## visualization

May 13, 2024

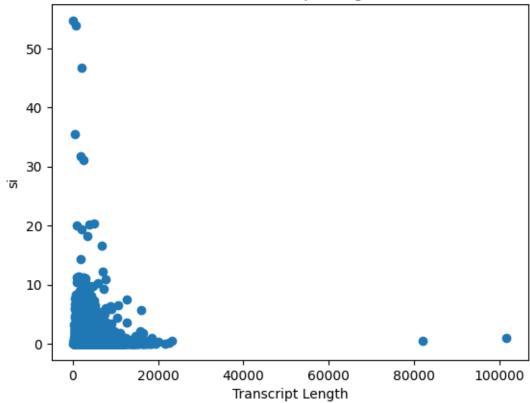
## 1 Visualizations and Analysis for the results obtained by Gibbs Sampling

```
[]: import numpy as np
     import matplotlib.pyplot as plt
[]: headers = ["Gene Name", "Transcript Length", "si", "koni", "koffi", "skoffi", "
     data = np.genfromtxt("result.txt", delimiter="\t", skip_header=1, dtype=None,_
      ⇔encoding=None, unpack=True)
     gene_names = data[0].astype(str)
     si = data[2].astype(float)
     transcript_length = data[1].astype(float)
     koni = data[3].astype(float)
     koffi = data[4].astype(float)
     skoffi = data[5].astype(float)
     exi = data[6].astype(float)
     #plot si vs transcript length
     plt.scatter(transcript_length, si)
     plt.xlabel("Transcript Length")
     plt.ylabel("si")
     plt.title("si vs Transcript Length")
     plt.show()
     # find the correlation between si and transcript length
     correlation = np.corrcoef(transcript_length, si)
     print("Correlation between si and transcript length: ", correlation[0,1])
     # find the correlation between koni and koffi
     correlation = np.corrcoef(koni, koffi)
     # plot koni vs koffi
     plt.scatter(koni, koffi)
     plt.xlabel("koni")
     plt.ylabel("koffi")
     plt.title("koni vs koffi")
```

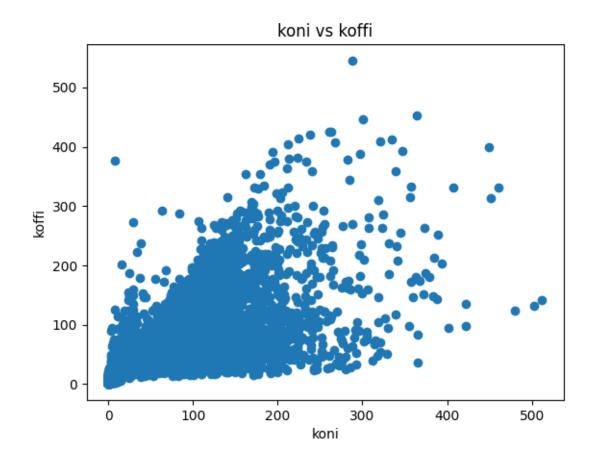
```
plt.show()
print("Correlation between koni and koffi: ", correlation[0,1])
# print covariance matrix between koni and koffi
covariance = np.cov(koni, koffi)
print("Covariance matrix between koni and koffi: ", covariance)
# make a correlation matrix between si, koni, koffi, skoffi, exi
data = np.array([transcript_length, si, koni, koffi, skoffi, exi])
correlation = np.corrcoef(data)
print("Correlation matrix between si, koni, koffi, skoffi, exi:\n ", u
 ⇔correlation)
# print the pairs with correlation higher than 0.5
for i in range(data.shape[0]):
    for j in range(i+1, data.shape[0]):
        if correlation[i,j] > 0.5:
            print("Correlation between ", headers[i+1], " and ", headers[j+1], "]

¬" is: ", correlation[i,j])
```

## si vs Transcript Length



Correlation between si and transcript length: 0.005717807729022441



```
Correlation between koni and koffi: 0.8055642726213191
Covariance matrix between koni and koffi: [[2487.67251349 1823.73771177]
 [1823.73771177 2060.30329132]]
Correlation matrix between si, koni, koffi, skoffi, exi:
  [[ 1.
                0.00571781 0.00285494 0.00265964 0.01047814 0.0052909 ]
 [ 0.00571781 1.
                         -0.06186055 0.03469592 0.44635901 0.86715441]
 [ 0.00285494 -0.06186055 1.
                                      0.80556427 -0.16189324 -0.06129007]
 [ 0.00265964  0.03469592  0.80556427  1.
                                                -0.15264258 -0.00414808]
 [ 0.01047814  0.44635901 -0.16189324 -0.15264258  1.
                                                             0.52235117]
 [ 0.0052909
              0.86715441 -0.06129007 -0.00414808 0.52235117 1.
                                                                       ]]
Correlation between si and exi is: 0.8671544101189494
Correlation between koni and koffi is:
                                          0.8055642726213191
Correlation between skoffi and exi is: 0.5223511698997131
```