

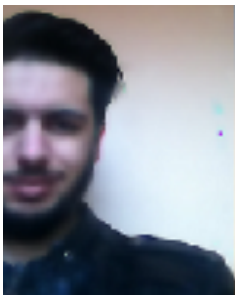
(AP)Affinity Propagation algorithm

Overview:

In statistics and data mining, **affinity propagation (AP)** is a clustering algorithm based on the concept of “message passing” between data points. Unlike clustering algorithms such as **k-means or k-medoids**, affinity propagation does **not require the number of clusters to be determined** or estimated before running the algorithm. Similar to k-medoids, affinity propagation finds “**exemplars**,” members of the input set that are **representative of clusters**

Algo:

- 1-compute similarities
- 2-compute responsibilities
- 3-compute availabilities



delphi & c++ programming

For more information:

contact us at: fb ..gitH ..or .. yt



<https://www.facebook.com/M.Aek.Progs.Angedevil.AD/>



<https://www.youtube.com/channel/UC6AhJIORlsp56XDwqfNSTsg>



<https://github.com/Angedevil-AD>

x: set of data points

N: size of dataset .. where ... $0 \leq i < N$ & $0 \leq k < N$

availability , similarity, responsibility matrix = $N \times N$

1- Similarity Equation:

$$s(i,k) = -\|x_i - x_k\|^2$$

Diagonal $s(i,i)$,initialized to the median similarity

2-Responsibility Equation:

responsibilities matrix initialized to 0

$$r(i,k) \leftarrow s(i,k) - \max_{k \neq k'} \{ a(i,k') + s(i,k') \}$$

3-Availabilities:

$$a(i,k) \leftarrow \min \left(0, r(k,k) + \sum_{i' \notin (i,k)} \max(0, r(i',k)) \right) \text{ for } i \neq k$$

$a(k,k)$ diagonal:

$$a(k,k) \leftarrow \sum_{i' \neq k} \max(0, r(i',k))$$

finally ,

algorithm terminate if clusters unchanged after infinite number of iteration (use break to finaliz the iteration loop) , or after predetermined number of iteration is reached

exemplar extracted from diagonal sum of r & a , where $r(i,i) + a(i,i) > 0$

lets explain the 3 equation with example:

dataset =

one dimension

0.90 ,0.15 ,0.62 ,0.8 ,0.27 ,0.18 ,0.30

len ...N= 7

similarities matrix s(i,k)

compute similarities:

as we know

$$s(i,k) = - || x_i - x_k ||^2$$

$$0 \leq i < N$$

$$0 \leq k < N$$

0.00 ,	-0.56 ,	-0.08 ,	-0.01 ,	-0.40 ,	-0.52 ,	-0.36 ,
-0.56 ,	0.00 ,	-0.22 ,	-0.42 ,	-0.01 ,	-0.00 ,	-0.02 ,
-0.08 ,	-0.22 ,	0.00 ,	-0.03 ,	-0.12 ,	-0.19 ,	-0.10 ,
-0.01 ,	-0.42 ,	-0.03 ,	0.00 ,	-0.28 ,	-0.38 ,	-0.25 ,
-0.40 ,	-0.01 ,	-0.12 ,	-0.28 ,	0.00 ,	-0.01 ,	-0.00 ,
-0.52 ,	-0.00 ,	-0.19 ,	-0.38 ,	-0.01 ,	0.00 ,	-0.01 ,
-0.36 ,	-0.02 ,	-0.10 ,	-0.25 ,	-0.00 ,	-0.01 ,	0.00

$$s(0,0) = -(0.90-0.90)^2 = 0.0$$

$$s(0,1) = -(0.90-0.15)^2 = -0.56$$

$$s(0,2) = -(0.90-0.62)^2 = -0.08$$

.

$$s(6,6) = -(0.30-0.30)^2 = 0.0$$

put result in Matrix NxN

fill Diagonal(s(i,i) with median value of similarity matrix:

0.00 , -0.56 , -0.08 , -0.01 , -0.40 , -0.52 , -0.36 ,
-0.56 , 0.00 , -0.22 , -0.42 , -0.01 , -0.00 , -0.02 ,
-0.08 , -0.22 , 0.00 , -0.03 , -0.12 , -0.19 , -0.10 ,
-0.01 , -0.42 , -0.03 , 0.00 , -0.28 , -0.38 , -0.25 ,
-0.40 , -0.01 , -0.12 , -0.28 , 0.00 , -0.01 , -0.00 ,
-0.52 , -0.00 , -0.19 , -0.38 , -0.01 , 0.00 , -0.01 ,
-0.36 , -0.02 , -0.10 , -0.25 , -0.00 , -0.01 , 0.00

before calc median value we must sort the matrix :

