

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) ATGGATTTATCCTTGCGCGCGCATTTTCGCTCATCTAGGTGTCTATGC

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)*