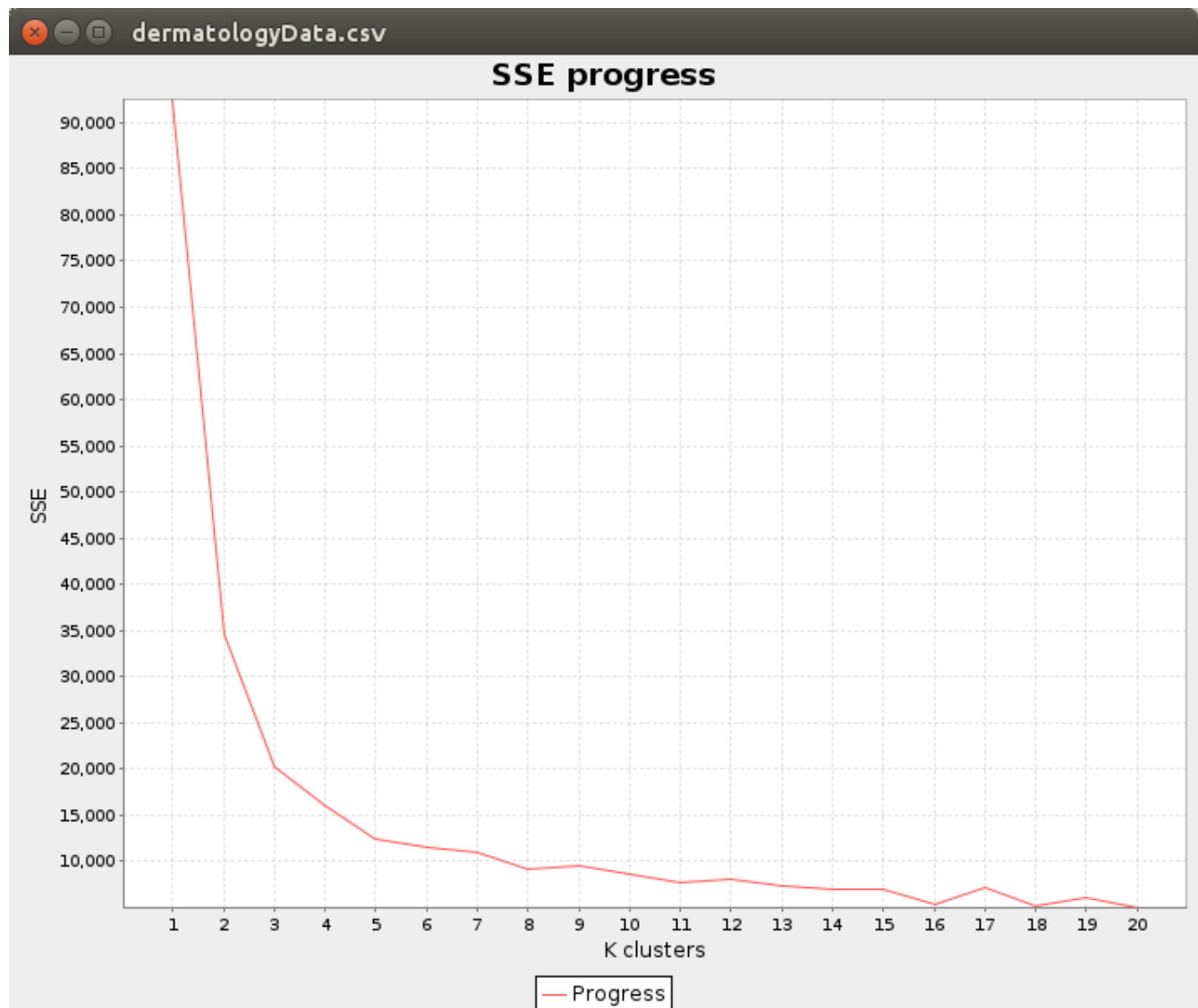


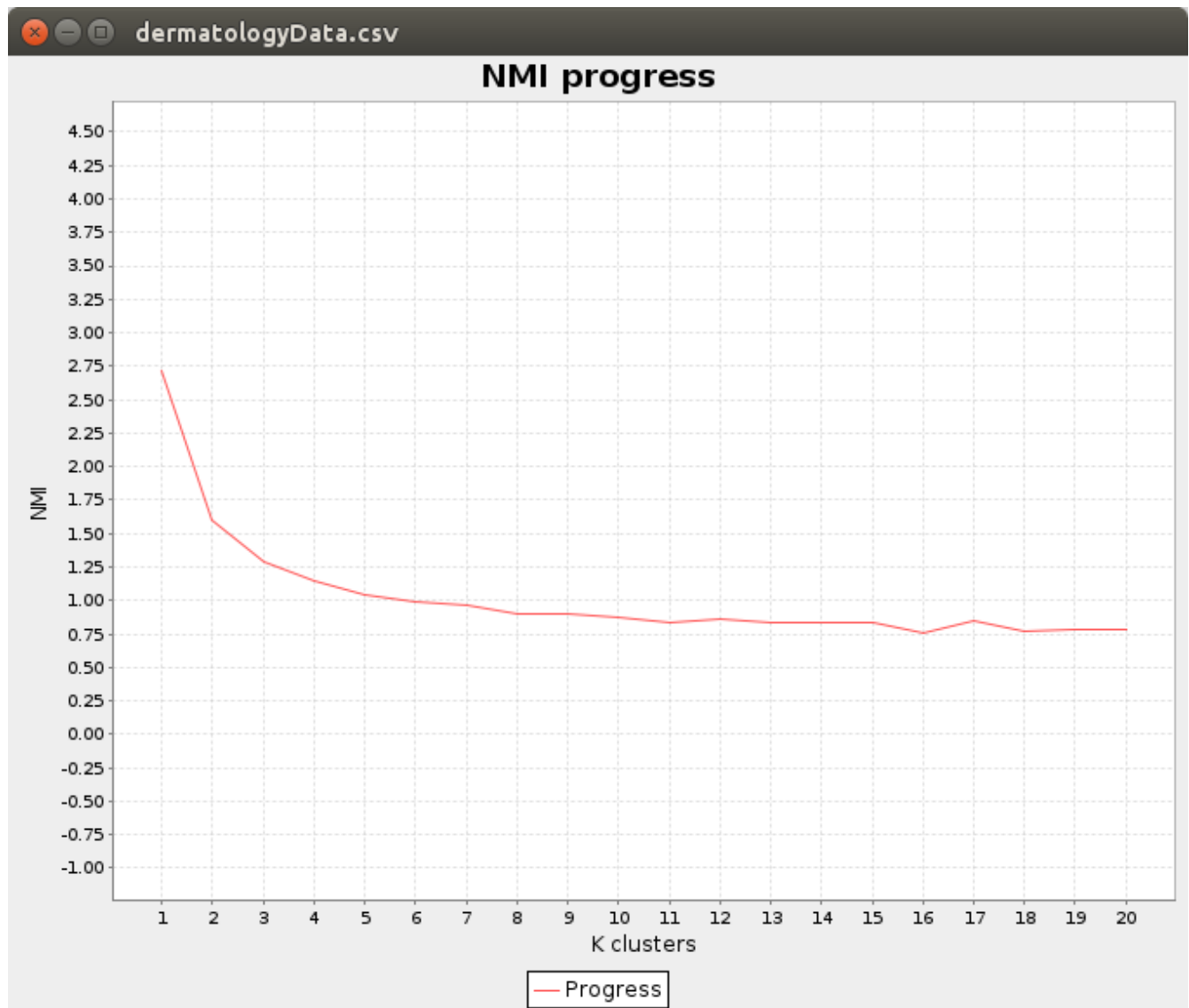
Shreyas Mahimkar

3) Kmeans Clustering::

SSE vs k and NMI vs k plots for each dataset ::

