

Clustering methods: Part 3

Cluster validation

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Part I:

Introduction

Cluster validation

Supervised classification:

- Ground truth class labels known
- Accuracy, precision, recall

Cluster analysis:

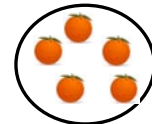
- No class labels

Validation need to:

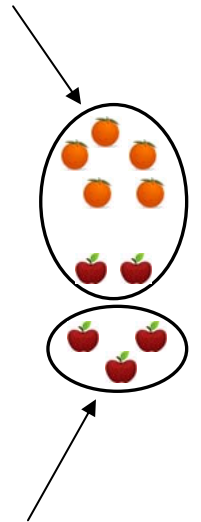
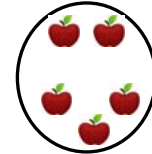
- Compare clustering algorithms
- **Solve the number of clusters**
- Avoid finding patterns in noise

Precision = $5/5 = 100\%$
Recall = $5/7 = 71\%$

Oranges:



Apples:

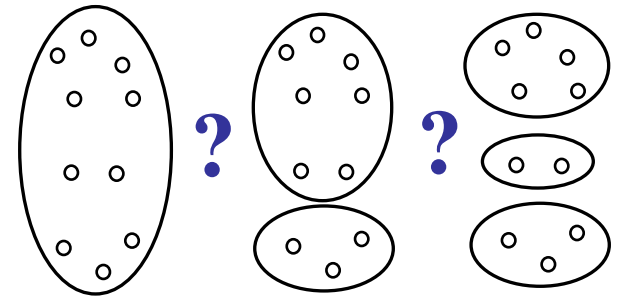


Precision = $5/5 = 100\%$
Recall = $3/5 = 60\%$

Measuring clustering validity

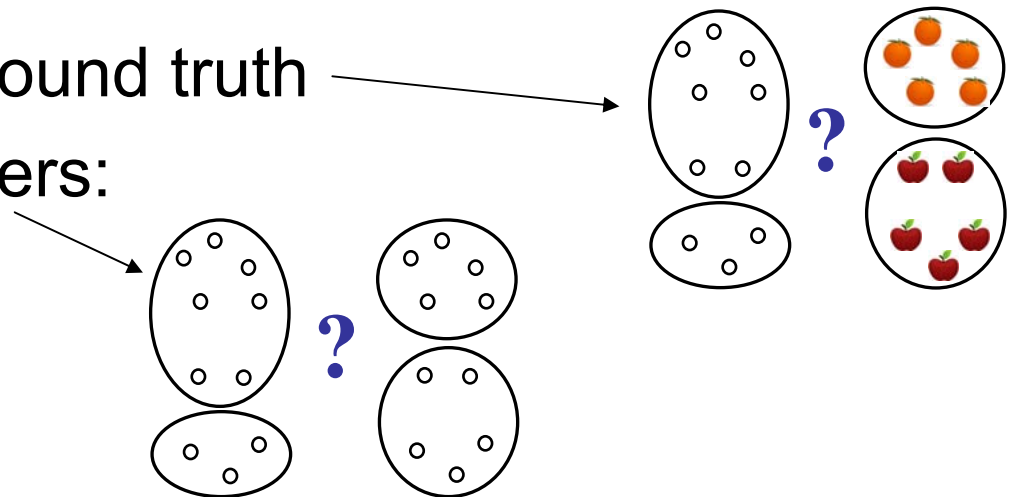
Internal Index:

- Validate *without* external info
- With different number of clusters
- Solve the number of clusters