0,00 0,00 0,00 0,00 0,00 0,00 0,00

0,00 0,00 0,00 0,00 -0,01 -0,01 -0,01

-0,01 -0,01 -0,01 -0,01 -0,01 -0,02 -0,02

-0,03 -0,03 -0,08 -0,08 -0,10 -0,10 -0,12

-0,12 -0,19 -0,19 -0,22 -0,22 -0,25 -0,25

-0,28 -0,28 -0,36 -0,36 -0,38 -0,38 -0,40

-0,40 -0,42 -0,42 -0,52 -0,52 -0,56 -0,56

median = sorted_similarities(N/2,N/2)
= -0.08

final similarities matrix :

-0.08 ,	-0.56 ,	-0.08 ,	-0.01 ,	-0.40 ,	-0.52 ,	-0.36 ,
-0.56 ,	-0.08 ,	-0.22 ,	-0.42 ,	-0.01 ,	-0.00 ,	-0.02 ,
-0.08 ,	-0.22 ,	-0.08 ,	-0.03 ,	-0.12 ,	-0.19 ,	-0.10 ,
-0.01 ,	-0.42 ,	-0.03 ,	-0.08 ,	-0.28 ,	-0.38 ,	-0.25 ,
-0.40 ,	-0.01 ,	-0.12 ,	-0.28 ,	-0.08 ,	-0.01 ,	-0.00 ,
-0.52 ,	-0.00 ,	-0.19 ,	-0.38 ,	-0.01 ,	-0.08 ,	-0.01 ,
-0.36 ,	-0.02 ,	-0.10 ,	-0.25 ,	-0.00 ,	-0.01 ,	-0.08

- we done with similarities matrix.
lets move to responsibilities and availabilities

these two matrix , must updated continuouly as long as the algorithm not terminated, therfore we have to determine the number of iteration , or use an infinite loop and break if clusters, unchanged

for i=0 to infinite
responsibilities(...)
availabilities(...)
if curent clusters = previous clusters ..break

-responsibility:

$r(i,k) = s(i,k) - \max(s(i,k') + a(i,k'))$ wherre k not equal k'

compute max value in $s(i,k') + a(i,k')$

$0 \leq i < N, 0 \leq k < N, 0 \leq k' < N$

for i = 0 , k = 0, k' = 0

k = k' ..we know that k must not equal to k' ..so this step will bypassed

move to

for i = 0 , k = 0, k' = 1

k <> k' ... fine

$s(i,k') = -0.56$...v(alue from similarities matrix)

$a(i,k') = 0$... availability is not yet filled ... is initialized to 0 for first use

$s+k = -0.56$

for i = 0 , k = 0, k' = 2

k <> k' ... fine

$s(i,k') = -0.08$

$a(i,k') = 0$

$s+k = -0.08$

for i = 0 , k = 0, k' = 3

k <> k' ... fine

$s(i,k') = -0.01$

$a(i,k') = 0$

$s+k = -0.01$

```
for i = 0 , k = 0, k' = 4
k<>k' ... fine
s(i,k') = -0.40
a(i,k') = 0
s+k=-0.40
```

```
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for i = 0 , k = 0, k' = 5
k<>k' ... fine
s(i,k') = -0.52
a(i,k') = 0
s+k=-0.52
```

```
-----
for i = 0 , k = 0, k' = 6
k<>k' ... fine
s(i,k') = -0.36
a(i,k') = 0
s+k=-0.36
-----
```

max value in : -0.56 , -0.08 , -0.01 , -0.40 , -0.52 , -0.36 is -0.01
maxval = -0.01

$r(i,k) = s(i,k) - \text{maxval} = -0.08 - (-0.01) = -0.07$

continue:

```
for i = 0 , k = 1, k' = 0
k<>k' ... fine
s(i,k') = -0.08
a(i,k') = 0
s+k=-0.08
```

```
-----
for i = 0 , k = 1, k' = 1
k=k' ... bypass
-----
```

```
for i = 0 , k = 1, k' = 2
k<>k' ... fine
s(i,k') = -0.08
a(i,k') = 0
s+k=-0.08
```

```

-----
for i = 0 , k = 1, k' = 3 k<>k' ... fine
s(i,k') = -0.01
a(i,k') = 0
s+k=-0.01

```

```

.
.continue until

```

```

-----
for i = 0 , k = 1, k' = 6 k<>k' ... fine
s(i,k') = -0.36
a(i,k') = 0
s+k=-0.36

```

max value in : -0.08, -0.08 , -0.01 , -0.40 , -0.52 , -0.36 is -0.01 maxval = -0.01

$r(i,k) = s(i,k) - \text{maxval} = -0.56 - (-0.01) = -0.55$

```

continue until
for i = 6 , k = 6, k' = 6

