RGI PYTHON SCRIPT INSTRUCTION

TIANCI LI

There are five arguments that are required

-h, --help show this help message and exit

[-i\_card\_json] path to card.jason

[-i\_model] path to homology model

[-i\_main] path to fasta file

[-o\_prefix] ouput prefix

[-t] read,contig,protein,wgs

If couldn’t open maybe try different shebang.

The output file will be “prefix”.json and “prefix”.txt. All other files that contains “temp” are useless.