1) Dermatology Data



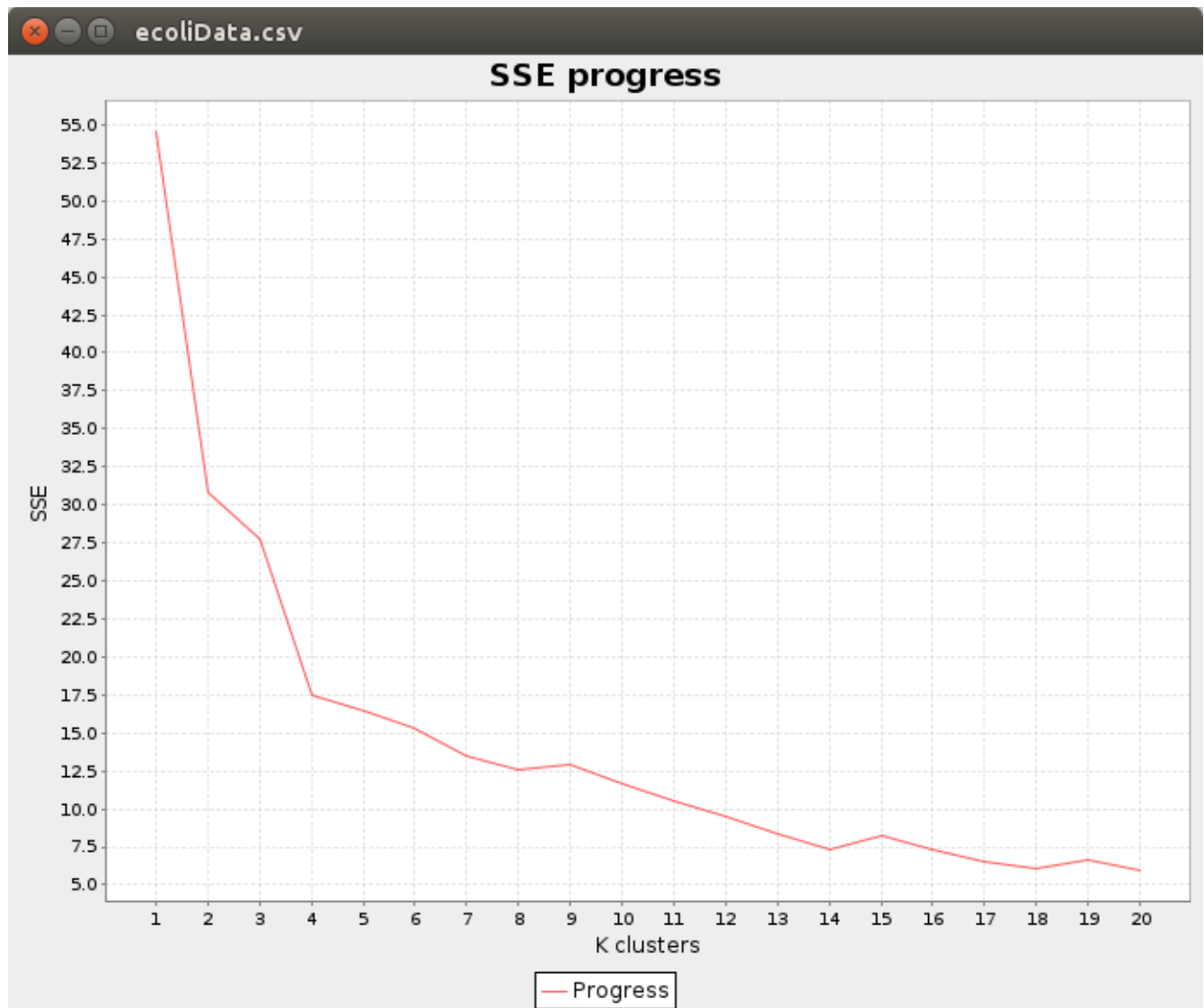


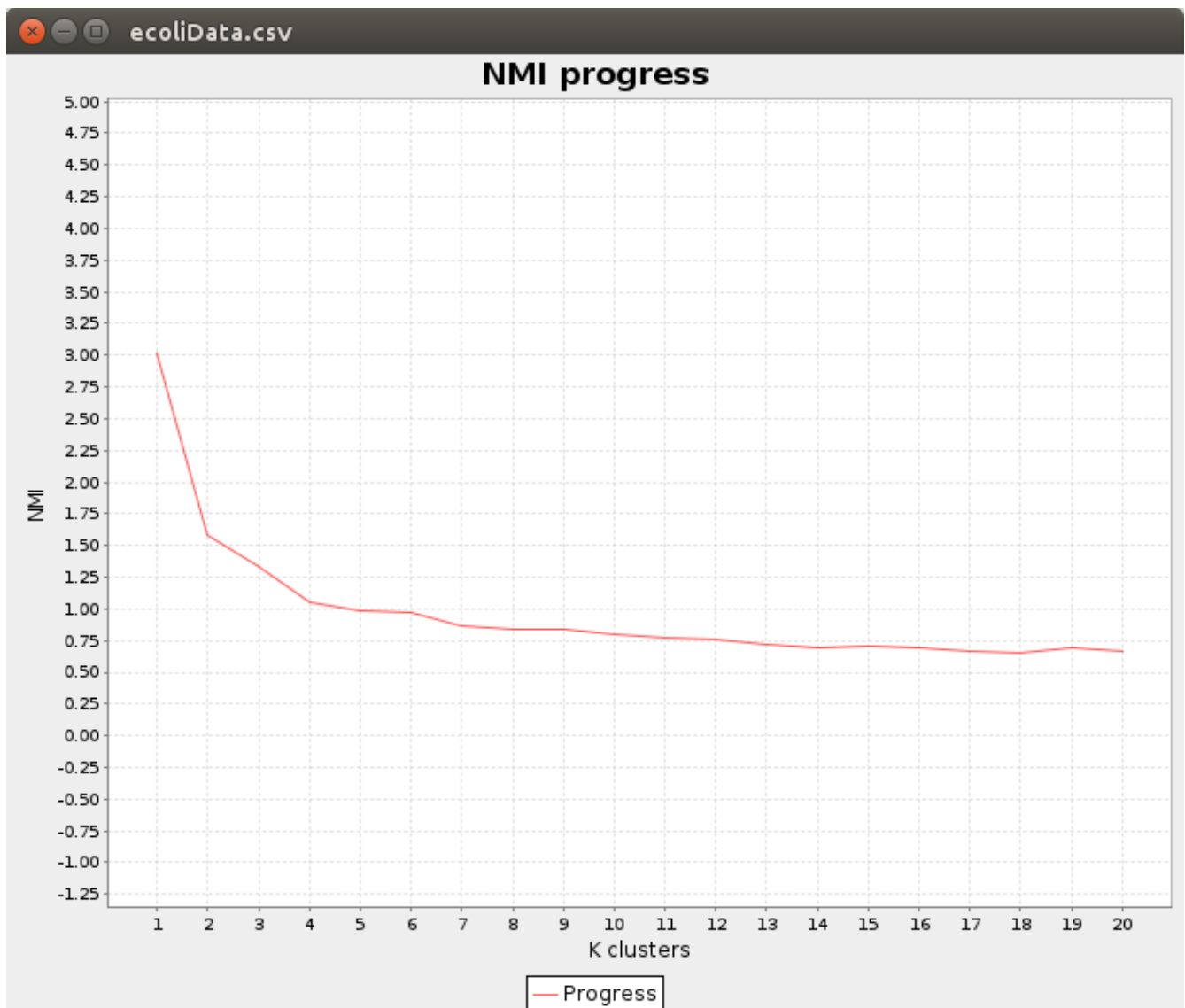
For $k = 6$

Optimal k base on the above plot is 5 (Both NMI and SSE).

