MATLAB **EXPO** 

November 13–14, 2024 | Online

#### The CLASSIX Story

#### Developing the same algorithm in MATLAB and Python simultaneously

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MathWorks





A data clustering method that is

- fast
- memory efficient
- conceptually very simple
- non-iterative and 100% deterministic
- easy to tune with two hyperparameters
- computing fully explainable clustering results

Currently two implementations

- Python <a href="https://github.com/nla-group/classix">https://github.com/nla-group/classix</a>
- MATLAB <u>https://github.com/nla-group/classix-matlab</u>



## Here are some data points we want to cluster.



# We compute the first principal component.



## And enumerate the data along that direction.





## And enumerate the data along that direction.

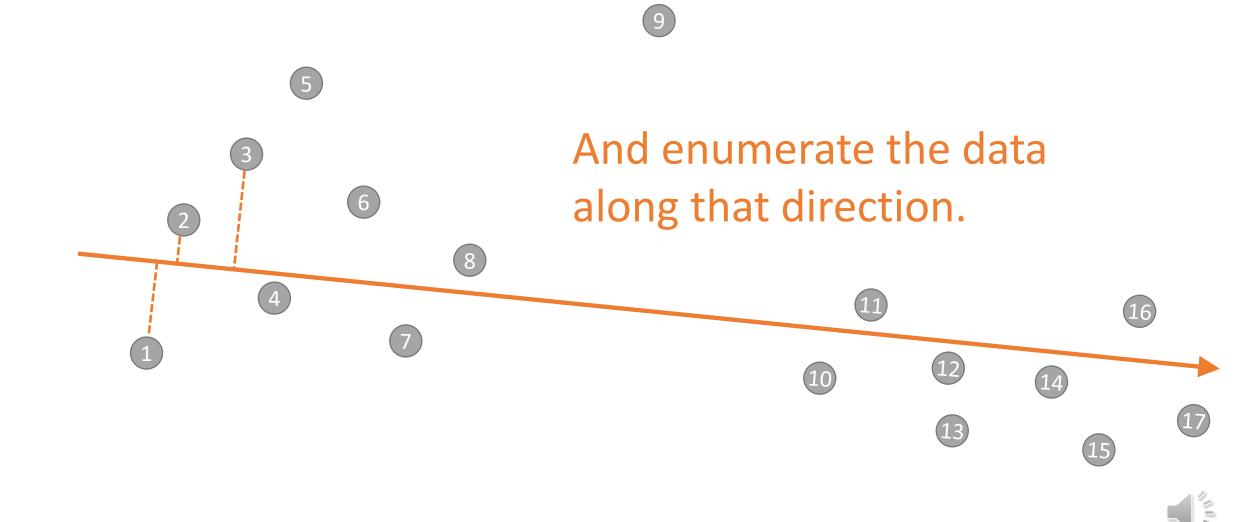




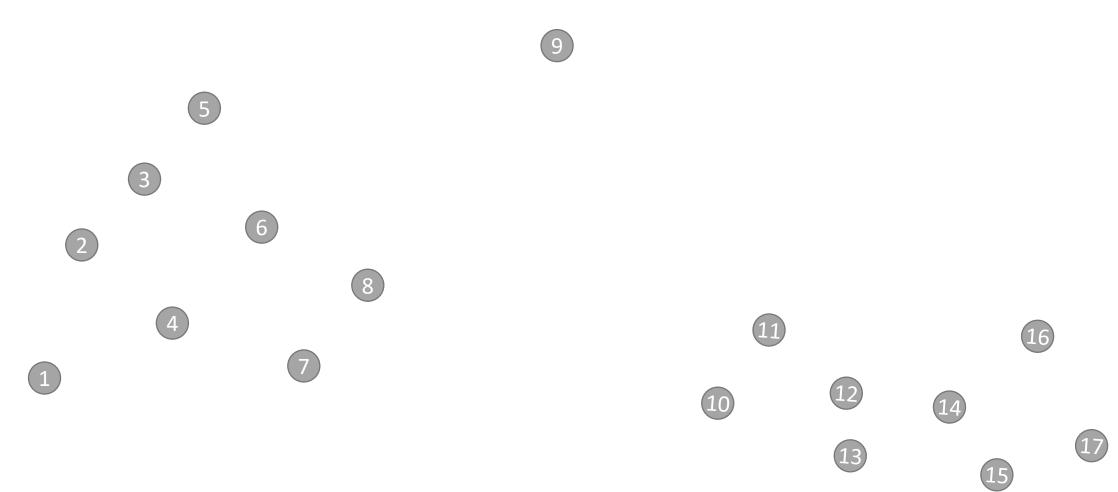
## And enumerate the data along that direction.







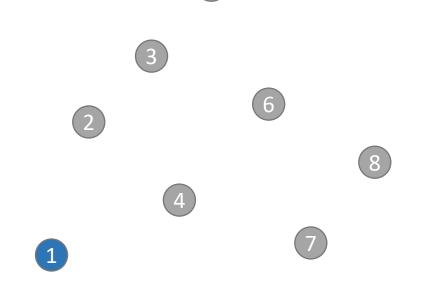




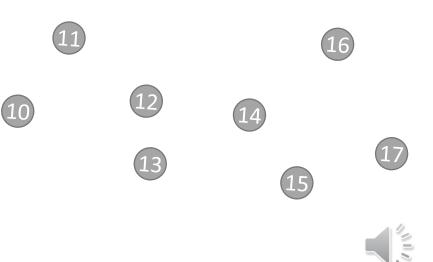




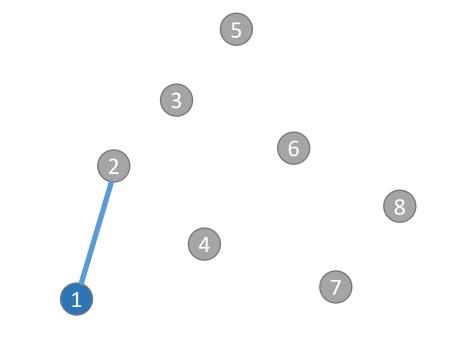
#### Starting from the first point



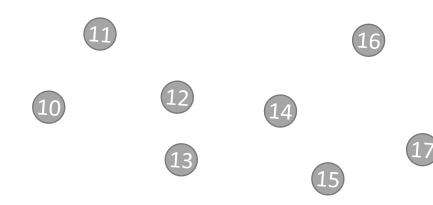
(5)



