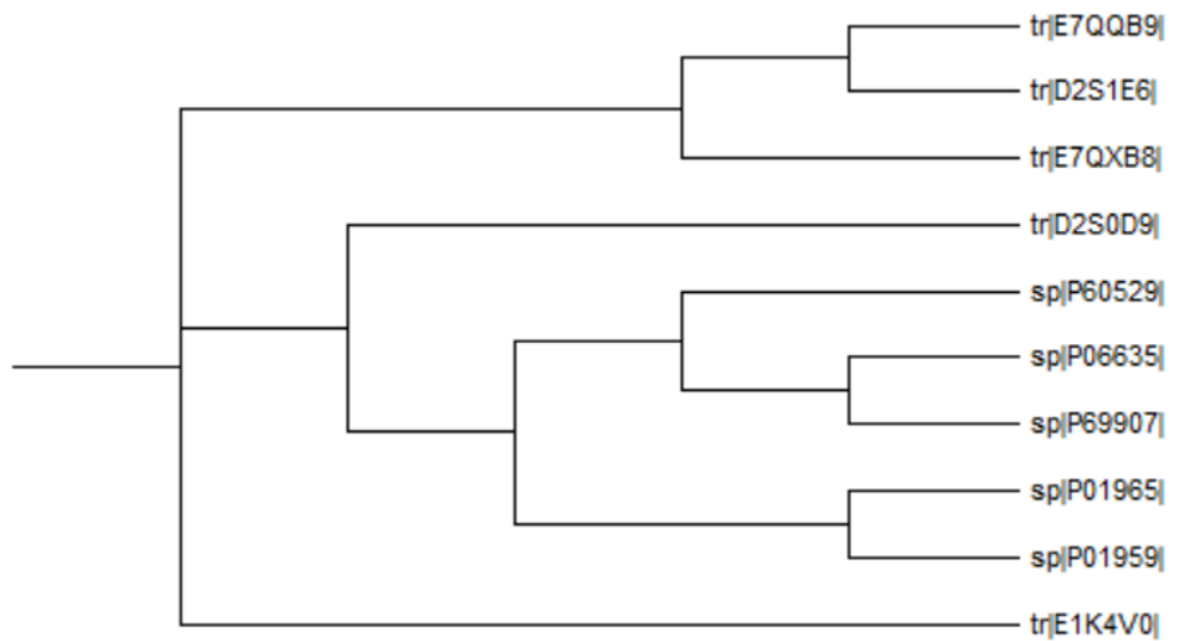


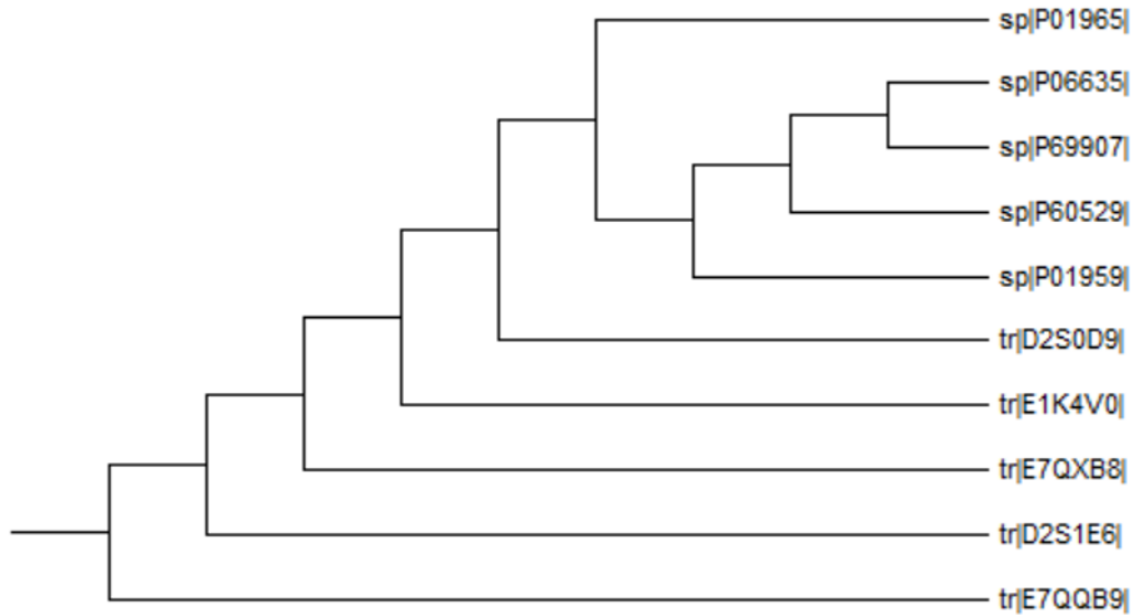
1. For file

tim.dat-