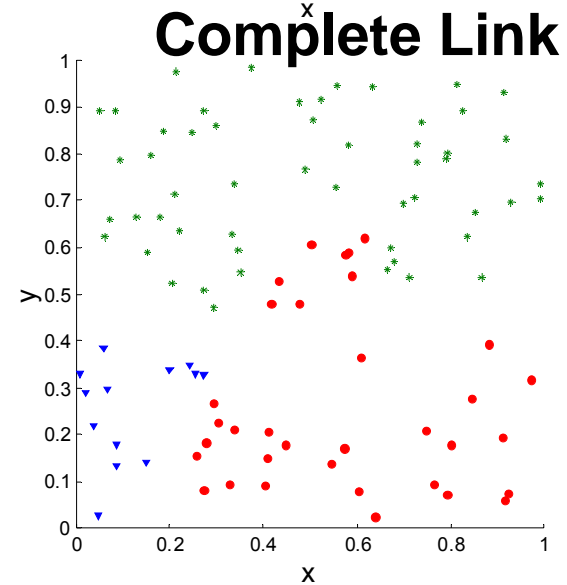
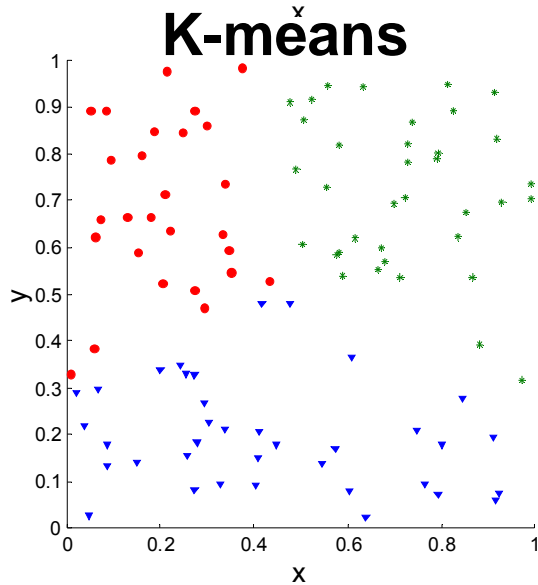
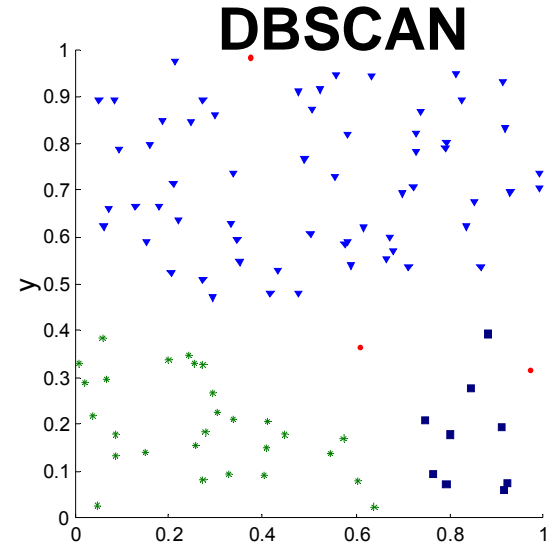
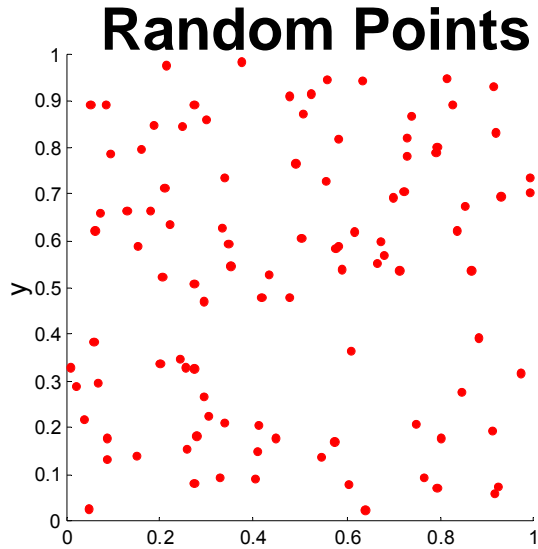


External Index

- Validate against ground truth
- Compare two clusters:
(how similar)