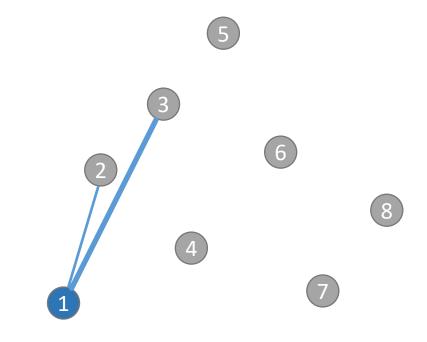




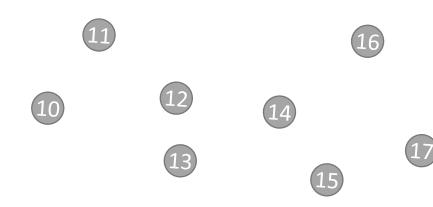




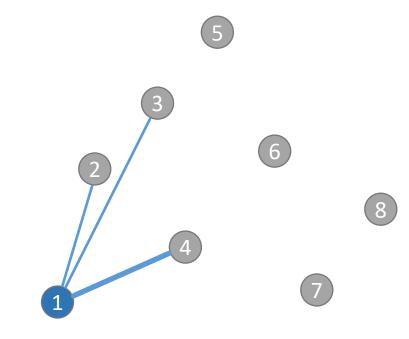




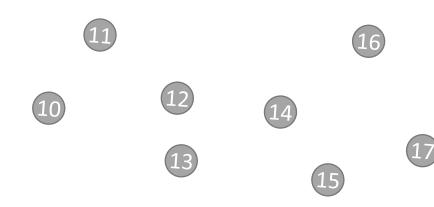




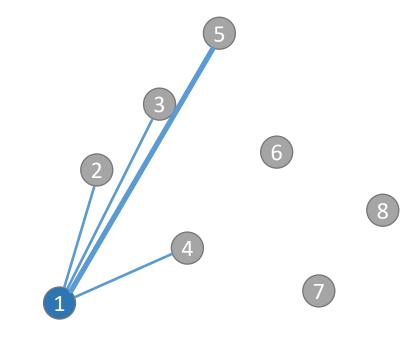




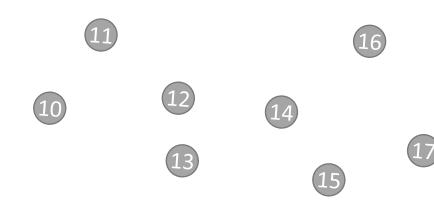




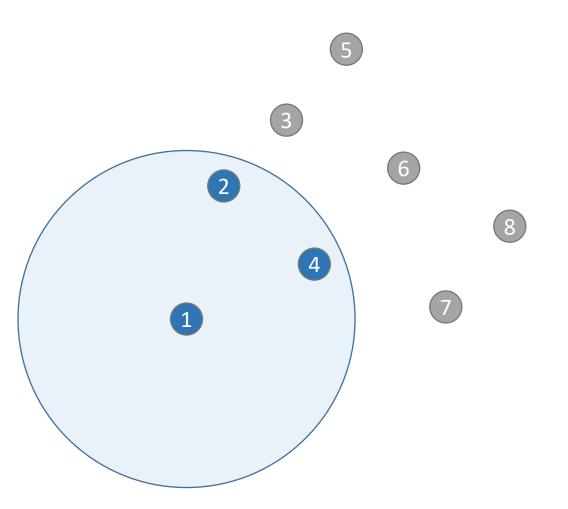






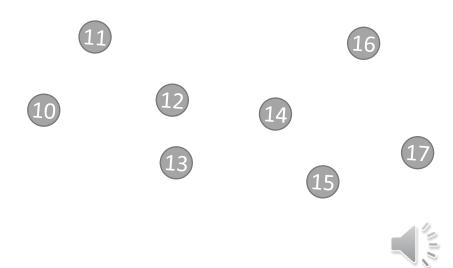




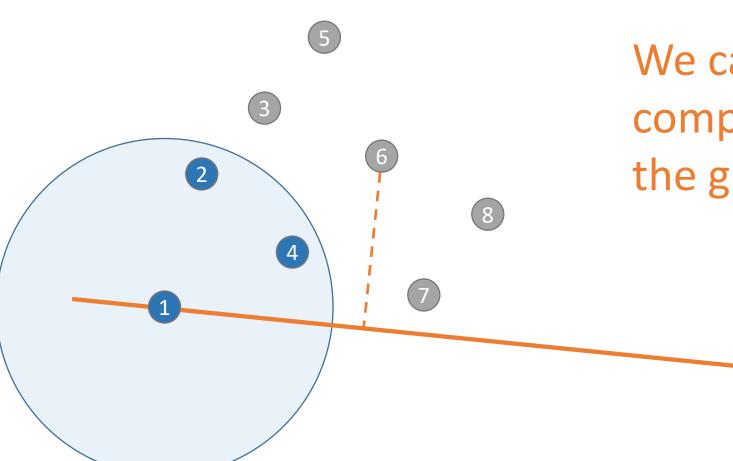




## All points within a predefined radius become part of a group.





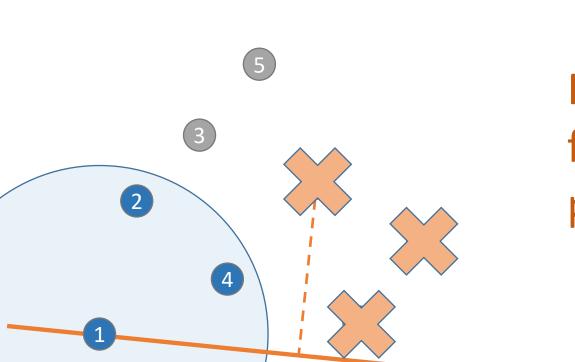




We can use the principal component to terminate the group search.

(16)

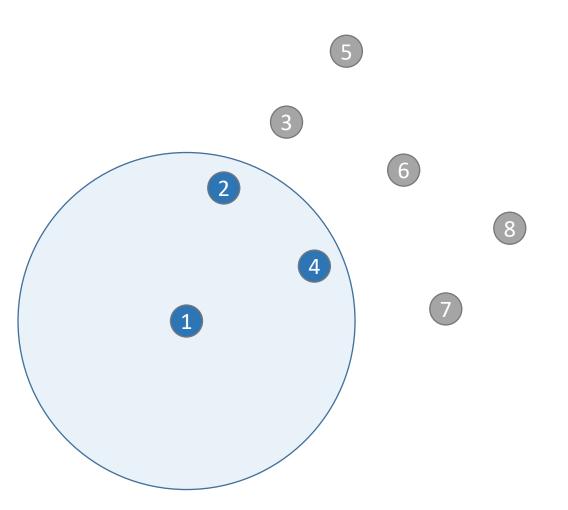






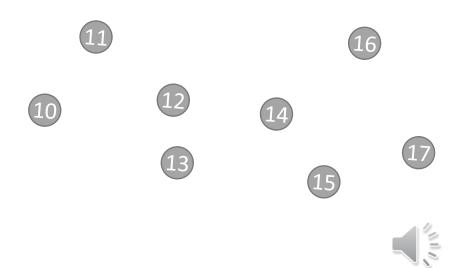
#### Point 6 and all the following ones cannot be part of the first group.