NMI:0.9896762049031579 sse:11449.901306748103 iteration:38

2) Ecoli Data



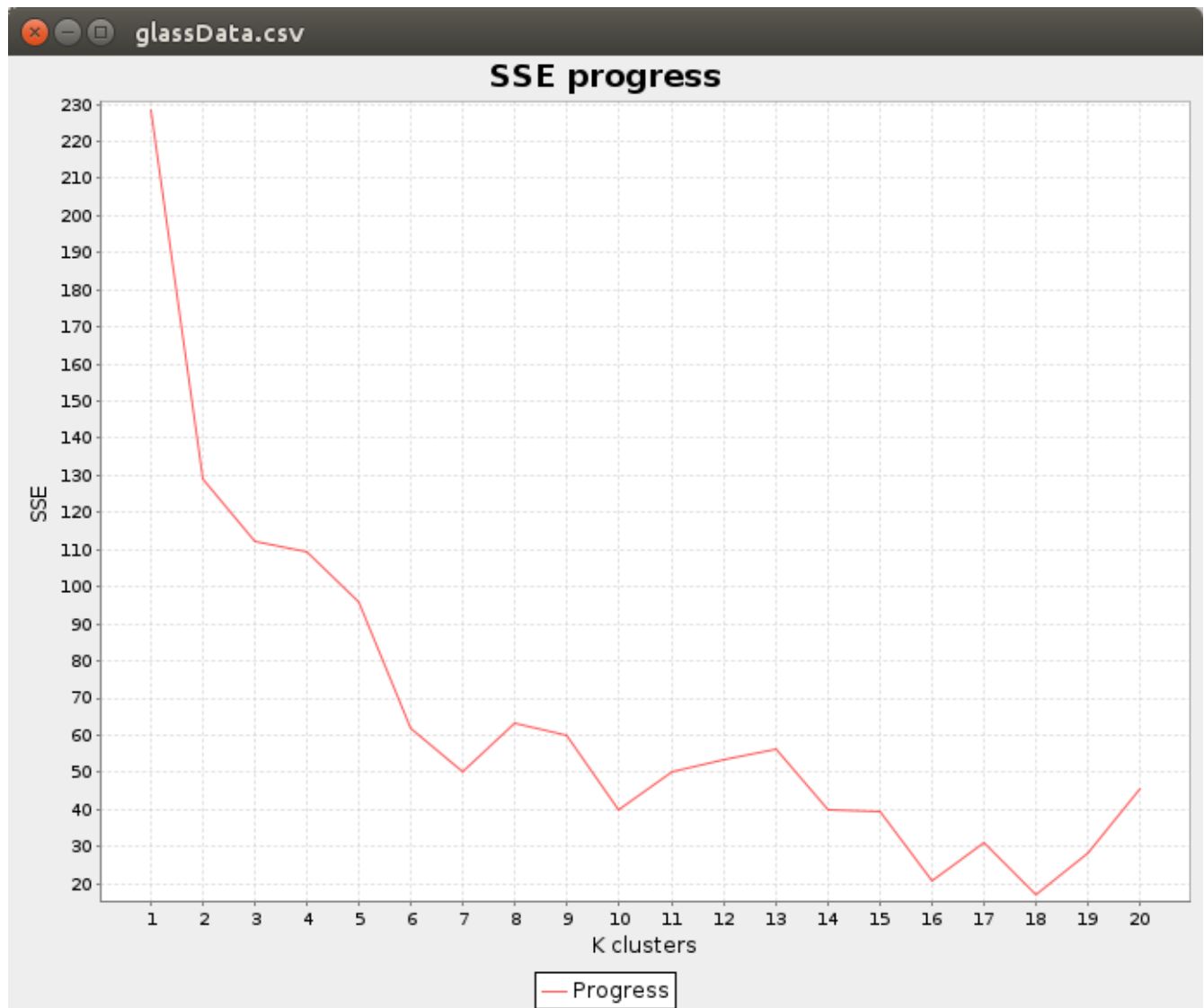


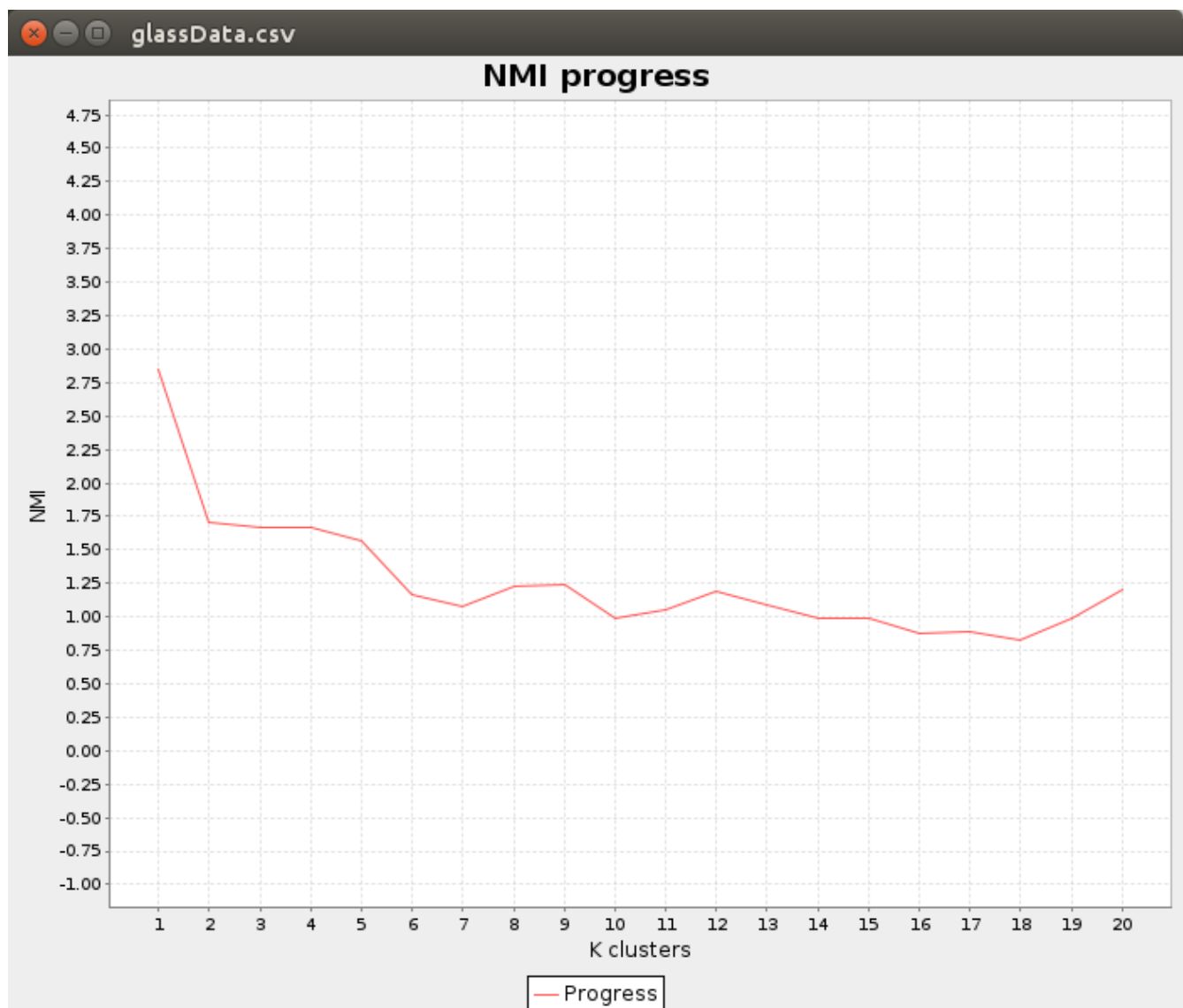
Optimal K base on SSE and NMI :: between 4 and 8. As we can see after k = 4 the exponential drop (knee) is not exponential anymore.