Clustering of random data

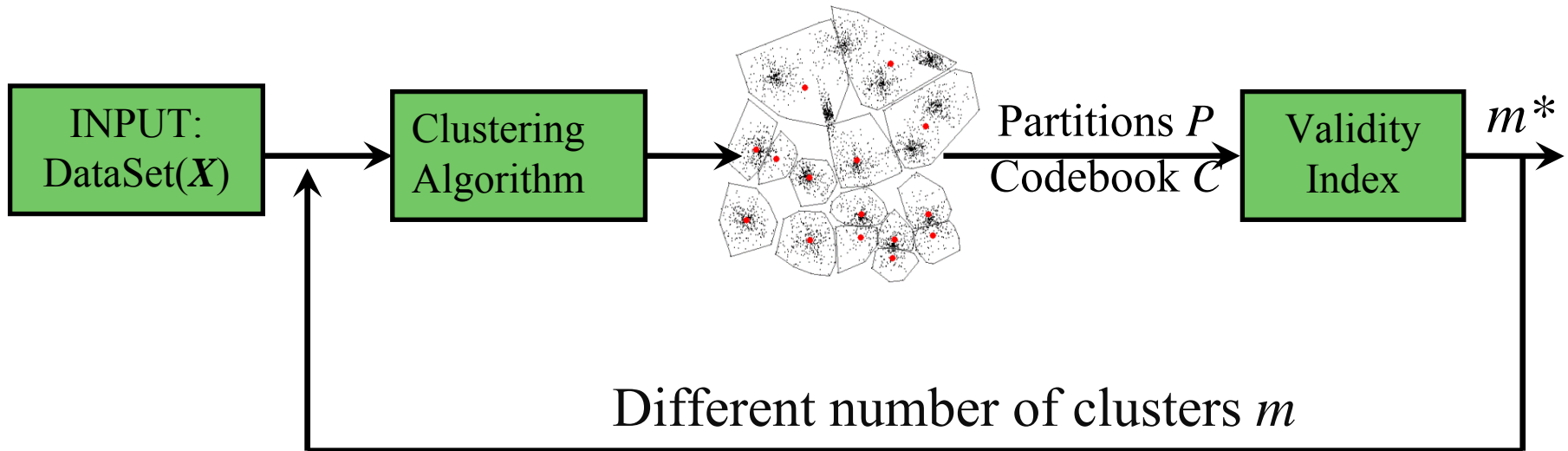


Cluster validation process

1. Distinguishing whether non-random structure actually exists in the data (one cluster).
2. Comparing the results of a cluster analysis to external ground truth (class labels).
3. Evaluating how well the results fit the data *without* reference to external information.
4. Comparing two different clustering results to determine which is better.
5. Determining the number of clusters.

Cluster validation process

- **Cluster validation** refers to procedures that evaluate the results of clustering in a **quantitative** and **objective** fashion. [Jain & Dubes, 1988]
 - How to be “quantitative”: To employ the measures.
 - How to be “objective”: To validate the measures!