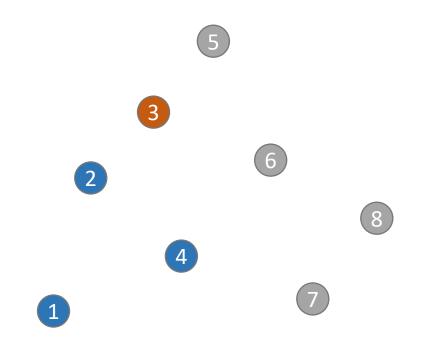




## No need for range query data structures to implement this.

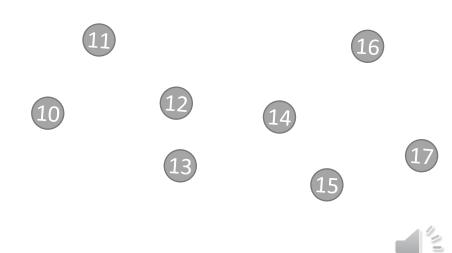




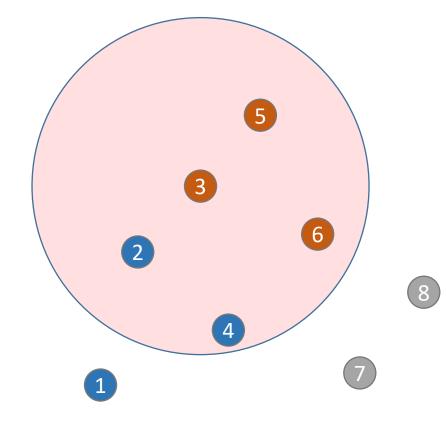




## We continue with the first unassigned data point.

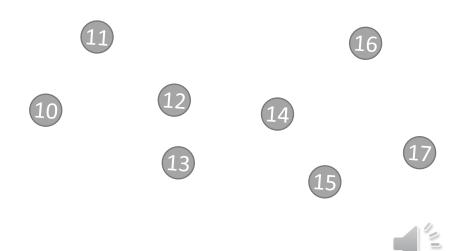




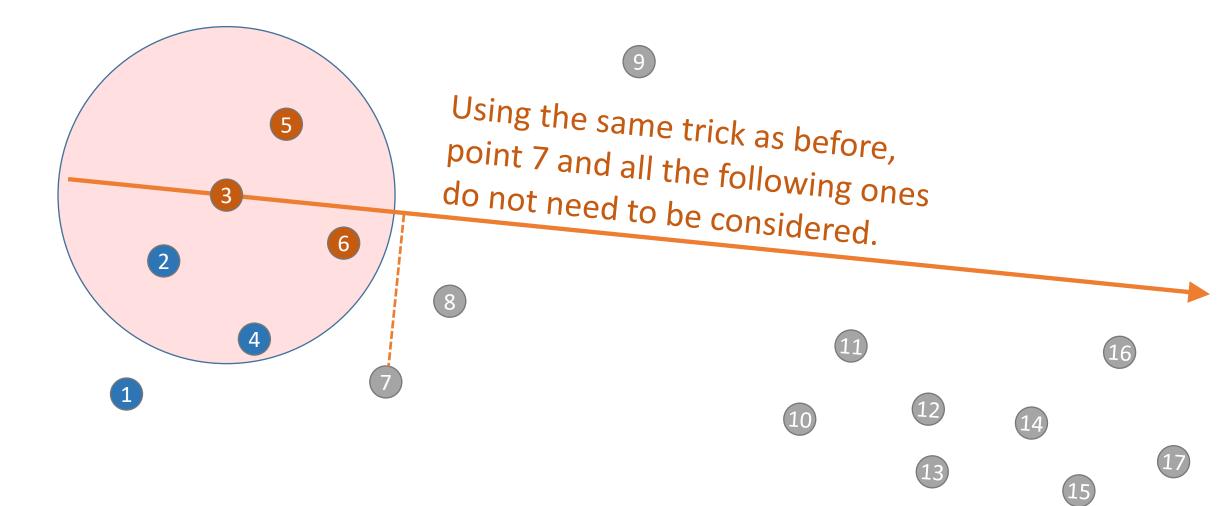




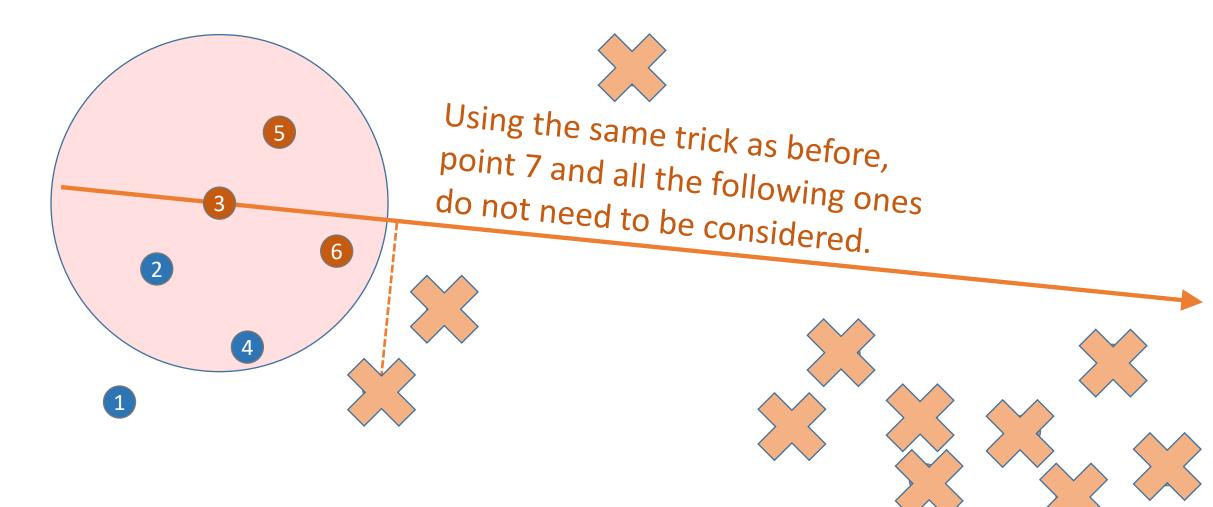
## And group all data points within the predefined radius.



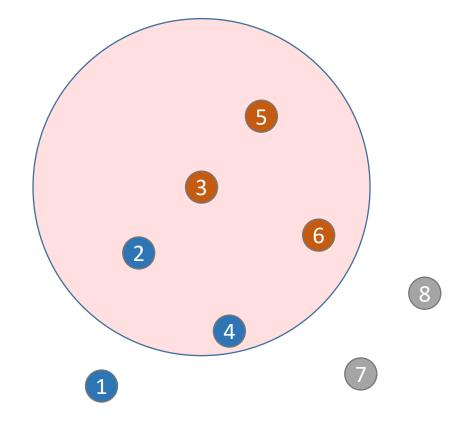






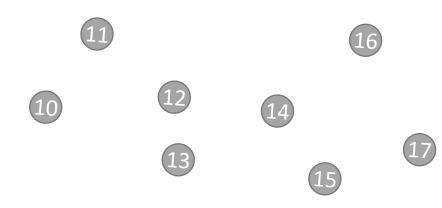




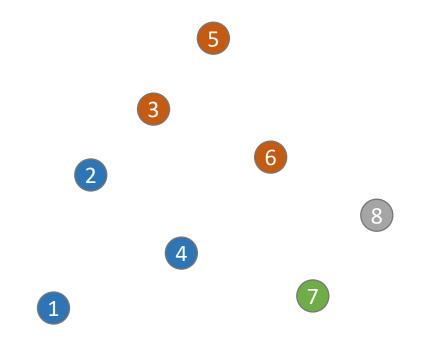




This early search termination keeps the number of distance calculations low.

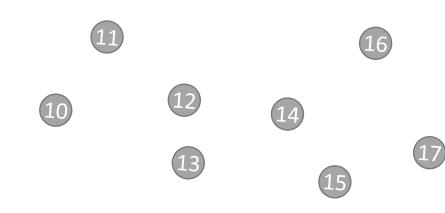




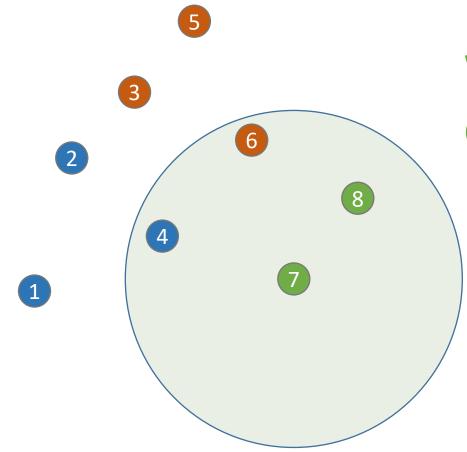




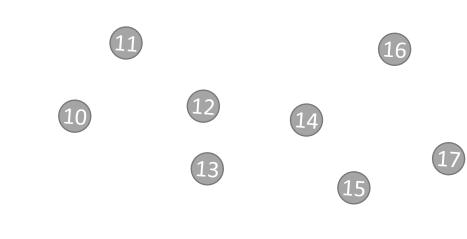
## We continue grouping the data points in their order.