```

put all result in matrix, and before do that , we need apply dump factor on result, where dump factor [0...1]

ex: dump factor = 0.3
new matrix = (1-dump factor) * current matrix + dump factor * previous matrix for
matrix[0][0] = (1-0.3)*0.07 - 0.3*0 = -0.05

responsibilities matrix:

[-0.05 , -0.39 , -0.05 , 0.05 , -0.27 , -0.36 , -0.25 , -0.39 , -0.05 , -0.15 , -0.30 , -0.01 , 0.01 , -0.02 , -0.03 , -0.13 , -0.03 , 0.03 , -0.06 , -0.11 , -0.05 , 0.02 , -0.29 , -0.02 , -0.05 , -0.19 , -0.26 , -0.17 , -0.28 , -0.01 , -0.09 , -0.20 , -0.05 , -0.01 , 0.01 , -0.36 , 0.01 , -0.13 , -0.27 , -0.01 , -0.05 , -0.01 , -0.25 , -0.02 , -0.07 , -0.17 , 0.01 , -0.01 , -0.05]
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now compute availabilities:

-availabilities:

$a(i,k) = \min(0, r(k,k) + \sum (\max(0, r(i',k))) \text{ } i \text{ not equal to } k \text{ and } i' \text{ not belong to } (i,k)$

compute max value in $0, r(i',k)$

$0 \leq i < N, \quad 0 \leq k < N, \quad 0 \leq i' < N$

for $i = 0, \quad k = 0, \quad k' = 0$

$i = i'$ bypass

for $i = 0, \quad k = 0, \quad i' = 1$

$i < i' \dots$ fine

$r(i',k) = -0.39 \dots$ (value from responsibilities matrix)

$\max(0, -0.39) = 0$

for $i = 0, \quad k = 0, \quad i' = 2 \quad i < i'$

\dots fine

$r(i',k) = -0.03$

$\max(0, -0.03) = 0$

for $i = 0, \quad k = 0, \quad i' = 3 \quad i < i'$

\dots fine

$r(i',k) = 0.02$

$\max(0, 0.02) = 0.02$

for $i = 0, \quad k = 0, \quad i' = 4$

$i < i' \dots$ fine

$r(i',k) = -0.28$

$\max(0, -0.28) = 0$

for $i = 0, \quad k = 0, \quad i' = 5$

$i < i' \dots$ fine

$r(i',k) = -0.36$

$\max(0, -0.36) = 0$

for $i = 0, \quad k = 0, \quad i' = 6$

$i < i' \dots$ fine

$r(i',k) = -0.25$

$\max(0, -0.25) = 0$

sum all result : $0 + 0 + 0.02 + 0 + 0 + 0 = 0.02$

add sum to $r(k,k) \dots r(k,k) = r(0,0) = -0.05$

$-0.05 + 0.02 = -0.03$

$a(i,k) = \min(0, -0.03) = -0.03 < \text{-----}$

now set diagonal $a(k,k)$ $a(k,k) = \sum(\max(0, r(i',k)))$ $\sum(\max(0, r(i',k))) = 0.02$
 $a(k,k) = 0.02$

don't forget to apply dump factor to availabilities matrix
 continue until
 for $i = 6$, $k = 6$, $i' = 6$

you get as result :

availabilities matrix

[0.01 , -0.03 , -0.02 , -0.01 , -0.03 , -0.03 , -0.03 , -0.02 , 0.00 , -0.02 , 0.00 , -0.03 , -0.04 , -0.03 , -0.02 , -0.03 , 0.00 , 0.00 , -0.03 , -0.03 , -0.03 , -0.03 , -0.03 , -0.02 , 0.06 , -0.03 , -0.03 , -0.03 , -0.02 , -0.03 , -0.02 , 0.00 , 0.01 , -0.03 , -0.04 , -0.02 , -0.04 , -0.02 , 0.00 , -0.03 , 0.01 , -0.03 , -0.02 , -0.03 , -0.02 , 0.00 , -0.04 , -0.03 , 0.00]
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we done with the first iteration

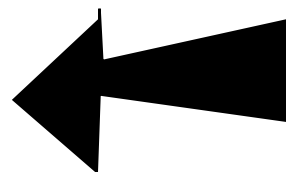
calc availabilities + responsibilities & extract exemplar from diagonal. avail-
 abilities + responsibilities matrix:

-0.04 , -0.42 , -0.07 , 0.04 , -0.30 , -0.39 , -0.28 ,
 -0.42 , -0.05 , -0.18 , -0.30 , -0.04 , -0.03 , -0.05 ,
 -0.05 , -0.17 , -0.03 , 0.03 , -0.09 , -0.14 , -0.08 ,
 -0.02 , -0.32 , -0.04 , 0.01 , -0.22 , -0.29 , -0.20 ,
 -0.30 , -0.04 , -0.11 , -0.20 , -0.05 , -0.04 , -0.03 ,
 -0.38 , -0.03 , -0.16 , -0.27 , -0.04 , -0.05 , -0.04 ,
 -0.27 , -0.05 , -0.09 , -0.17 , -0.03 , -0.04 , -0.05 ,

check if any element of the diagonal = positive ?

