## Dale Heck

CS 417/ Dr. Kranz

Assignment 3 outputs

## **Question 1**

```
In [2]: runfile('/Users/loganheck/untitled0.py', wdir='/Users/loganheck')
Count for 'Right' (case-insensitive): 7
Count for 'Rights' (case-insensitive): 5
Top 25 words:
the -> 133
of -> 106
to -> 70
and -> 61
for -> 33
import -> 29
a -> 27
our -> 27
in -> 22
their -> 22
has -> 21
us -> 20
he -> 19
declaration -> 18
archives -> 17
nbsp -> 17
them -> 15
by -> 13
documents -> 13
is -> 13
these -> 13
have -> 12
that -> 12
all -> 11
it -> 11
```

## **Question 3**

```
In [7]: runfile('/Users/loganheck/Documents/Intro to object
  oriented programming/untitled2.py', wdir='/Users/loganheck/
  Documents/Intro to object oriented programming')
  Enter a string:
  3
  You entered: 3
  Number of characters: 1

In [8]: runfile('/Users/loganheck/Documents/Intro to object
  oriented programming/assign.3, question3.py', wdir='/Users/
  loganheck/Documents/Intro to object oriented programming')
  Enter a string:
  123445
  You entered: 123445
  Number of characters: 6
```

## **Question 4**

```
In [9]: runfile('/Users/loganheck/Documents/Intro to object
oriented programming/untitled3.py', wdir='/Users/loganheck/
Documents/Intro to object oriented programming')
Original DNA (first 60 chars):
TGGCCAGTAGATCTTCCCAACATAGCCTAGCTGGACATATTCACTAAACCGAACAATCTA
Mutated DNA (first 60 chars):
TGGCCAGTGGATCTTCCCAACATAGTCTAGCTGTAGATAATCACTACACCGAACAATCTA
Before mutation:
Frequency Table (showing first 16 positions):
A: [0, 0, 0, 0, 0, 1, 0, 1, 0, 1, 0, 0, 0, 0, 0]
C: [0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1]
G: [0, 1, 1, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0]
T: [1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 1, 1, 0]
After mutation:
Frequency Table (showing first 16 positions):
A: [0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0]
C : [0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1]
G: [0, 1, 1, 0, 0, 0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 0]
T: [1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 1, 1, 0]
```