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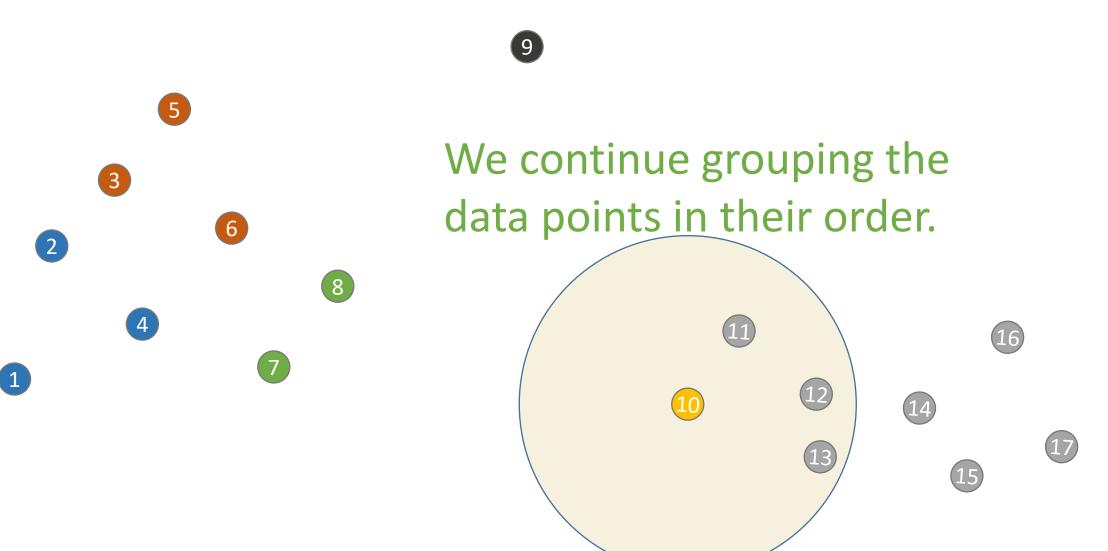
**CLASSIX** Fast and Explainable Clustering

## We continue grouping the data points in their order.

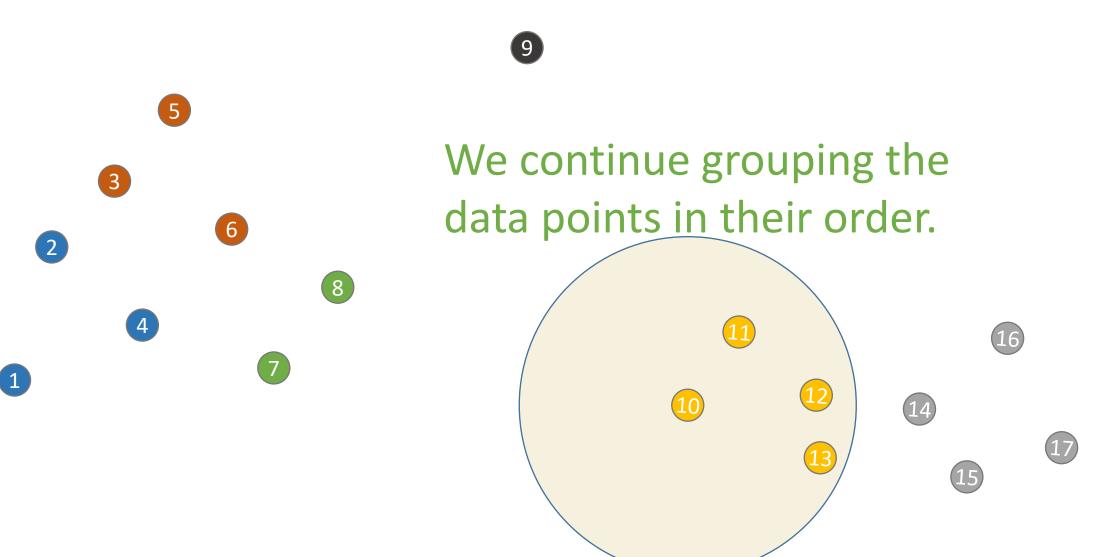
(12)

(14)

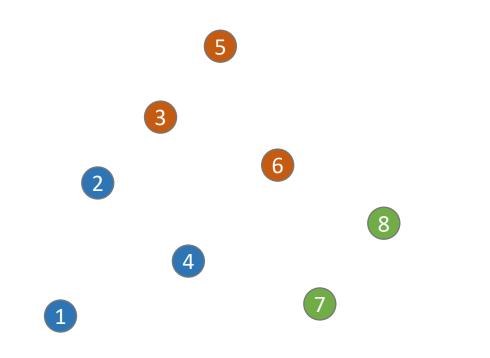




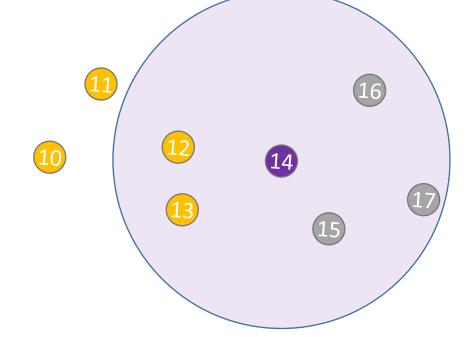




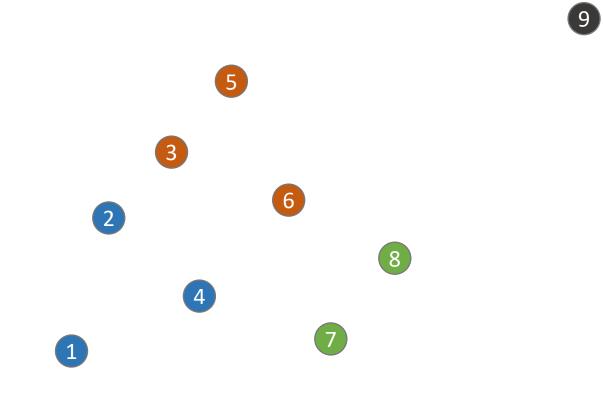




### We continue grouping the data points in their order.

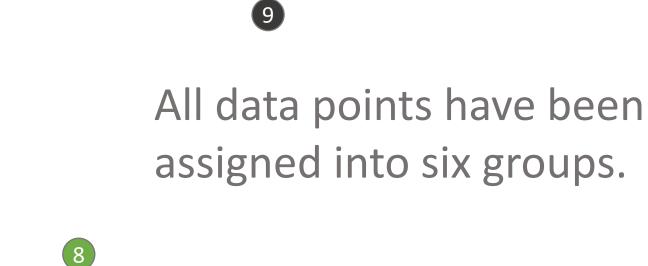


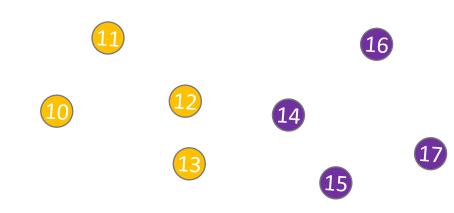












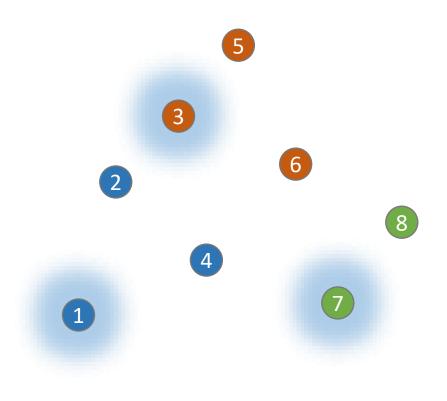




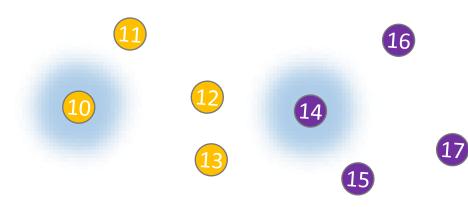
Now we just have to merge these groups into clusters.



