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College of CUNY Problem 1: Coin Change (a) The  
greedy algorithm for coin change can be described  
as:  $G(n) = 1 + G(n - c)$  where  $c$  is the largest coin value  
less or equal to  $n$ .  $G(n)$  if  $n > 0$  then let  $c$  be largest coin  
value  $n$  return  $1 + G(n - c)$  else return 0*

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Introduction to Bioinformatics Lopresti BioS 10  
October 2010 Slide 13. HHMI. Howard Hughes Medical  
Institute. Sequencing a Genome. Most genomes are  
enormous (e.g., 10<sup>10</sup> base pairs in case of human).*

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*Current sequencing technology, on the other hand, only allows biologists to determine ~10<sup>3</sup> base pairs at a time.*

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*Introduction to Bioinformatics Algorithms Homework 4 Solution Saad Mneimneh, Computer Science, Hunter College of CUNY Problem 1: Spliced alignments (a) Consider the Exon Chaining problem in the case where all intervals have the same weight. For this setting, the best chain is obviously the one that has the maximum number of non-overlapping intervals.*

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*An Introduction to Bioinformatics Algorithms, MIT Press, Cambridge, Mass. (slides below from [www.bioalgorithms.info](http://www.bioalgorithms.info)) Molecular Biology (Ch 3) DNA Mapping (Ch 4) Brute Force Motif Searching (Ch 4) Genome Rearrangements (Ch 5) Alignment (Ch 6) Edit*

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*1998 (10) is an excellent practical guide to bioinformatics; Mount, 2001 (76) excels in showing the connections between biological problems and bioinformatics techniques; and Bourne and Weissig,*



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*2002 (15) focuses on protein*

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