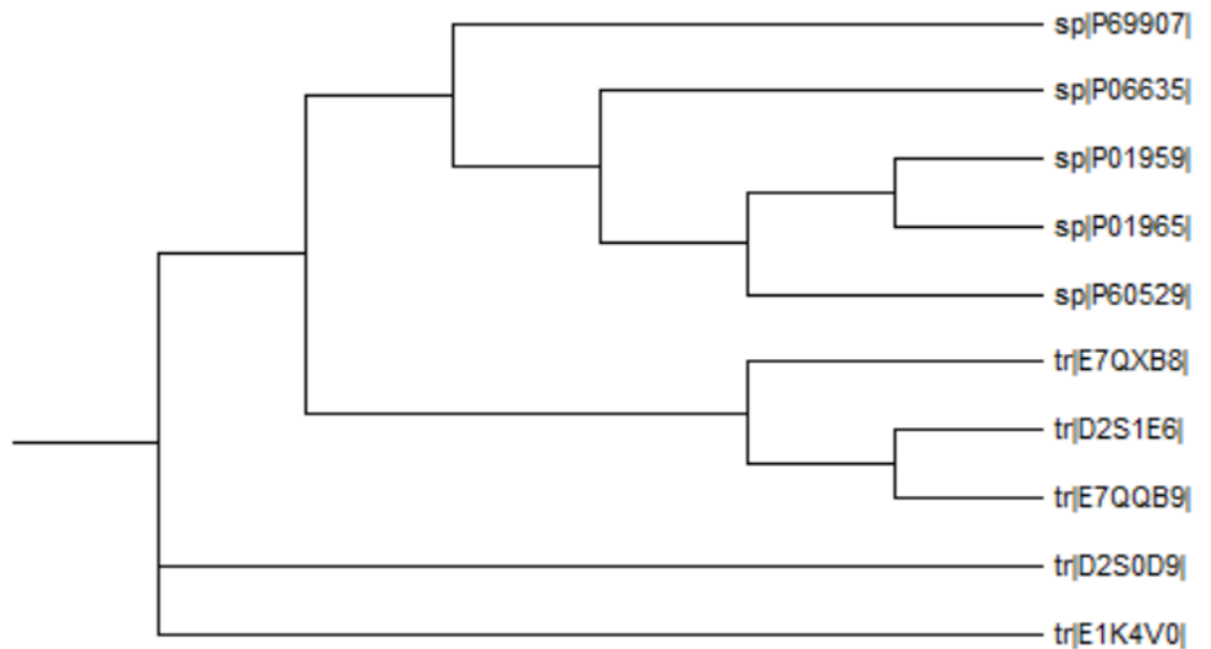
Outtree obtained from proml:



The output from `consense`:

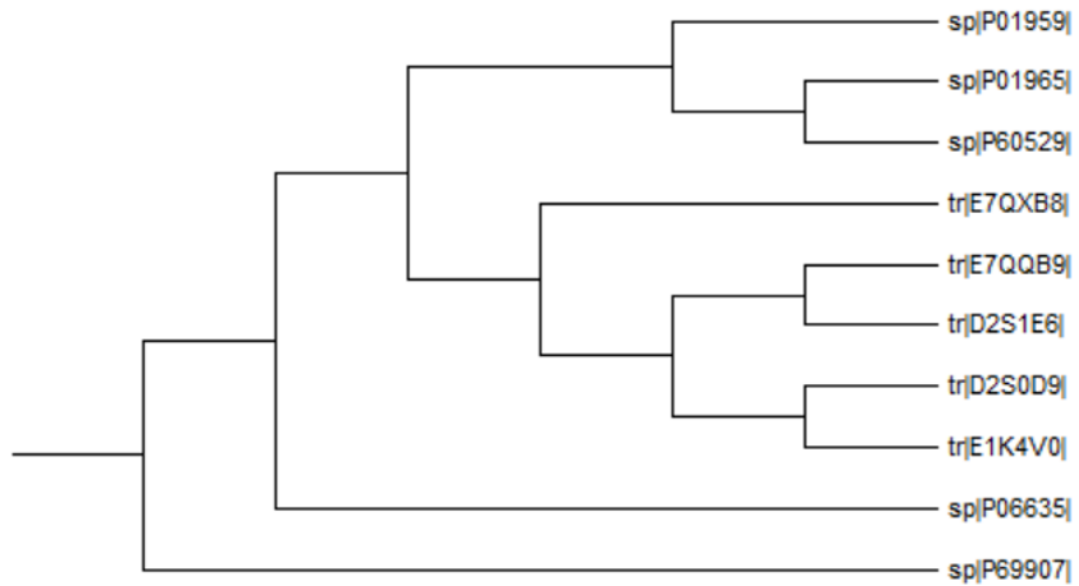


The output from neighbor:



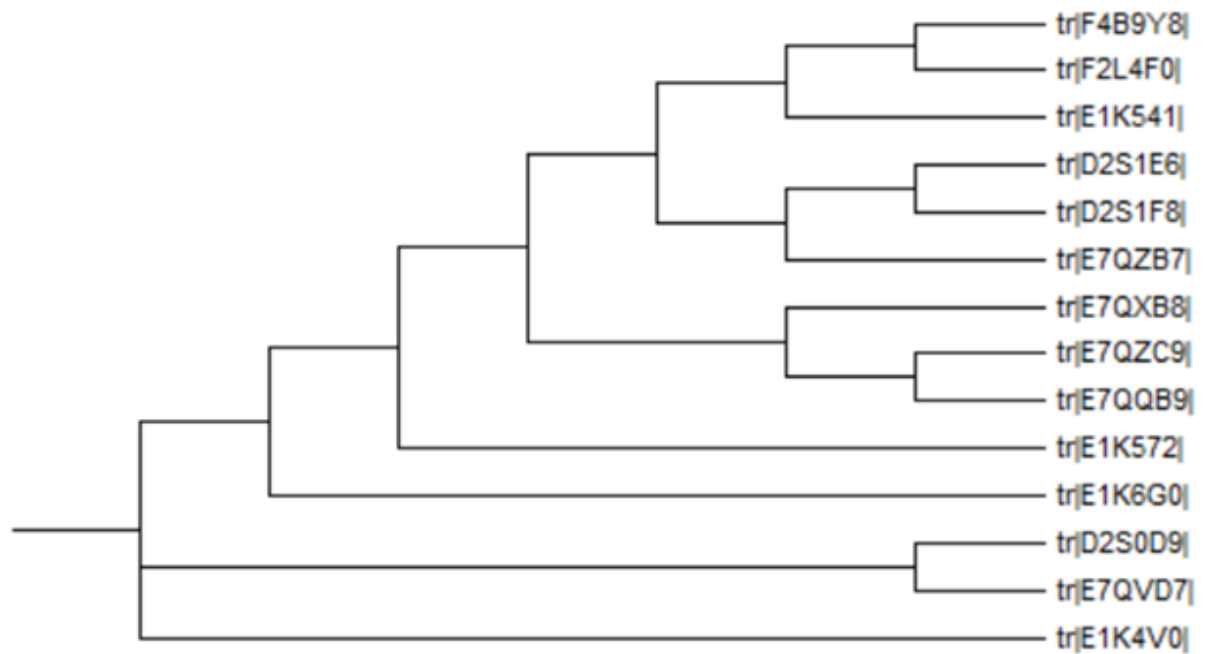
Input file work1-nj-tree:

The output from `consense`:

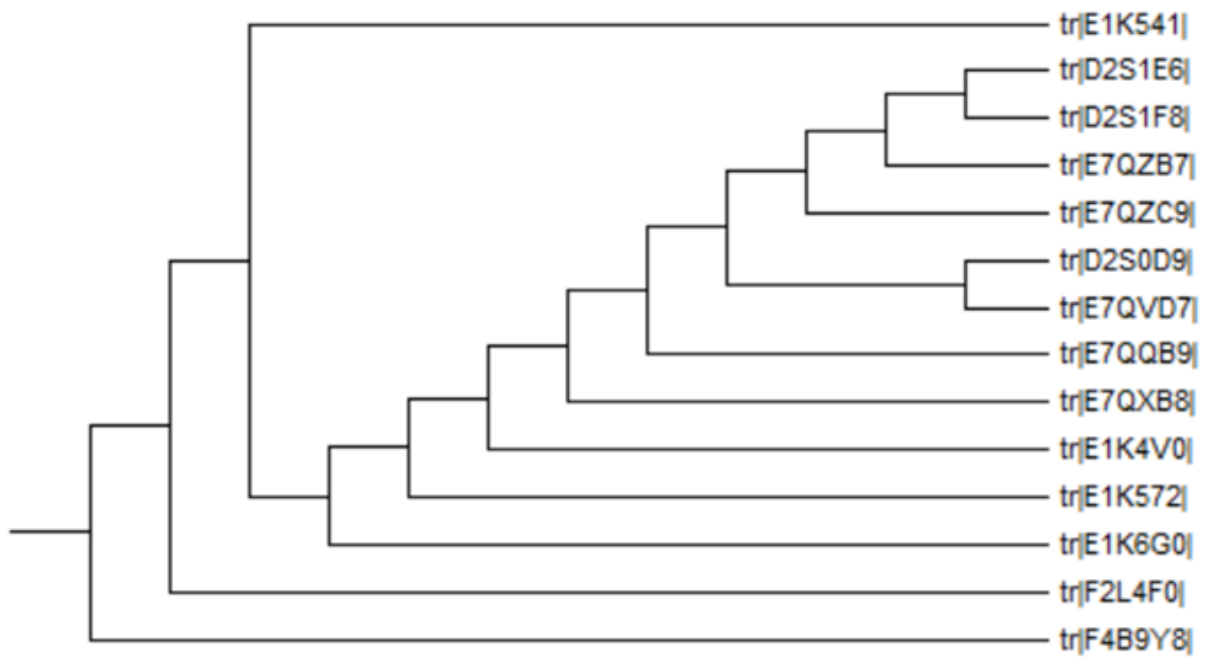


For file **tim.dat.txt**

Output obtained from proml:

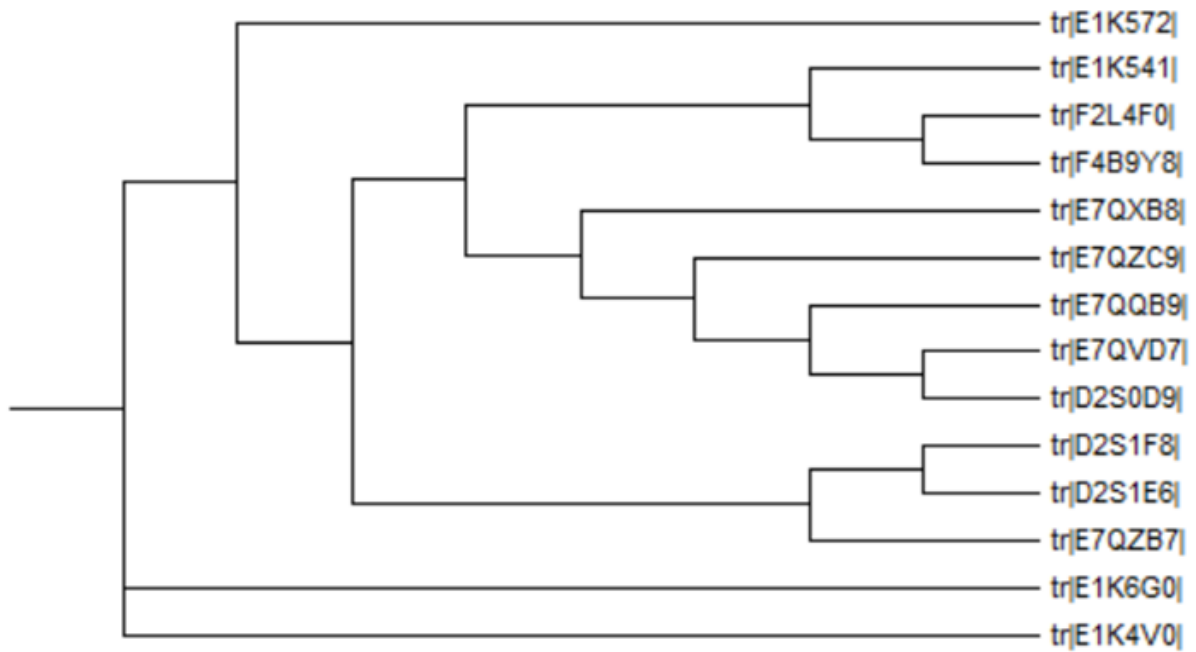


Output from consensus:



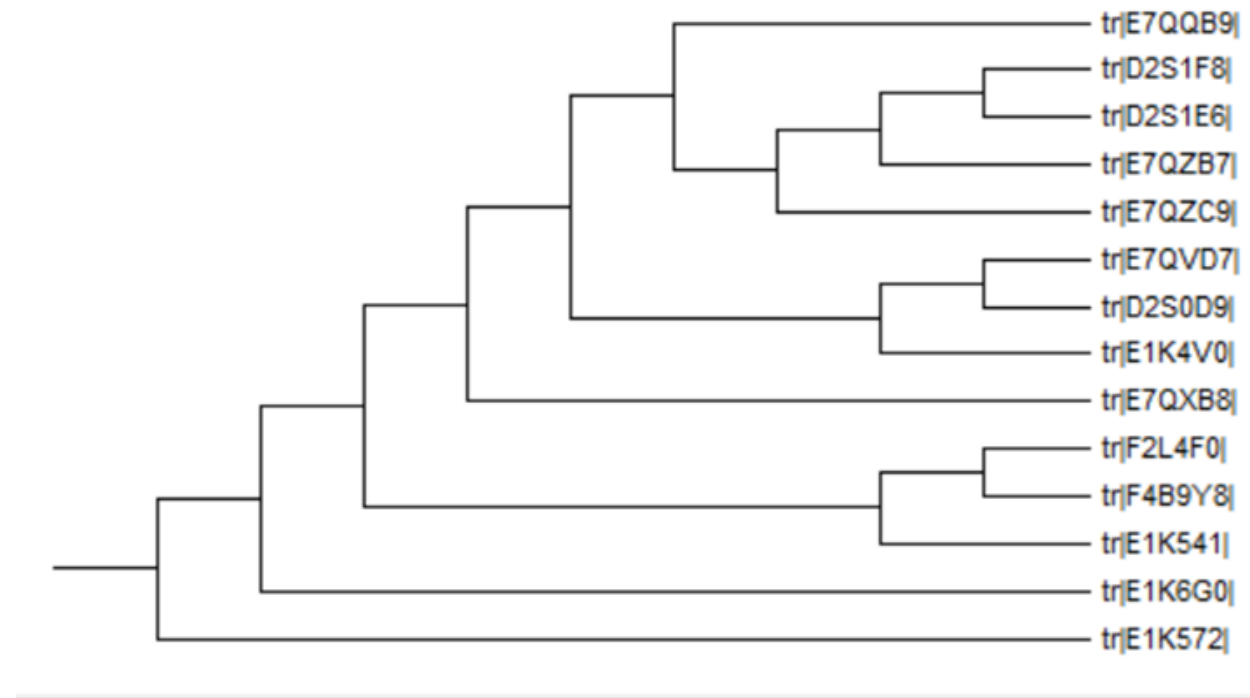
Input file work2-protldist:

The output from neighbor:



Input file work-nj-tree:

The output obtained from `consense`:



2.

```
import math as m
```

```
import pandas as pd
```

```
import numpy as np
```

```
def Q2(seq):
```

AA = ['A', 'C', 'D', 'E', 'F', 'G', 'H', 'I', 'K', 'L', 'M', 'N', 'P', 'Q', 'R', 'S', 'T', 'V', 'W', 'Y']

```
N, L = len(seq), len(seq[0])
```

```
a_m, w_m = [[0 for _ in range(L)] for __ in range(20)], [[0.000 for _ in range(L)] for __ in range(20)]
```

```
p = [str(_+1) for _ in range(27)]
```

```

for i in range(N):
    for j in range(L):
        a_m[AA.index(A[i][j])][j] += 1
data1 = np.array(a_m)

for k in range(20):
    for l in range(L):
        w_m[k][l] = float('%0.2f'%(math.log((a_m[k][l]+0.05)/(0.05 * (N+1))))))
data2 = np.array(w_m)

print('Weight Matrix for the given sequences:\nrows ---> amino acids\tand\tcolumns --->
positional occurances')

print(pd.DataFrame(data2, AA, p))

return pd.DataFrame(data2, AA, p)

```

```

Seq = ['MVLSPADKTNVKGKVGAAHAGEYGAAAW',
'MKRLPADPPCVKGKVKAKAGDYGATTW',
'MALSAADKTNVKS KVGGHAGEYGAATS',
'MVLSAADKTNVKS KAGGNAGEWWAAAW',
'MVLSAADKTNVKS KVLNAGEFGAAAW',
'ALLPIRTTYHKK CASGHIPEEKDLNNV',
'DEASSLKGH HIKKLEADALLIPLSASS']

```

Q2(Seq)

	1	2	3	4	5	6	7	8	9	10	...	18	19	20	21	22	23	24	25	26	27
A	0.97	0.97	0.97	-2.08	2.03	2.54	-2.08	-2.08	-2.08	-2.08	...	0.97	2.54	-2.08	-2.08	-2.08	-2.08	2.54	2.54	2.03	-2.08
C	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	...	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08
D	0.97	-2.08	-2.08	-2.08	-2.08	-2.08	2.54	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	0.97	-2.08	0.97	-2.08	-2.08	-2.08	-2.08
E	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	0.97	2.54	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08
F	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	-2.08
G	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	-2.08	-2.08	...	-2.08	-2.08	2.54	-2.08	-2.08	2.32	-2.08	-2.08	-2.08	-2.08
H	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	1.63	...	1.63	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08
I	-2.08	-2.08	-2.08	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	-2.08	...	0.97	-2.08	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08
K	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	0.97	2.32	-2.08	-2.08	...	0.97	-2.08	-2.08	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	-2.08
L	-2.08	0.97	2.54	0.97	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	...	-2.08	0.97	0.97	-2.08	-2.08	0.97	0.97	-2.08	-2.08	-2.08
M	2.54	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08
N	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	2.32	...	1.63	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	0.97	-2.08
P	-2.08	-2.08	-2.08	0.97	1.63	-2.08	-2.08	0.97	0.97	-2.08	...	-2.08	0.97	-2.08	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	-2.08
Q	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08
R	-2.08	-2.08	0.97	-2.08	-2.08	0.97	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08
S	-2.08	-2.08	-2.08	2.54	0.97	-2.08	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	-2.08	0.97	1.63
T	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	0.97	2.32	-2.08	...	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	1.63	-2.08
V	-2.08	2.03	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97
W	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	...	-2.08	-2.08	-2.08	-2.08	0.97	0.97	-2.08	-2.08	-2.08	2.32
Y	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	-2.08	0.97	-2.08	...	-2.08	-2.08	-2.08	-2.08	2.03	-2.08	-2.08	-2.08	-2.08	-2.08

20 rows × 27 columns