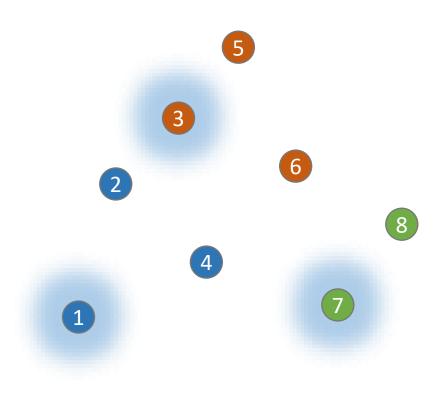




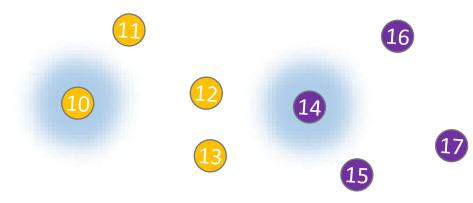
#### This can be done efficiently by using the starting points of each group.



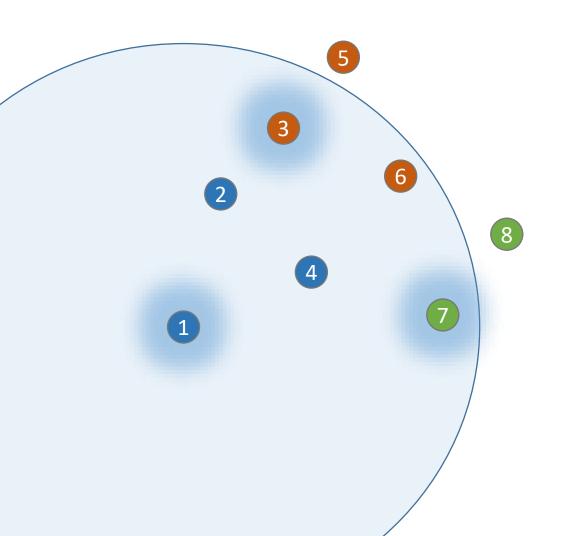




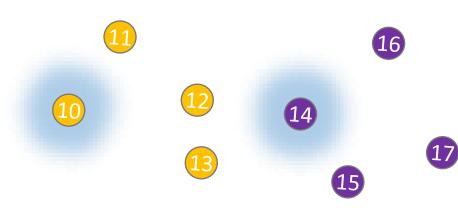
#### The starting points are already sorted and we can again use early search termination.



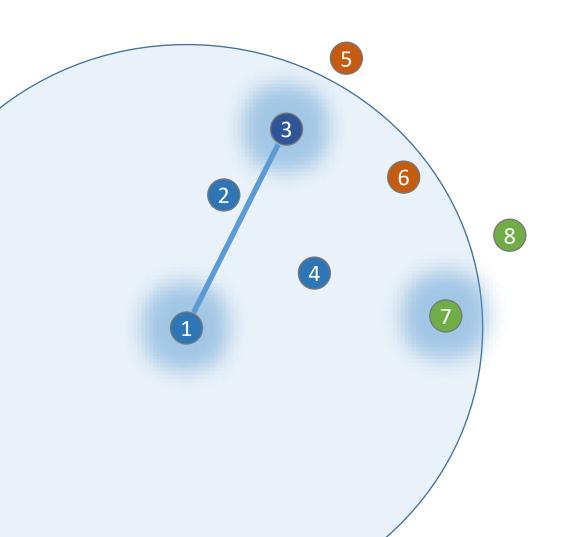




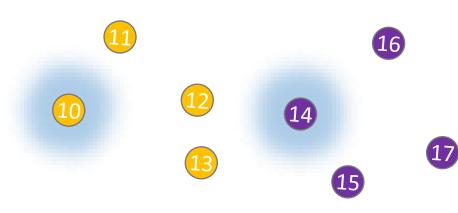
#### We merge two starting points within 1.5x the predefined radius.



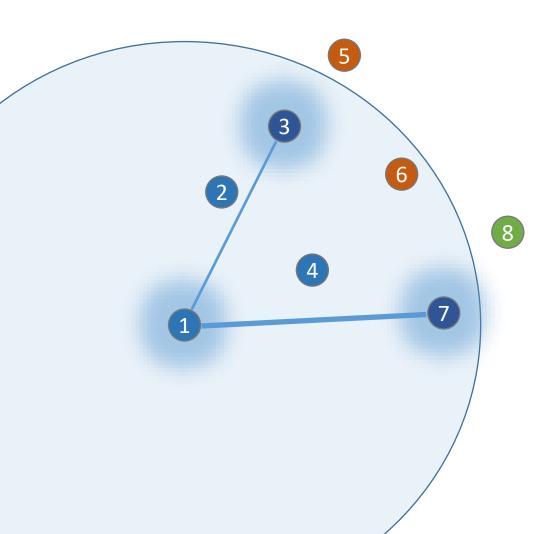




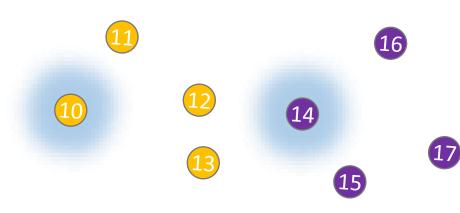
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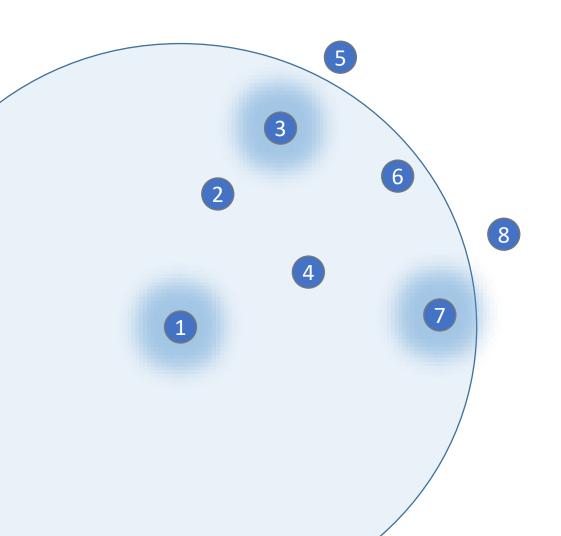