 -0.04 , -0.42 , -0.07 , 0.04 , -0.30 , -0.39 , -0.28 , -0.42 , -0.05 , -0.18 , -0.30 , -0.04 , -0.03 , -0.05 , -0.05 , -0.17 , -0.03 , 0.03 , -0.09 , -0.14 , -0.08 , -0.02 , -0.32 , -0.04 , 0.01 , -0.22 , -0.29 , -0.20 , -0.30 , -0.04 , -0.11 , -0.20 , -0.05 , -0.04 , -0.03 , -0.38 , -0.03 , -0.16 , -0.27 , -0.04 , -0.05 , -0.04 , -0.27 , -0.05 , -0.09 , -0.17 , -0.03 , -0.04 , -0.05 ,

0.01 = positive

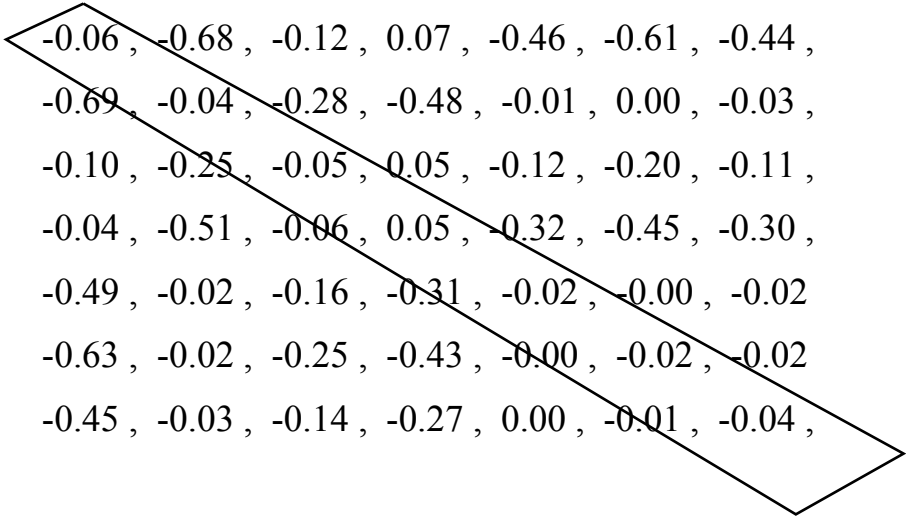


$(3,3) = 0.01$

exemplar = 0.01

keep updating availabilities & responsibilities by passing to iteration 2,3,4....until n

check result in iteration 2:



-0.06 , -0.68 , -0.12 , 0.07 , -0.46 , -0.61 , -0.44 ,
-0.69 , -0.04 , -0.28 , -0.48 , -0.01 , 0.00 , -0.03 ,
-0.10 , -0.25 , -0.05 , 0.05 , -0.12 , -0.20 , -0.11 ,
-0.04 , -0.51 , -0.06 , 0.05 , -0.32 , -0.45 , -0.30 ,
-0.49 , -0.02 , -0.16 , -0.31 , -0.02 , -0.00 , -0.02
-0.63 , -0.02 , -0.25 , -0.43 , -0.00 , -0.02 , -0.02
-0.45 , -0.03 , -0.14 , -0.27 , 0.00 , -0.01 , -0.04 ,

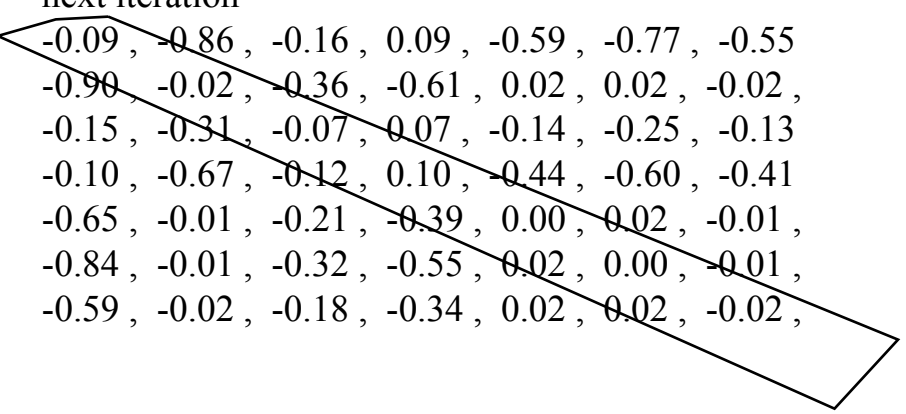
exemplar = 0.05

currnt exemplars not = previous exemplar

0.05 not = 0.01

continue iteration to check if exemplar change nor not ... if not change ... break the iteration loop

