

Exercise 1 The human genome consists of 23 pairs of chromosomes. Perform a literature search to find out how many chromosomes there are in the genomes of the following organisms:

- (a) Chimpanzee (*Pan troglodytes*)
- (b) Dog (*Canis familiaris*)
- (c) Mouse (*Mus musculus*)
- (d) Chicken (*Gallus gallus*)
- (e) Cow (*Bos taurus*)
- (f) Zebrafish (*Danio rerio*)
- (g) Yeast (*Saccharomyces cerevisiae*)
- (h) Rice (*Oryza sativa*).

Exercise 2 Give the reverse complement (i.e. Watson-Crick complementary sequence in reverse order) of the following sequences:

- (a) TAGCTGATCTATCTCGCGCGCCGCATTATCGTATTCTCAGTGATCTCT
- (b) ATGGATTATCCTTGCAGCGCGCATTTCGCTCATCTAGGTGTCTATGC

Exercise 3 Compute the binomial coefficients nC_r for $0 \leq n \leq 5$ and $0 \leq r \leq n$

Exercise 4 Implement the quick sort algorithm in your favorite programming language.

Exercise 5 Write an algorithm to calculate selected Fibonacci number in your favorite programming language.

Exercise 6 Write an algorithm to compute the entropy of a DNA sequence. Implement the algorithm in your favorite programming language. Obtain a DNA sequence and use your program to calculate the entropy of the sequence. Report the results.

Exercise 7 Write a program to read in FASTA file, reverse complement the sequence(s) in the file and write the results into another well-formed FASTA file. Do not allow the lines of the output file to exceed 60 characters in length (except possibly for the defline)