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Sequencing a Genome. Most
genomes are enormous (e.g., 10¹⁰

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base pairs in case of human). Current sequencing technology, on the other hand, only allows biologists to determine ~10³ base pairs at a time.

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1998 (10) is an excellent practical

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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

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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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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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represents a great opportunity to bolster knowledge on the biological data science since information grew up exponentially, it is necessary to count with tools offered by bioinformatics to handle this kind of information.

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