next iteration



-0.09 , -0.86 , -0.16 , 0.09 , -0.59 , -0.77 , -0.55
-0.90 , -0.02 , -0.36 , -0.61 , 0.02 , 0.02 , -0.02 ,
-0.15 , -0.31 , -0.07 , 0.07 , -0.14 , -0.25 , -0.13
-0.10 , -0.67 , -0.12 , 0.10 , -0.44 , -0.60 , -0.41
-0.65 , -0.01 , -0.21 , -0.39 , 0.00 , 0.02 , -0.01 ,
-0.84 , -0.01 , -0.32 , -0.55 , 0.02 , 0.00 , -0.01 ,
-0.59 , -0.02 , -0.18 , -0.34 , 0.02 , 0.02 , -0.02 ,

exemplar = 0.10,0.00,0.00

currnt exemplars not = previous exemplar

0.10,0.00,0.00 not = 0.05

continue iteration :
after 48 iteration we get

-0.16 , -1.29 , -0.27 , 0.16 , -0.94 , -1.22 , -0.82 ,
-1.38 , 0.00 , -0.53 , -0.89 , -0.03 , -0.03 , -0.00 ,
-0.27 , -0.44 , -0.11 , 0.11 , -0.24 , -0.41 , -0.16 ,
-0.62 , -1.43 , -0.62 , 0.41 , -1.13 , -1.37 , -1.02 ,
-1.25 , -0.19 , -0.55 , -0.82 , -0.03 , -0.21 , -0.16 ,
-1.53 , -0.16 , -0.72 , -1.06 , -0.21 , -0.03 , -0.19 ,
-0.91 , 0.00 , -0.25 , -0.49 , -0.03 , -0.03 , -0.00 ,

exemplars : 0.00, 0.41

next iteration

-0.16 , -1.29 , -0.27 , 0.16 , -0.94 , -1.22 , -0.82 ,
-1.38 , 0.00 , -0.53 , -0.89 , -0.03 , -0.03 , -0.00 ,
-0.27 , -0.44 , -0.11 , 0.11 , -0.24 , -0.41 , -0.16 ,
-0.62 , -1.43 , -0.62 , 0.41 , -1.13 , -1.37 , -1.02 ,
-1.25 , -0.19 , -0.55 , -0.82 , -0.03 , -0.21 , -0.16 ,
-1.53 , -0.16 , -0.72 , -1.06 , -0.21 , -0.03 , -0.19 ,
-0.91 , 0.00 , -0.25 , -0.49 , -0.03 , -0.03 , -0.00 ,

exemplars: 0.00, 0.41
as you see after 49 iteration
exemplars wan't change.....
algorithm terminate

exemplar 0.00 position = 1
exemplar 0.41 position = 3

clustering

1- get max(similarities[1] , similarities[3])

for each i

for each k

if similarities[i][1] > similarities[i][3] ... dataset(i) is belong to cluster(1)

if similarities[i][1] < similarities[i][3] ... dataset(i) is belong to cluster(3)

similarities

-0.08 , -0.56 , -0.08 , -0.01 , -0.40 , -0.52 , -0.36 ,
-0.56 , -0.08 , -0.22 , -0.42 , -0.01 , -0.00 , -0.02 ,
-0.08 , -0.22 , -0.08 , -0.03 , -0.12 , -0.19 , -0.10 ,
-0.01 , -0.42 , -0.03 , -0.08 , -0.28 , -0.38 , -0.25 ,
-0.40 , -0.01 , -0.12 , -0.28 , -0.08 , -0.01 , -0.00 ,
-0.52 , -0.00 , -0.19 , -0.38 , -0.01 , -0.08 , -0.01 ,
-0.36 , -0.02 , -0.10 , -0.25 , -0.00 , -0.01 , -0.08

for i=0

-0.56 < -0.01

dataset(0) -> cluster 3

for i=4

-0.01 > -0.28

dataset(4) -> cluster 1

for i=1

-0.08 > -0.42

dataset(1) -> cluster 1

for i=5

-0.00 > -0.38

dataset(5) -> cluster 1

for i=2

-0.22 < -0.03

dataset(2) -> cluster 3

for i=6

-0.02 > -0.25

dataset(6) -> cluster 1

for i=3

-0.41 < -0.08

dataset(3) -> cluster 3

cluster(1) = 0.15, 0.27, 0.18, 0.30

cluster(3) = 0.90, 0.62, 0.8

lets consider another exemple this time we work on 2-D ...points(x,y)

len: 25

(0,0) (0,2) (0,4) (0,6) (1,1)

