# **UNIX Assignment**

## **Data Inspection**

#### Attributes of fang\_et\_al\_genotypes

here is my snippet of code used for data inspection

By inspecting this file I learned that:

- 1. point 1
- 2. point 2
- 3. point 3

or

- point 1
- point 2
- point 3

### Attributes of snp position.txt

here is my snippet of code used for data inspection

By inspecting this file I learned that:

- 1. point 1
- 2. point 2
- 3. point 3

or

- point 1
- point 2
- point 3

## **Data Processing**

#### **Maize Data**

here is my snippet of code used for data processing

Here is my brief description of what this code does

### **Teosinte Data**

here is my snippet of code used for data processing

Here is my brief description of what this code does