATAATCTCCCATATTGTA-ACTT-ACTACTCCGGAAAAAAAGAACCATTTGGAT ${\tt ACGAGCATATTTCACCTCCGCTACCATAATCATCGCTATCCCCACCGGCGTCAAAGTATTTAGCTGACTCGCCACACTCCACGGAAGCAATATGA}$ ${\tt AAT-GA-TCTGCT-GCAGTGCTCTGAGCCCTAGGATTCATCTTTCTTTTCACCGTAGGTGGCCTGACTGGCATTGTATTAGCAAACTCATCACTA}$ GACATCGTACTACACGACACGTACTACGTTGT-AGCCCACTTCCACTATGTCCTATCAATAGGAGCTGTATTTGCCATCAT-A-GGAGGCTTCAT TTCTTCCCACACACTTTCTCGGCCTATCCGGAATGCCCCG-ACGTTACTCGGACTACCCCGATGCATACACCACATGAAACATCCTATCATCTG TA-GGCTC-ATTCATTCTCTAACAGCAGTAATATTAATAATT-TTCATG-ATT-TGAGAAGCCTTCGCTT-CG-AAGCGAAAAGTCCTAAT-AG A-CATAAAATCTAGA


```
a) Base Coses
3) Recusive Definition:
4) Forward Solution
          L [F, F, F, F, T, F, T, F, T, ]
         L - [len(5)]
         L[1] = del (4(1))
```

Figure 2: Page 2

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                                                                                                                                 0
                                                                                                                                 00
  2) Base Coses;
           L[1,0] - 1 iFF
   H) Forward
                      [F(*[i] -= g[o]);
L[i][o] = 1
                 if (x[i] - y[i]);
L[i][j] - L[i-i][j-q + 4
d(nm)
                  rt:1[j] - 6
            result . L[40[07][00[1]]
           while ( L[ind[0]-1][ind[1]-1] >0):
```

Figure 3: Page 3

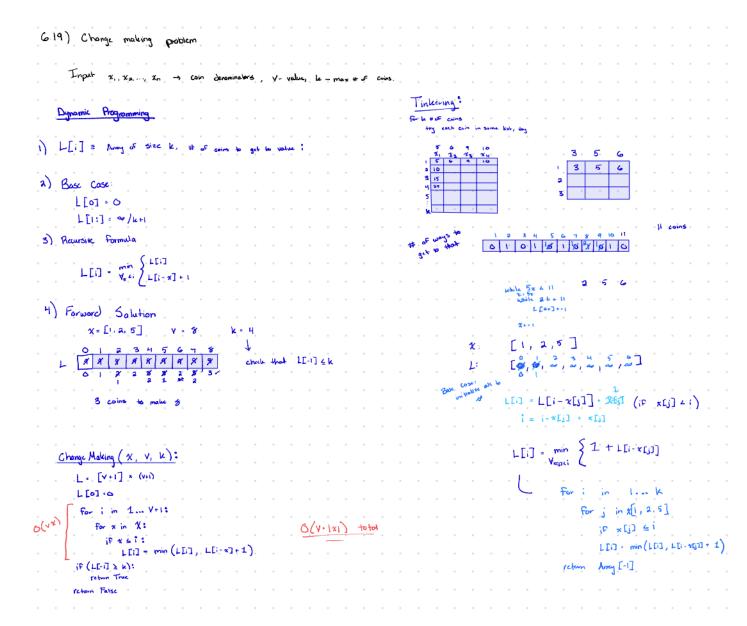


Figure 4: Page 3