(1,3) (1,5) (2,0) (2,2) (2,4)

(2,6) (3,1) (3,3) (3,5) (4,0)

(4,2) (4,4) (4,6) (5,1) (5,3)

(5,5) (6,0) (6,2) (6,4) (6,6)

similarities after setting diagonal:

len = 625

-16 -4 -16 -36 -2 -10 -26 -4 -8 -20 -40 -10 -18 -34 -16 -20 -32 -52 -26 -34 -50 -36 -40 -52 -72
-4 -16 -4 -16 -2 -2 -10 -8 -4 -8 -20 -10 -10 -18 -20 -16 -20 -32 -26 -26 -34 -40 -36 -40 -52
-16 -4 -16 -4 -10 -2 -2 -20 -8 -4 -8 -18 -10 -10 -32 -20 -16 -20 -34 -26 -26 -52 -40 -36 -40
-36 -16 -4 -16 -26 -10 -2 -40 -20 -8 -4 -34 -18 -10 -52 -32 -20 -16 -50 -34 -26 -72 -52 -40 -36
-2 -2 -10 -26 -16 -4 -16 -2 -2 -10 -26 -4 -8 -20 -10 -10 -18 -34 -16 -20 -32 -26 -26 -34 -50
-10 -2 -2 -10 -4 -16 -4 -10 -2 -2 -10 -8 -4 -8 -18 -10 -10 -18 -20 -16 -20 -34 -26 -26 -34
-26 -10 -2 -2 -16 -4 -16 -26 -10 -2 -2 -20 -8 -4 -34 -18 -10 -10 -32 -20 -16 -50 -34 -26 -26
-4 -8 -20 -40 -2 -10 -26 -16 -4 -16 -36 -2 -10 -26 -4 -8 -20 -40 -10 -18 -34 -16 -20 -32 -52
-8 -4 -8 -20 -2 -2 -10 -4 -16 -4 -16 -2 -2 -10 -8 -4 -8 -20 -10 -10 -18 -20 -16 -20 -32
-20 -8 -4 -8 -10 -2 -2 -16 -4 -16 -4 -10 -2 -2 -20 -8 -4 -8 -18 -10 -10 -32 -20 -16 -20
-40 -20 -8 -4 -26 -10 -2 -36 -16 -4 -16 -26 -10 -2 -40 -20 -8 -4 -34 -18 -10 -52 -32 -20 -16
-10 -10 -18 -34 -4 -8 -20 -2 -2 -10 -26 -16 -4 -16 -2 -2 -10 -26 -4 -8 -20 -10 -10 -18 -34
-18 -10 -10 -18 -8 -4 -8 -10 -2 -2 -10 -4 -16 -4 -10 -2 -2 -10 -8 -4 -8 -18 -10 -10 -18
-34 -18 -10 -10 -20 -8 -4 -26 -10 -2 -2 -16 -4 -16 -26 -10 -2 -2 -20 -8 -4 -34 -18 -10 -10
-16 -20 -32 -52 -10 -18 -34 -4 -8 -20 -40 -2 -10 -26 -16 -4 -16 -36 -2 -10 -26 -4 -8 -20 -40
-20 -16 -20 -32 -10 -10 -18 -8 -4 -8 -20 -2 -2 -10 -4 -16 -4 -16 -2 -2 -10 -8 -4 -8 -20
-32 -20 -16 -20 -18 -10 -10 -20 -8 -4 -8 -10 -2 -2 -16 -4 -16 -4 -10 -2 -2 -20 -8 -4 -8
-52 -32 -20 -16 -34 -18 -10 -40 -20 -8 -4 -26 -10 -2 -36 -16 -4 -16 -26 -10 -2 -40 -20 -8 -4
-26 -26 -34 -50 -16 -20 -32 -10 -10 -18 -34 -4 -8 -20 -2 -2 -10 -26 -16 -4 -16 -2 -2 -10 -26
-34 -26 -26 -34 -20 -16 -20 -18 -10 -10 -18 -8 -4 -8 -10 -2 -2 -10 -4 -16 -4 -10 -2 -2 -10
-50 -34 -26 -26 -32 -20 -16 -34 -18 -10 -10 -20 -8 -4 -26 -10 -2 -2 -16 -4 -16 -26 -10 -2 -2
-36 -40 -52 -72 -26 -34 -50 -16 -20 -32 -52 -10 -18 -34 -4 -8 -20 -40 -2 -10 -26 -16 -4 -16 -36
-40 -36 -40 -52 -26 -26 -34 -20 -16 -20 -32 -10 -10 -18 -8 -4 -8 -20 -2 -2 -10 -4 -16 -4 -16
-52 -40 -36 -40 -34 -26 -26 -32 -20 -16 -20 -18 -10 -10 -20 -8 -4 -8 -10 -2 -2 -16 -4 -16 -4
-72 -52 -40 -36 -50 -34 -26 -52 -32 -20 -16 -34 -18 -10 -40 -20 -8 -4 -26 -10 -2 -36 -16 -4 -16