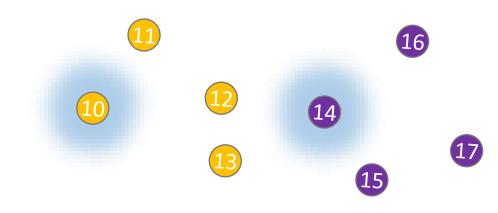
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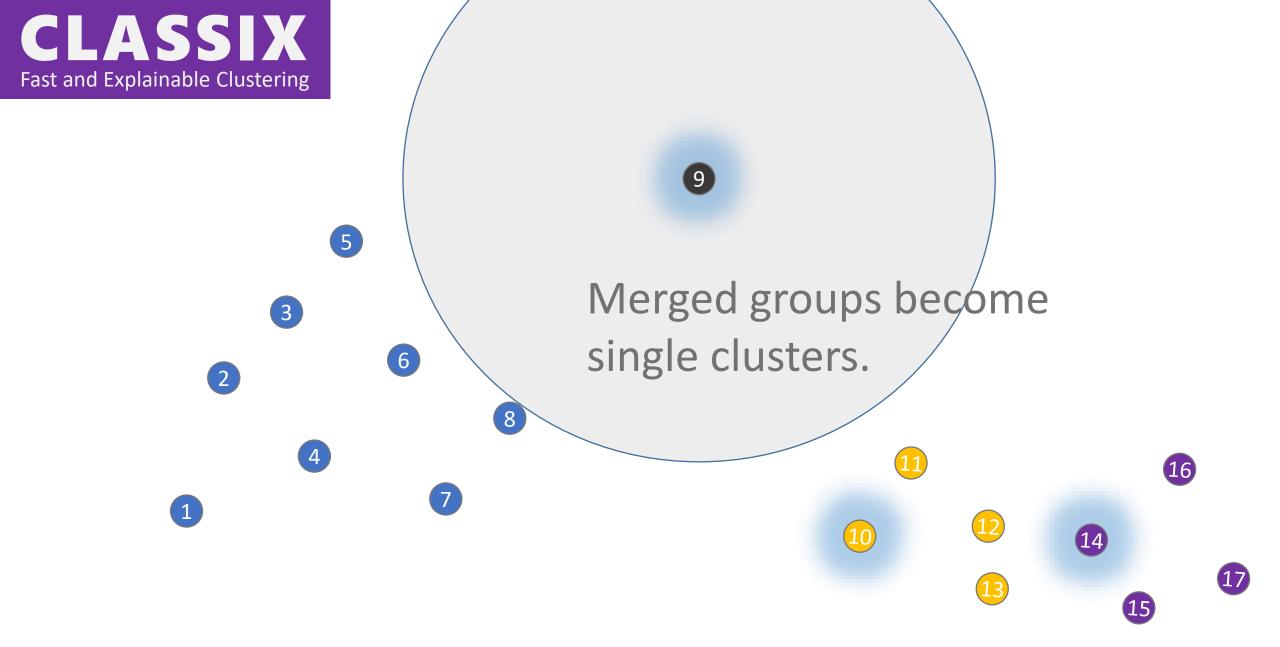




