

BIOL 5153, Practical Programming for Biologists
Assignment #5
Due Friday, 20 March 2020, by 11:59 PM.

Write a Python script called 'parseGFF.py' that opens and reads the file, 'watermelon.gff'. Extract the gene name, store it in a list, then print an alphabetized list of the genes at the end.

For the first entry, you would extract 'cox1' and print it. You can skip all the 'similar to . . . ' entries.

We will continue to expand and update this script in the coming weeks.