K = 5 – Class number

NMI:0.9696986830920746 sse:15.3123443054913 iteration:17

3) GlassData





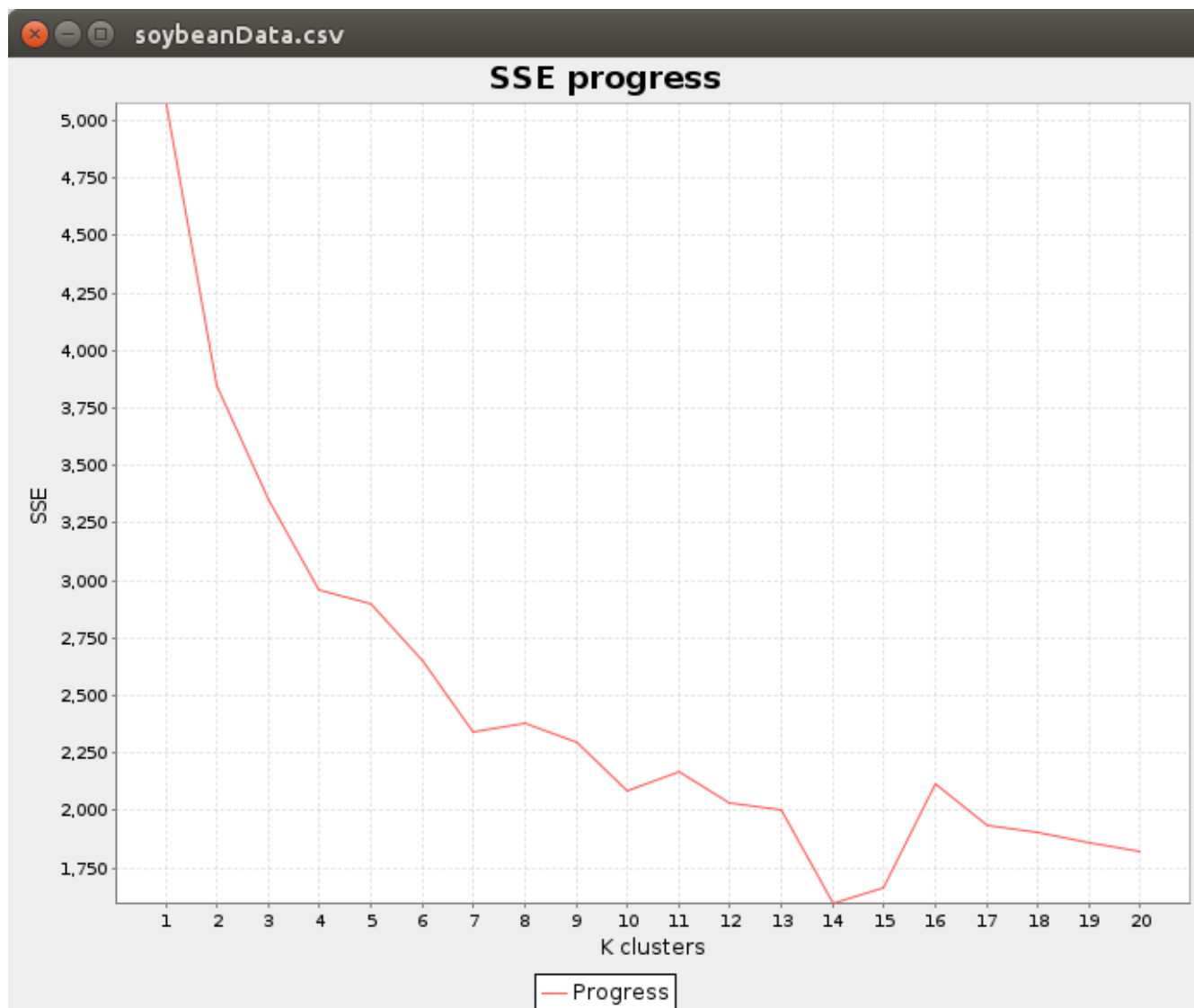
Based on SSE : Best $k = 7$. Exponential drop stops at 7.

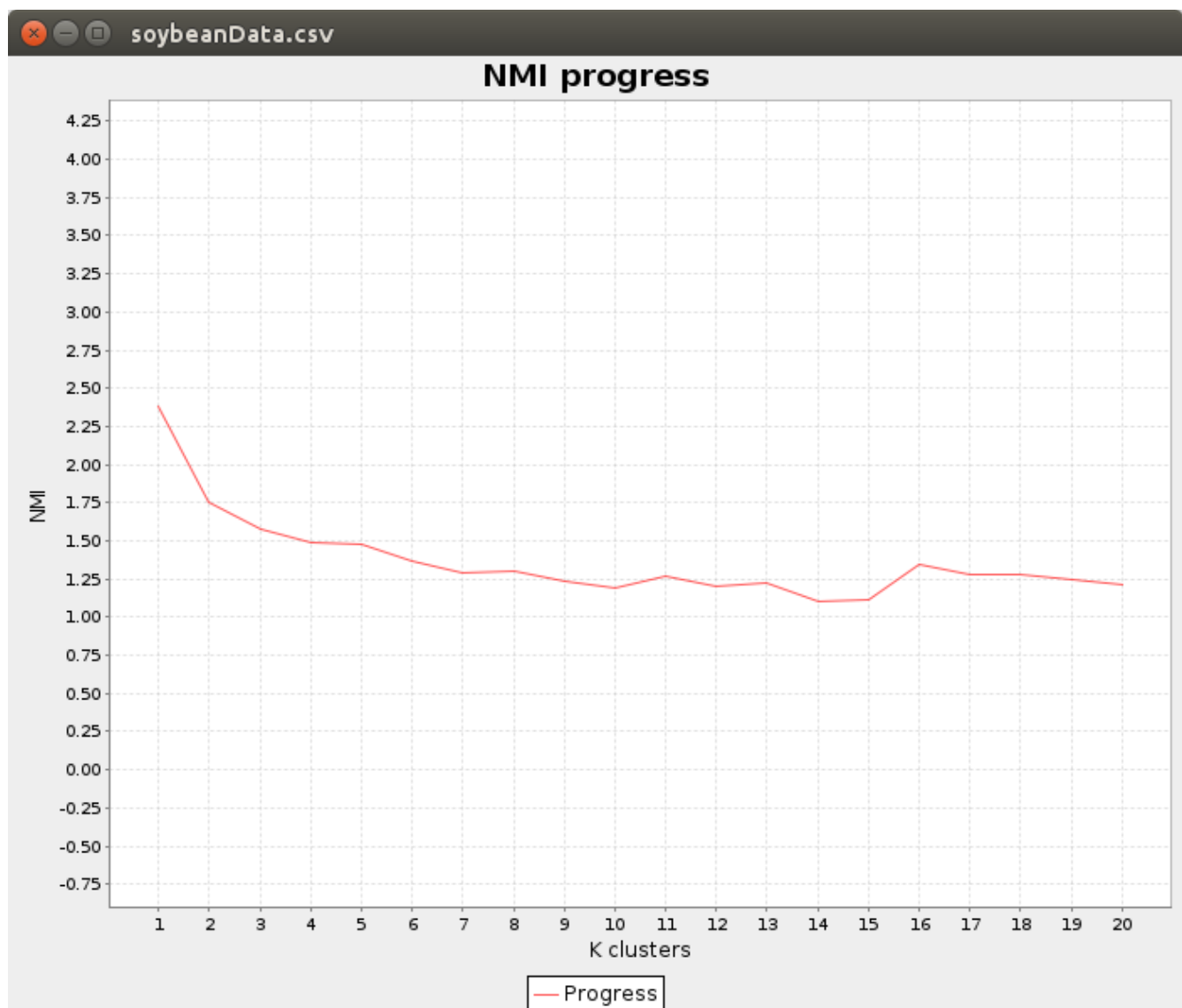
Based on NMI : $k=2$.