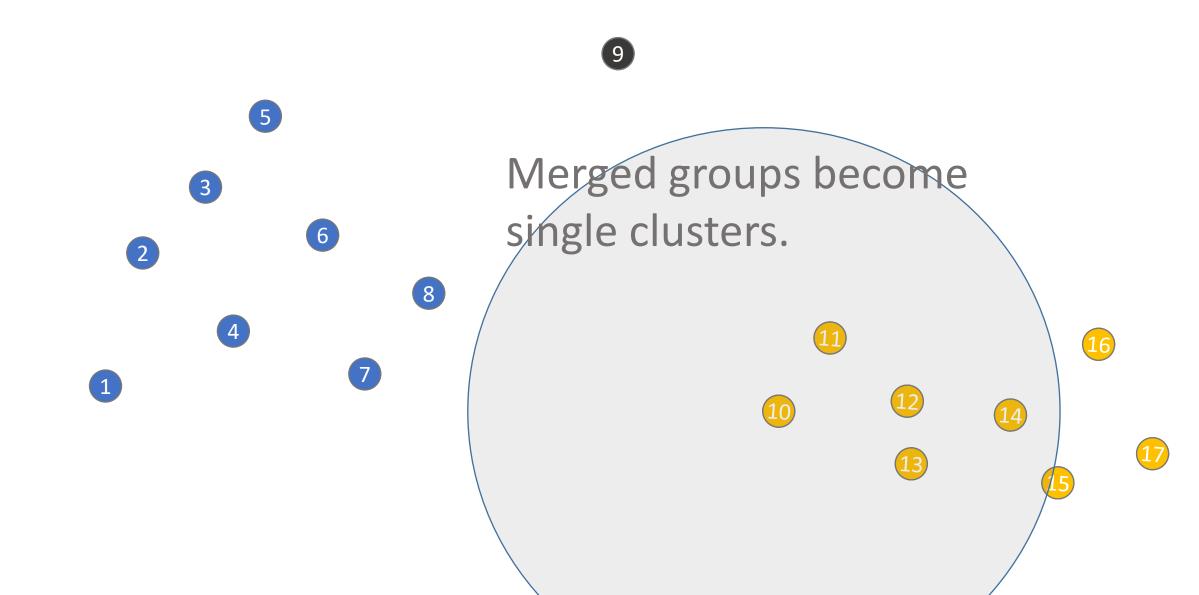


# Merged groups become single clusters.

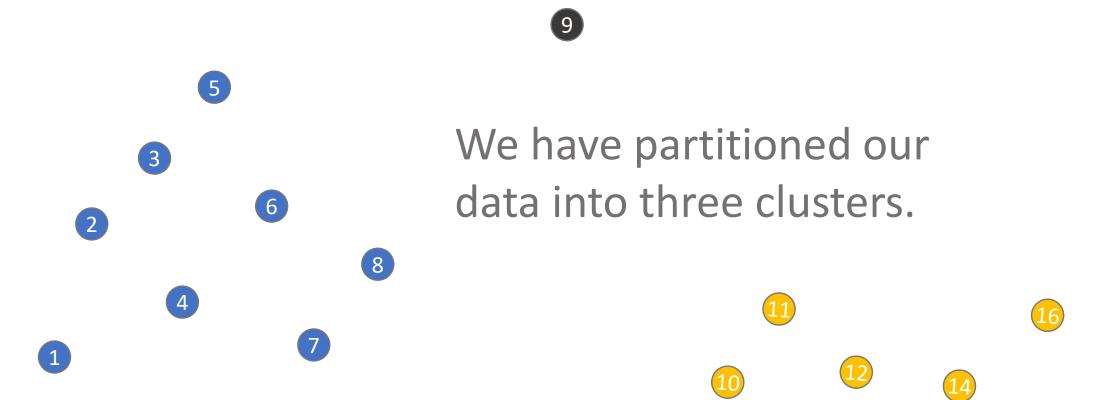








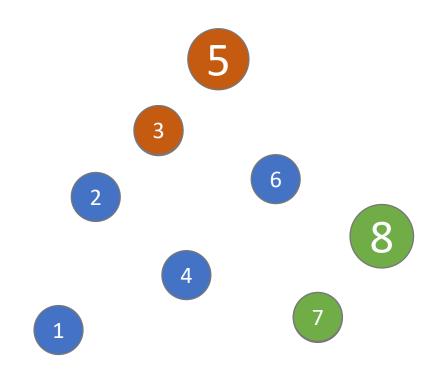




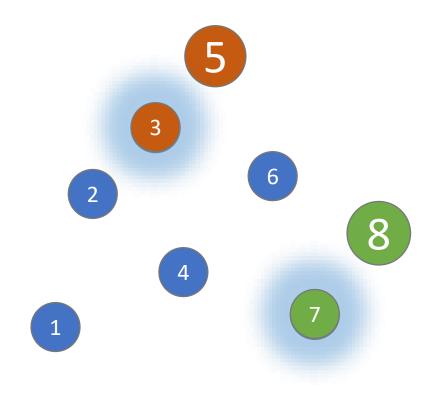


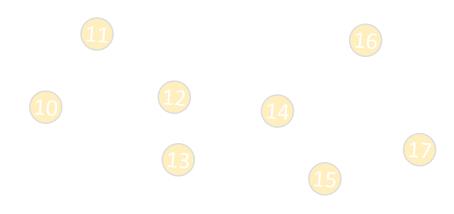




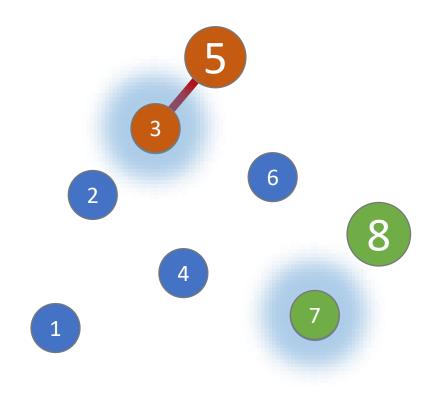


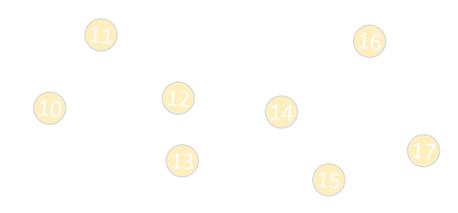




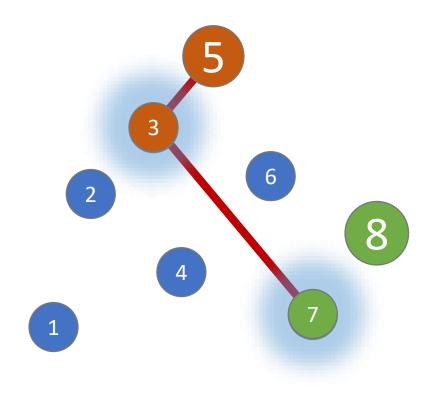




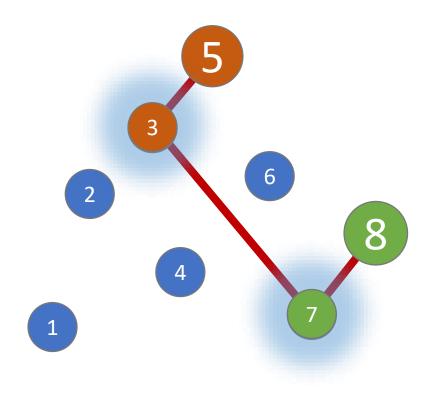


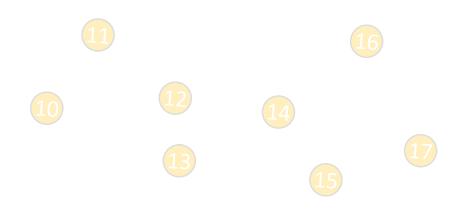




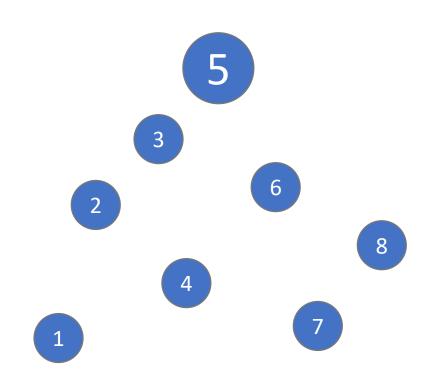




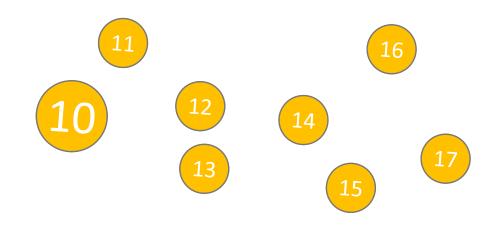






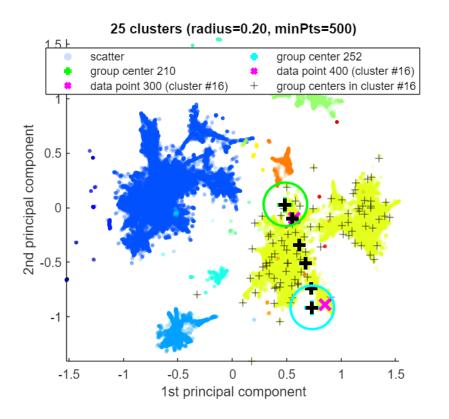


And there is no such path between points 5 and 10, hence different clusters.





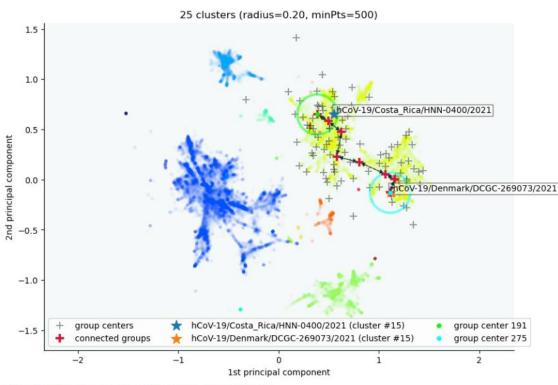
### MATLAB demo Clustering 5.7M RNA sequences of coronavirus



	CLASSIX	DBSCAN (5%)
Parameters	(0.2, 500)	(0.1, 1)
Runtime (s)	6.7	493
Clusters	25	39
AMI	0.61	0.60



### Same in Python Clustering 5.7M RNA sequences of coronavirus



Data point hCoV-19/Costa\_Rica/HNN-0400/2021 is in group 191. Data point hCoV-19/Denmark/DCGC-269073/2021 is in group 275. Both groups were merged into cluster #15.

The two groups are connected via groups 191 <-> 210 <-> 239 <-> 230 <-> 258 <-> 272 <-> 279 <-> 275.

	CLASSIX.py	HDBSCAN
Parameters	(0.2, 500)	(180000, 5)
Runtime (s)	7.7	4080
Clusters	25	4
AMI	0.61	0.59



### MATLAB vs Python

#### Timing comparison on URI machine learning datasets

	Dim	Size	#Classes	CLASSIX.m	CLASSIX.mex	CLASSIX.py
Banknote	4	1372	2	0.044	0.031	0.078
Dermatology	34	366	6	0.019	0.017	0.047
Ecoli	7	336	7	0.015	0.012	0.028
Glass	9	214	26	0.009	0.009	0.013
Iris	4	150	4	0.008	0.007	0.010
Seeds	7	210	3	0.013	0.010	0.029
Wine	13	178	2	0.011	0.010	0.020
Phoneme	256	4509	4	20.861	6.195	5.369
VDU Signals	2	2028780	11	1.028	1.034	2.860



## How did we get there?

A tale of developing the same algorithm in two languages simultaneously

### Step 0: Original version of CLASSIX

Developed in Python with PhD student Xinye Chen between 2021–2022

First arXiv preprint in February 2022 and GitHub release

Used inefficient disjoint set data structure for keeping track of clusters

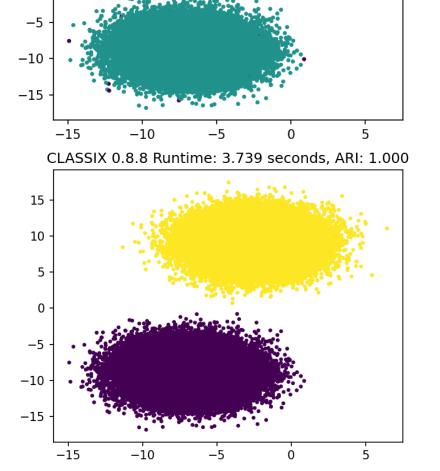
Still significantly faster than e.g. DBSCAN



### Step 0: Original version of CLASSIX

```
print('DBSCAN (sklearn 1.5.2)')
st = time()
clustering = DBSCAN(eps=3, min_samples=5).fit(X)
print(' Runtime:', time()-st, 'seconds')
print(' ARI: _', ari(clustering.labels_, y))
print('CLASSIX.py version 0.8.8')
st = time()
clx = <u>CLASSIX(radius=0.2, minPts=5)</u>
clx.fit(X)
print(' Runtime:', time()-st, 'seconds')
```

print(' ARI: ', ari(clx.labels , y))