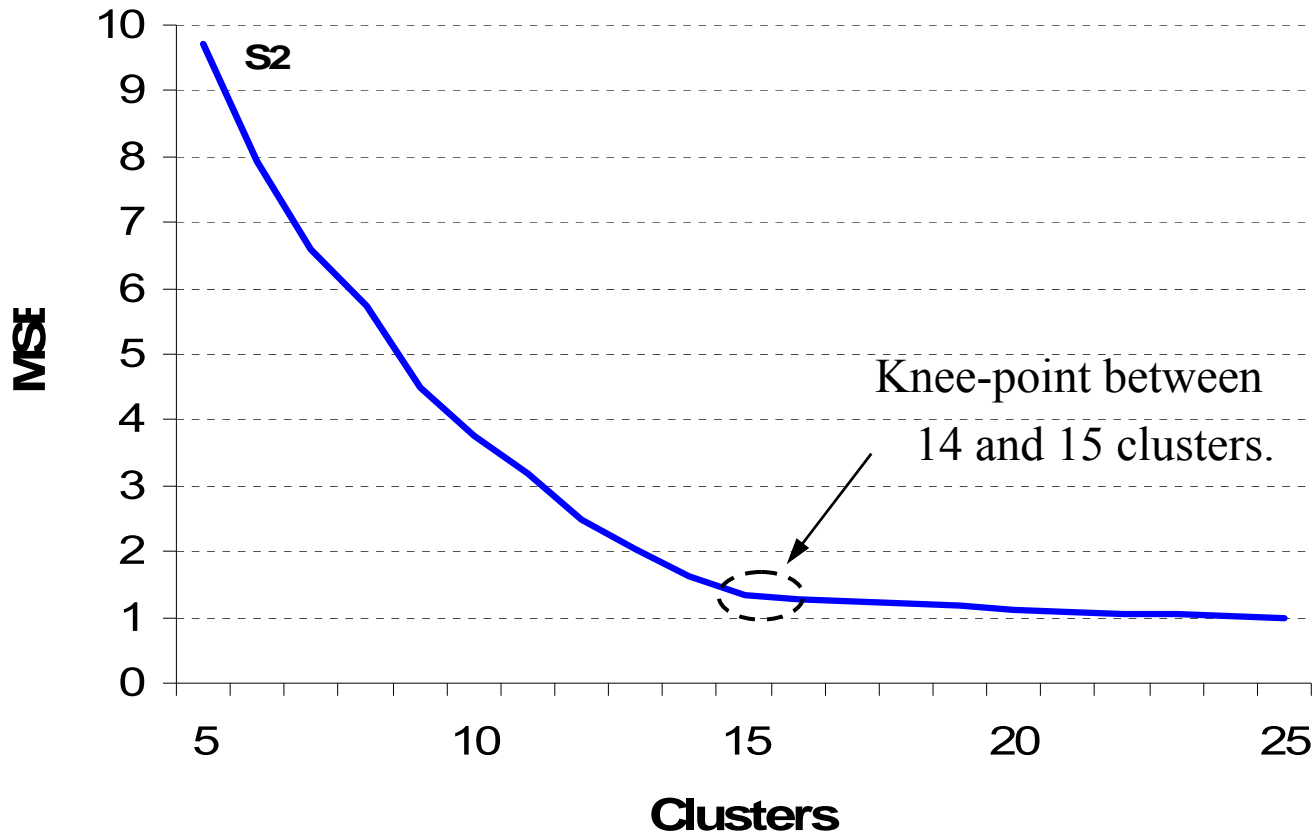
Part II:
Internal indexes

Internal indexes

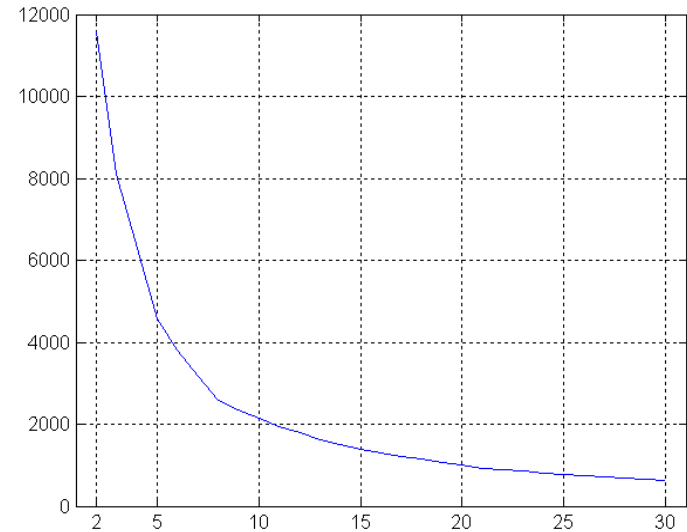
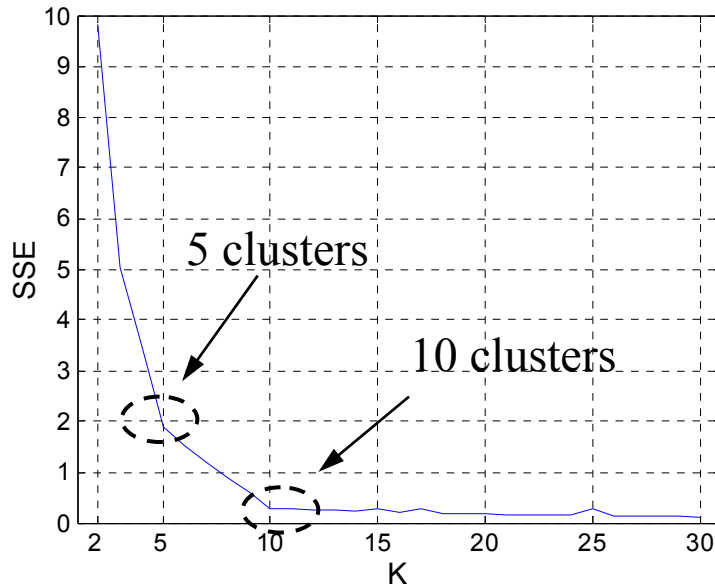
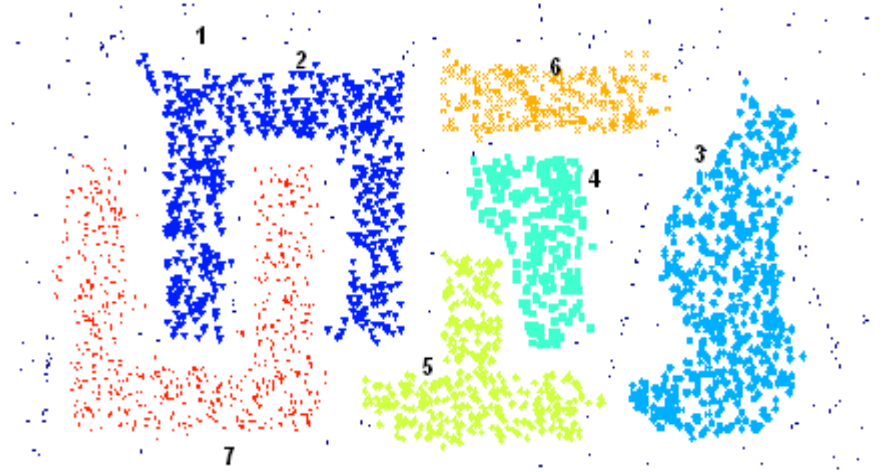
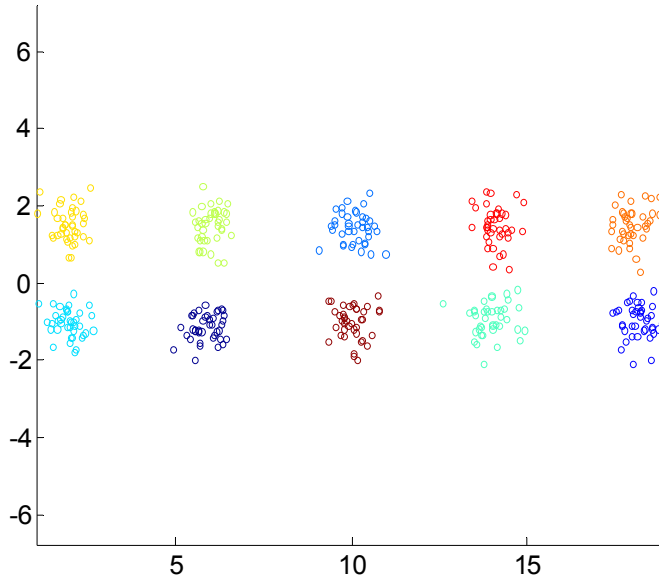
- Ground truth is rarely available but unsupervised validation must be done.
- Minimizes (or maximizes) internal index:
 - Variances of within cluster and between clusters
 - Rate-distortion method
 - F-ratio
 - Davies-Bouldin index (DBI)
 - Bayesian Information Criterion (BIC)
 - Silhouette Coefficient
 - Minimum description principle (MDL)
 - Stochastic complexity (SC)

Sum of squared errors

- The more clusters the smaller the value.
- Small knee-point near the correct value.
- But how to detect?