$k = 6$

NMI:1.1622090104655665 sse:61.81117417861708 iteration:16

4) SoyBean Data



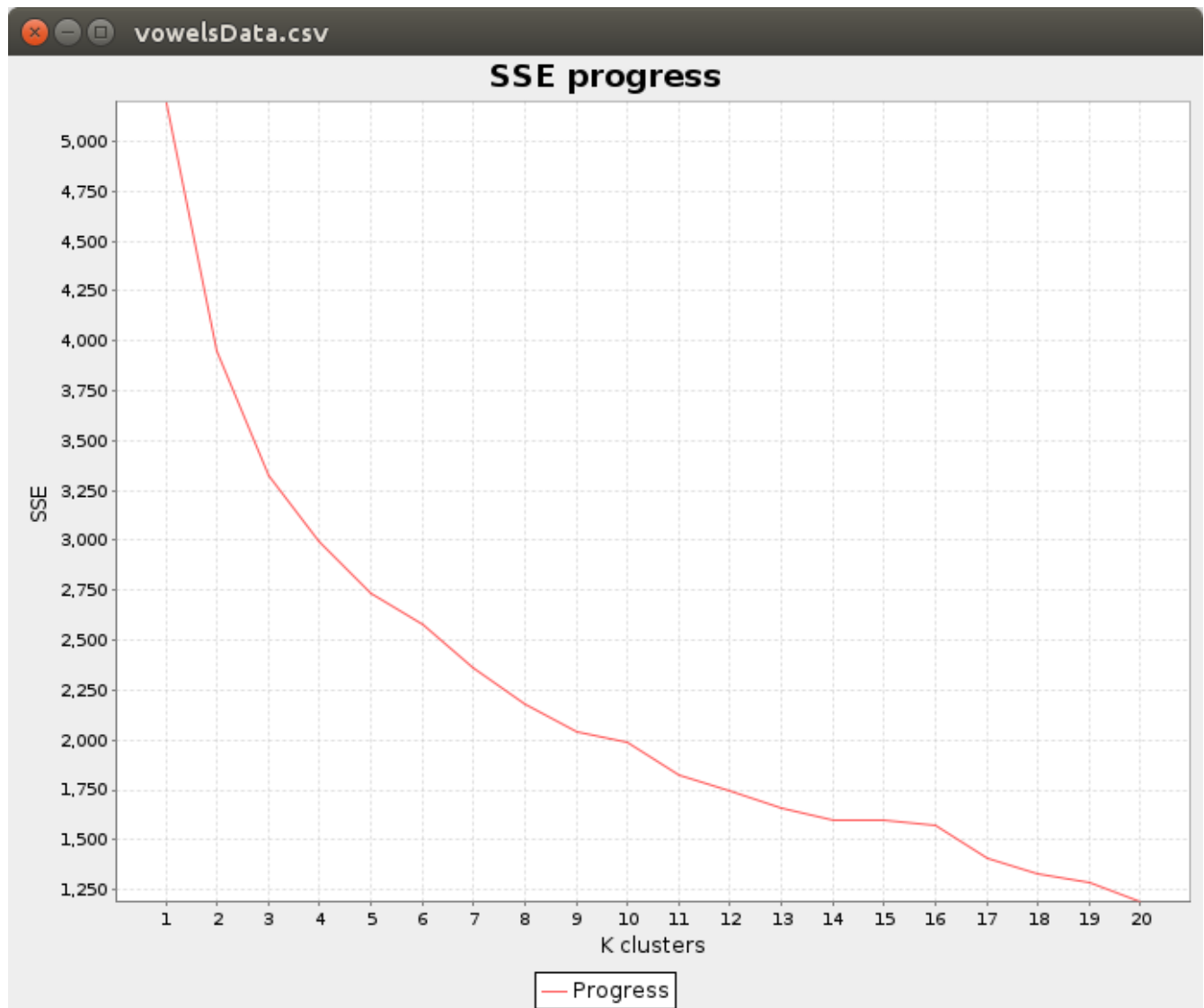


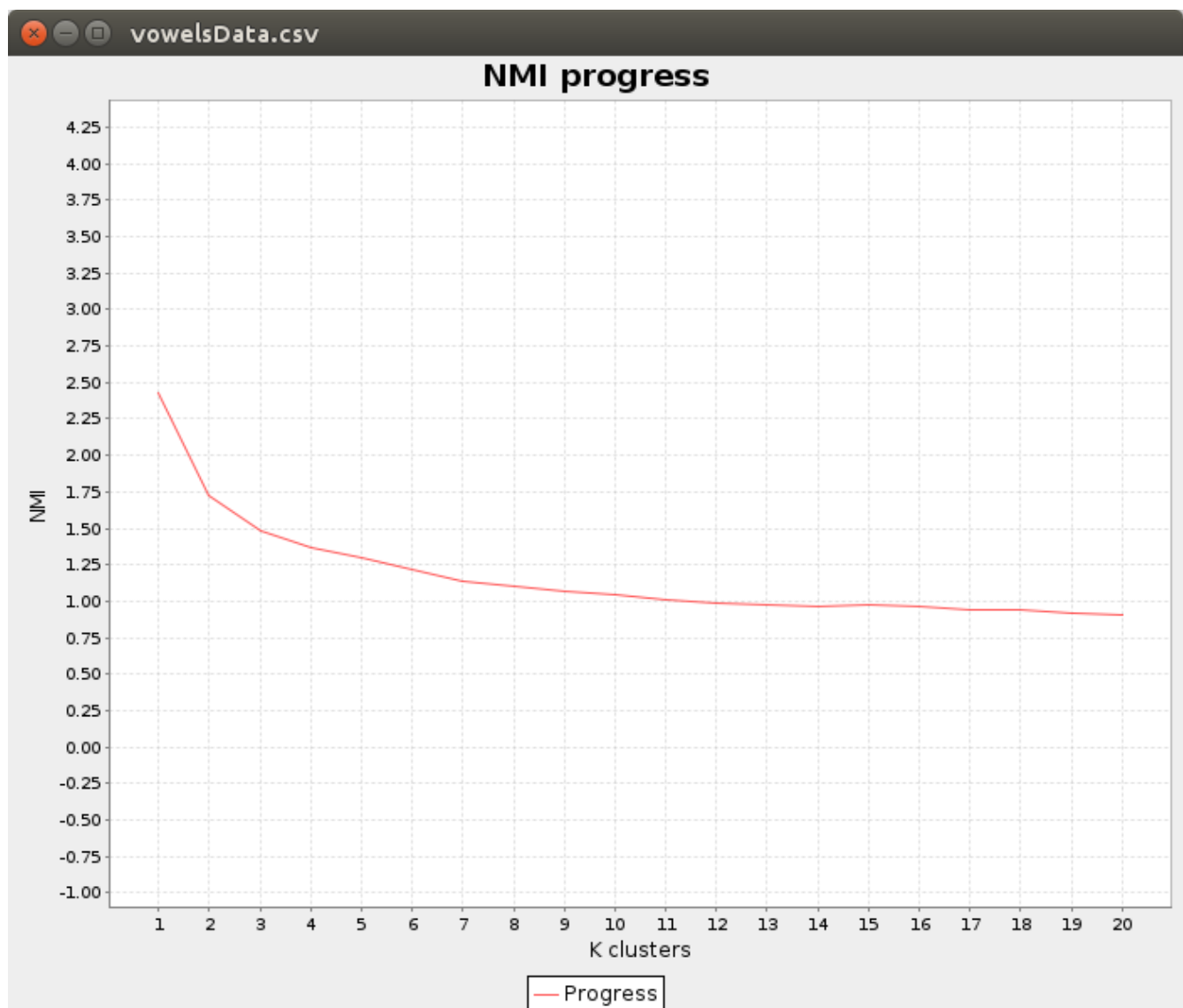
k = 14 according to SSE:

k = 15

NMI:1.1184126663796718 sse:1665.668453373021 iteration:24

5) Vowels Data

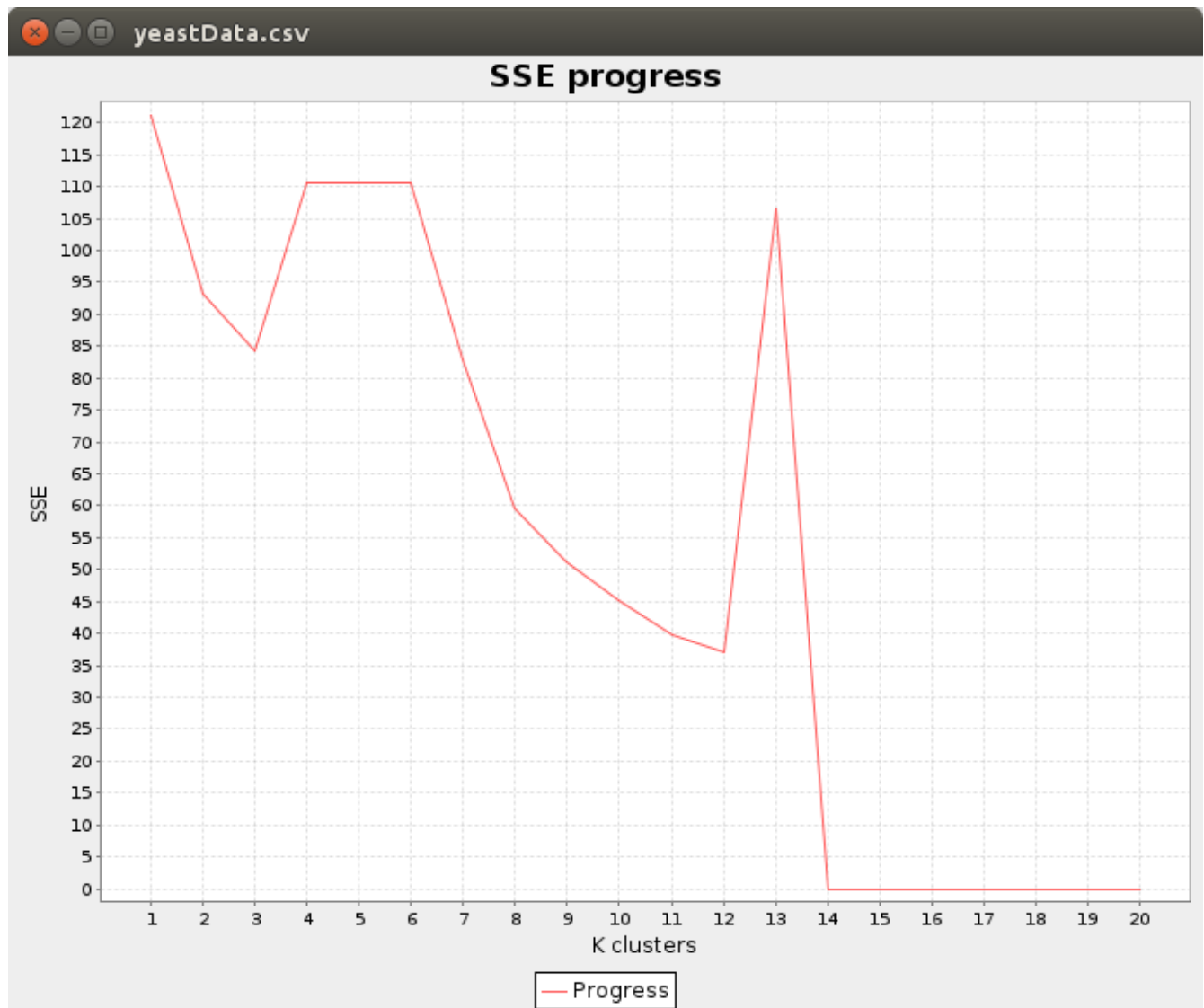


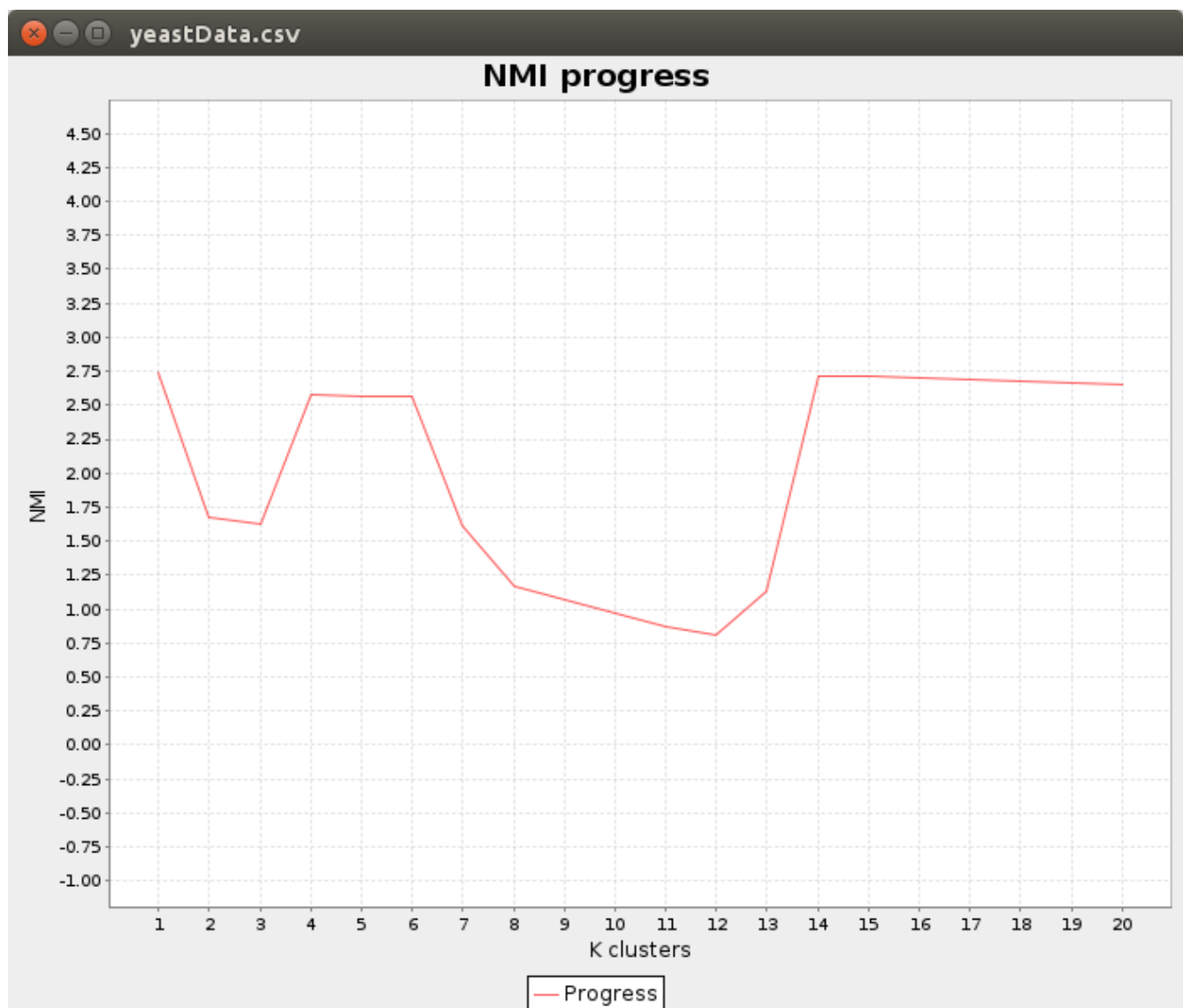


k = 11

NMI:1.0118774989164114 sse:1825.882806346687 iteration:64

6) Yeast Data





k = 9

NMI:1.0730482121277236 sse:51.03846248432911 iteration:46

Based on only Kmeans Algorithm

Datasets	Optimal k based on SSE	Optimal k based on NMI	Num classes	NMI value for k = numClasses	SSE value for k= numClasses
Dermatology	4	4	5	0.96	15.31
Ecoli	5	5	6	0.98	11449.90
Glass	7	2	6	1.16	61.81
SoyBean	14	5	15	1.11	1665.66
Vowels	10	4	11	1.01	1825.88
Yeast	3	2	9	1.07	51.03

Extra Credit for initializing cluster centroids.

Initiate cluster centroids such that if $k = 2$, select the two centroids in the dataset which are farthest from each other.

<http://cs.nyu.edu/courses/spring11/G22.2580-001/lec5.html>

Furthest Point First (FPF) Algorithm

[Geraci et al., 2006]

```
{ V := random point in S;
CENTERS := { V };
for (U in S) {
    LEADER[U] := V;
    D[U] = d(U,V);
}
for (I = 1 to K) {
    X := the point in S-CENTERS with maximal value of D[X];
    C := emptyset;
    for (U in S-CENTERS)
        if (d(U,X) < D[U]) add X to C;
    X := medoid(C,X);
    add X to CENTERS;
    for (U in S-CENTERS) {
        if (d(U,X) < D[U]) {
            LEADER[U] := X;
            D[U] := d(U,X)
        }
    }
}
return(CENTERS)}
```