DBSCAN Runtime: 28.344 seconds, ARI: 1.000

15

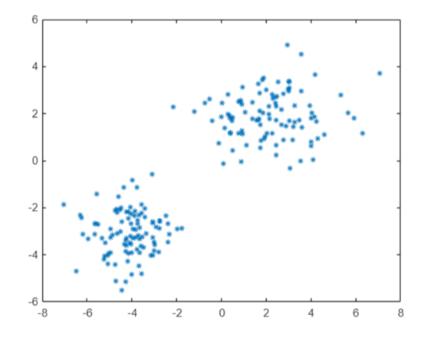
10 -

5

0

### Step 1: Using interoperability to run CLASSIX.py in MATLAB

rng('default') % For reproducibility mu1 = [2 2]; % Mean of the 1st cluster sigma1 = [2 0; 0 1]; % Covariance of the 1st cluster mu2 = [-4 -3]; % Mean of the 2nd cluster sigma2 = [1 0; 0 1]; % Covariance of the 2nd cluster r1 = mvnrnd(mu1,sigma1,100);| r2 = mvnrnd(mu2,sigma2,100); X = [r1; r2]; plot(X(:,1),X(:,2),"\*",MarkerSize=5);



Calling CLASSIX is straightforward. We don't even need to convert the MATLAB array X to a Numpy array as it's all done automatically.

clx = py.classix.CLASSIX(radius=0.3, verbose=0); clx = clx.fit(X);

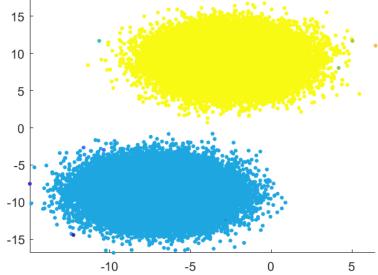
clx.explain(plot=false);

CLASSIX clustered 200 data points with 2 features. The radius parameter was set to 0.30 and winnes was set to 0. MATLAB Meets Python: Amplifying Research Impact with Cross-Platform Integration - MATLAB (mathworks.com)

### Step 2: Write native MATLAB version based on Python original

Profiler						- 0
PROFILER						
→ Back Fint Profile → Forward		code to run and time		Start Profiling		
LE NAVIGATE	SEARCH VIEW		PROFILE			
Function Name			Function Type			Calls
classix_mike>classix_mike.fast_ago	glomerate		Class method			30566
						l
Lines that take the most time	9					
Line Number	Code		Calls	Total Time (s)	% Time	Time Plot
25	<pre>parent_of_s2 = findParent(s2);</pre>		30566	0.205	49.6%	
<u>24</u>	<pre>parent_of_s1 = findParent(obj);</pre>		30566	0.146	35.4%	
<u>30</u>	end		30566	0.038	9.2%	•
27	if (parent_of_sl.data ~= parent_	_of_s2.data)	30566	0.002	0.5%	
<u>28</u>	<pre>parent_of_sl.parent = parent_of_</pre>	_s2;	3325	0.002	0.5%	
All other lines				0.020	4.9%	
Totals				0.414	100%	
<ul> <li>Children (called functions)</li> </ul>						
Function Name		Function Type	Calls	Total Time (s)	% Time	Time Plot
SET>SET.findParent	SET>SET.findParent Class method		61132	0.320	77.3%	
Salf time (built-ins overhead atc)	\ \			0.001	22.2%	

CLASSIX.m Runtime: 2.374 seconds, ARI 1.000

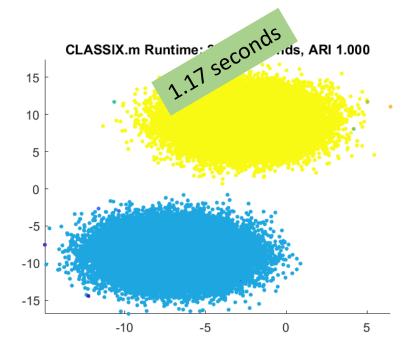


Function Name	Function Type	Calls	Total Time (s)	% Time	Time Plot
SET>SET.findParent	Class method	61132	0.320	77.3%	
Salf time (huilt-ins overhead etc.)			0 001	22.7%	

#### Large amount of time spent on disjoint set structure operations!

### Step 2: Write native MATLAB version based on Python original

Profiler							- 0
PROFILER							
int Profile Summary NAVIGATE Forward LE NAVIGATE	Q +SEARCH	Highlight VIEW	Enter code to run and time	PROFILE	Start Profiling		
Function Name				Function Type			Calls
<pre>classix_mike&gt;classix_mike.fast_ag</pre>	<u>glomerate</u>			Class method			30566
Lines that take the most time	e						
Line Number	Code			Calls	Total Time (s)	% Time	Time Plot
<u>25</u>	parent_of_s2 = fin	ndParent(s2	2);	30566	0.205	49.6%	
24	parent_of_s1 = fin	<pre>parent_of_s1 = findParent(obj);</pre>		30566	0.146	35.4%	—
<u>30</u>	end	end		30566	0.038	9.2%	-
27	if (parent_of_s1.0	data ~= pai	cent_of_s2.data)	30566	0.002	0.5%	
<u>28</u>	parent_of_s1.paren	nt = parent	_of_s2;	3325	0.002	0.5%	
All other lines					0.020	4.9%	I
Totals					0.414	100%	
Children (called functions)							
Function Name			Function Type	Calls	Total Time (s)	% Time	Time Plot
SET>SET.findParent			Class method	61132	0.320	77.3%	
Self time (built-ins overhead etc	١				0.004	22.7%	



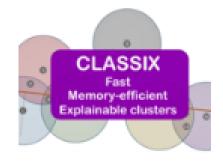
#### Large amount of time spent on disjoint set structure operations!

#### MATLAB profiler helped us to improve both MATLAB and Python versions

#### Step 3: Use insights to rewrite both Python and MATLAB versions

We used what we learned in Python to improve Mike's MATLAB code and got another significant speed-up.

The MATLAB version is currently faster than the Python package



#### Fast and explainable clustering with CLASSIX

Version 1.3 (23.8 MB) by Stefan Güttel

A fast and easy-to-use clustering method that provides explanations for the computed clusters. https://github.com/nla-group/classix-matlab

Follow

CLASSIX.py v0.8.8	CLASSIX.m v0.1	CLASSIX.m v1.0	CLASSIX.py v1.0.0	CLASSIX.m v1.3	CLASSIX.py v1.2.5
3.74 seconds	2.37 seconds	1.17 seconds	3.14 seconds	0.85 seconds	0.94 seconds
	not using minPts			using mex for submatrix-vector product	

#### Step 4: Use this work to help make a faster MATLAB (R2024b)

Bcols = 100000; A = rand(1,1000); B = randn(1000,Bcols); tic;C0 = A\*B; t0=toc; fprintf("Timing full A\*B: %.3e seconds\n",t0);

Timing full A\*B: 3.054e-02 seconds

tic;C1=A\*B(:,1:Bcols/2);t1=toc;

fprintf("Timing A\*B using only half of the columns of B: %.3e seconds\n",t1);fprintf("It is %.3fx faster to compute the full product",t1/t0)

Timing A\*B using only half of the columns of B: 2.176e-01 seconds It is 7.126x faster to compute the full product

tic;C2 = (A, 'none', B, 'none', 1, Bcols/2);t3=toc;
fprintf("Using mtimesColumns on the first half of the matrix B is %.3fx faster than doing the full product\n",t0/t3);

Using mtimesColumns on the first half of the matrix B is 1.982x faster than doing the full product

This is an undocumented function and is a proof of concept that will probably change in the future!



- Make a Python package available to MATLAB users using interoperability
- Rewrite the Python code as MATLAB code
- Iterate between Python package and MATLAB toolbox, using insights from one to drive improvements in the other. New algorithm is faster in both languages than it would have been otherwise
- BONUS: MATLAB itself is improved a little

#### https://github.com/nla-group/classix-matlab

Paper: X. Chen & S. Güttel. Fast and explainable clustering based on sorting. Pattern Recognition, 150: 110298, 2024.