INSTRUCTIONS TO USE LHT_CI.py

The file LHT_CI.py is a Python3 program that performs the calculations.

1. Data should be in a .csv file with only three columns, for instance:

n	h
50	0
50	0
50	0
49	0
49	0
49	0
49	0
48	0
46	0
44	28
44	28
44	10
43	118
41	111
	50 50 50 49 49 49 49 48 46 44 44 44

The **first column** are the units of time. The **second column** contains number of individuals alive at that unit of time. The **third column** contains offspring production in that unit of time.

2. Inside the code, modify the alpha required and change the file name:

```
alpha = 0.05  # 1-alpha is the confidence level of CI,
fnam = "test_data.csv"  # Change file name.
# (Put data file and Python program in same folder)
```

3. Run the program, an example output is:

Initial number of individuals N: 50 ----(Initial number of individuals)

Offspring size K: 2430 ----(Total offspring)

R0: 48.6 ----(R0, the basic reproductive number)

Longevity: [28.34 302.0644 23.5226 33.1574] -----(mean variance and CI for longevity)

Generation time: [26.884 151.77 26.3945 27.374] -----(mean variance and CI for gen. time)

r: [0.20209, 0.198870, 0.2055] ------(mean and CI for r)

lambda: [1.223964994, 1.22002366, 1.22814391] ------(mean and CI for lambda)

Saved to: test data added.csv ------(outcome was saved to this file)