As you see... a large matrix has been generated with len=625 where
 $25 * 25 = 625$

“assume that you input 1k set of data... the similarities size = $1k * 1k = 1mb$

1mb set of data give you 1TB.... $1mb * 1mb = 1000000 * 1000000 = 1000000000000$

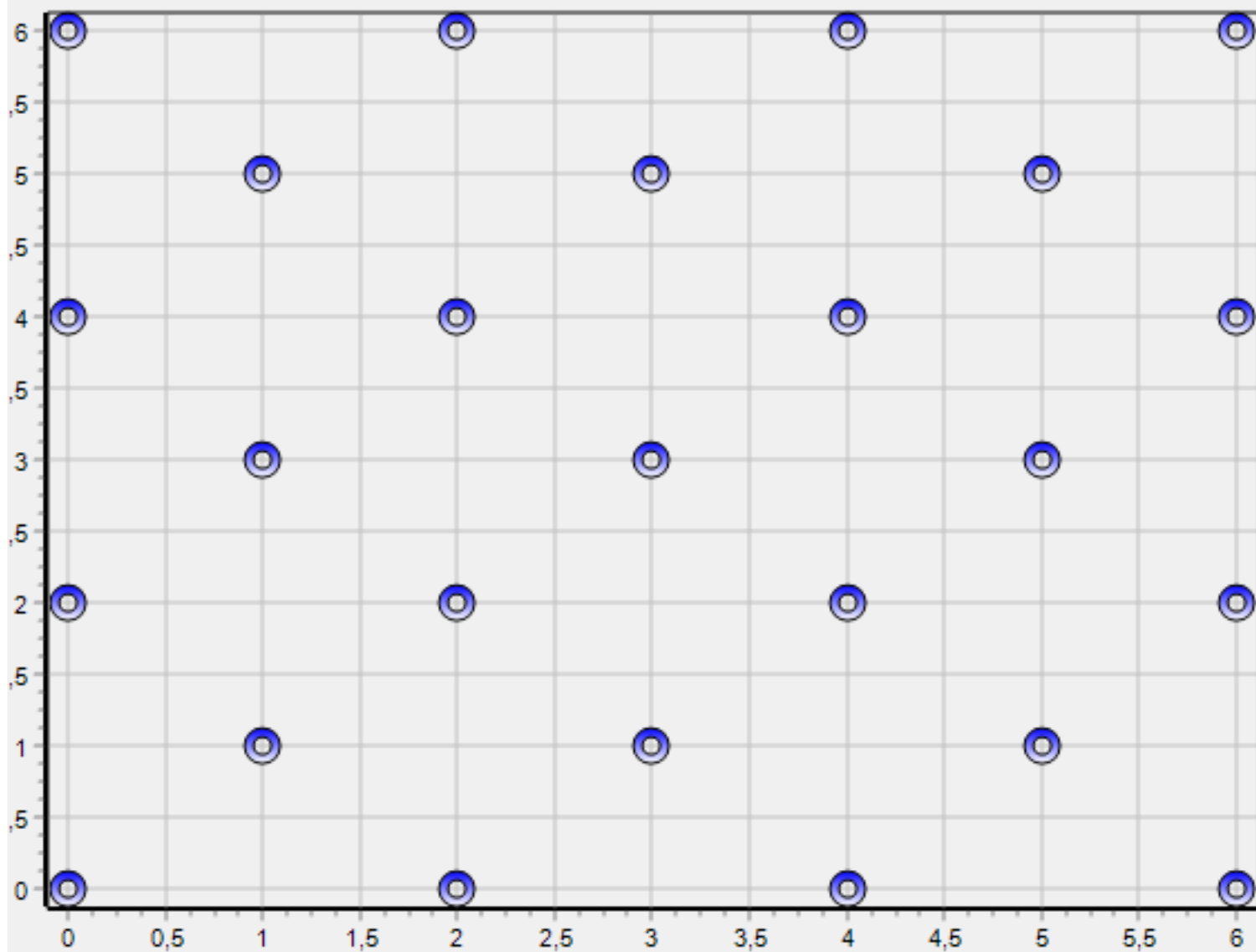
Space complexity is given by: $O(N^2)$

algorithm terminate after 27 iteration .. exemplar table:

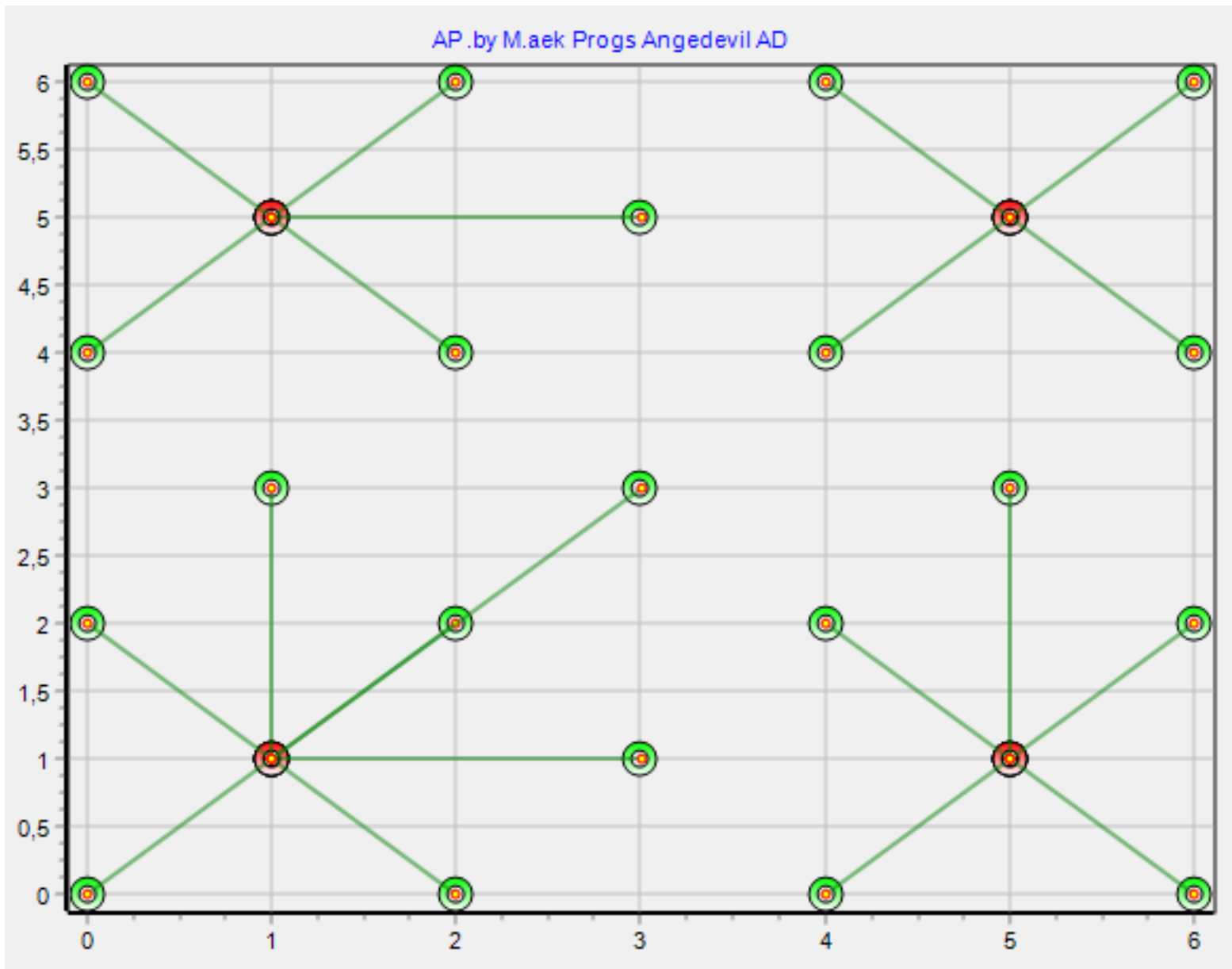
4,4,6,6,4, 4,6,4,4,6,
6,4,4,6,18,
18,20,20,18,18,
20,18,18,20,20

Time Complexity
 $O(N^2i)$

AP .by M.aek Progs Angedevil AD



dataset after affinity propagation



red point i = exemplars

affinity propagation doesn't need to determine the number of clusters.

affinity propagation is better than kmeans & k medoids on error minimizing and outliers

application:

- The inventors of affinity propagation showed it is better for certain computer vision and computational biology tasks

- Another recent application was in economics, when the affinity propagation was used to find some temporal patterns in the output multipliers of the US economy between 1997 and 2017

-and many other advantages