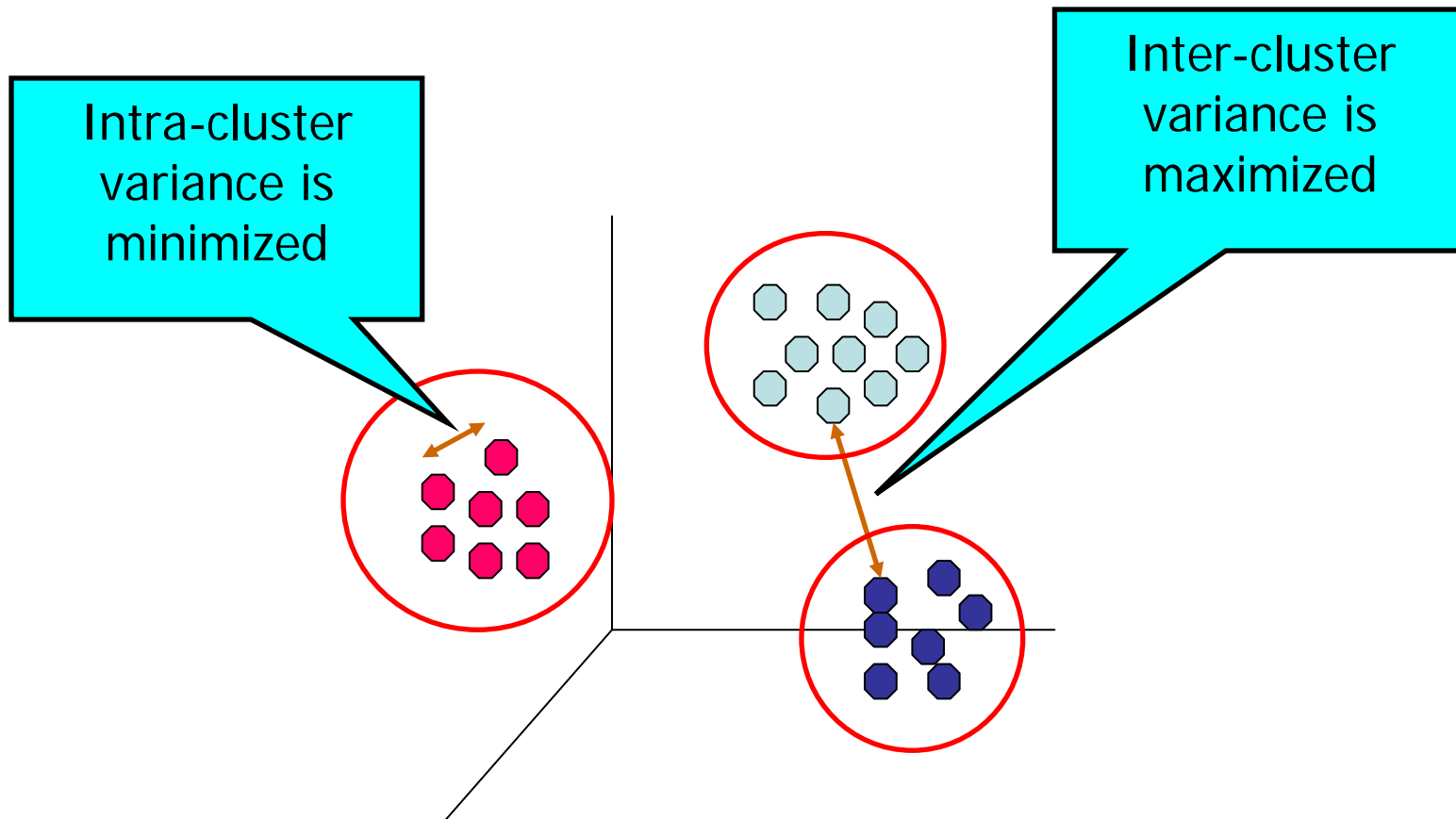


Sum of squared errors



From TSE to cluster validity

- Minimize within cluster variance (TSE)
- Maximize between cluster variance

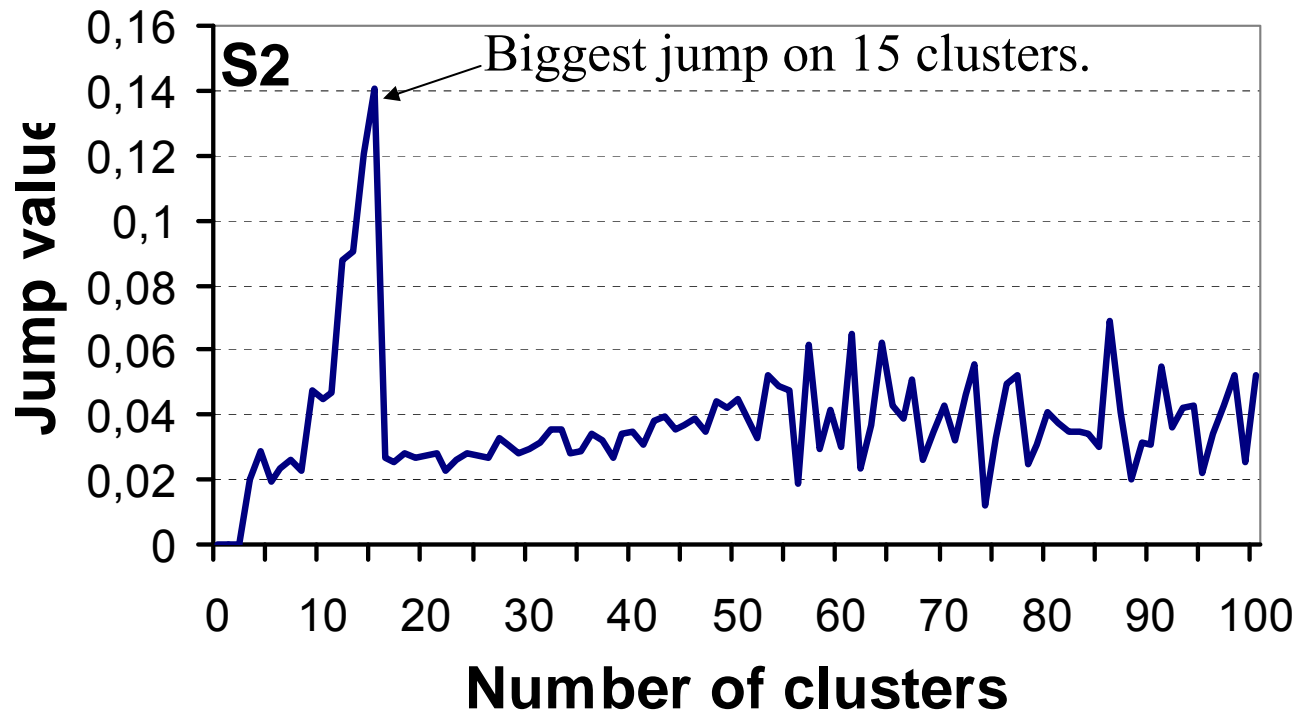


Jump point of TSE

(rate-distortion approach)

First derivative of powered TSE values:

$$J(k) = TSE(k)^{-d/2} - TSE(k-1)^{-d/2}$$



Cluster variances

Within cluster: $SSW(C, k) = \sum_{i=1}^N \|x_i - c_{p(i)}\|^2$

Between clusters: $SSB(C, k) = \sum_{j=1}^k n_j \|c_j - \bar{x}\|^2$

Total Variance of data set:

$$\sigma(X) = \underbrace{\sum_{i=1}^N \|x_i - c_{p(i)}\|^2}_{SSW} + \underbrace{\sum_{j=1}^k n_j \|c_j - \bar{x}\|^2}_{SSB}$$

WB-index

- Measures ratio of between-groups variance against the within-groups variance
- WB-index:

$$F = \frac{k \cdot \sum_{i=1}^N \|x_i - c_{p(i)}\|^2}{\sum_{j=1}^k n_j \|c_j - \bar{x}\|^2} = \frac{k \cdot SSW}{\sigma(X) - SSW}$$

SSB

Sum-of-squares based indexes

- SSW / k ----- Ball and Hall (1965)
- $k^2|W|$ ----- Marriot (1971)
- $\frac{SSB / k - 1}{SSW / N - k}$ ----- Calinski & Harabasz (1974)
- $\log(SSB/SSW)$ ----- Hartigan (1975)
- $d \log(\sqrt{SSW / (dN^2)}) + \log(k)$ ----- Xu (1997)

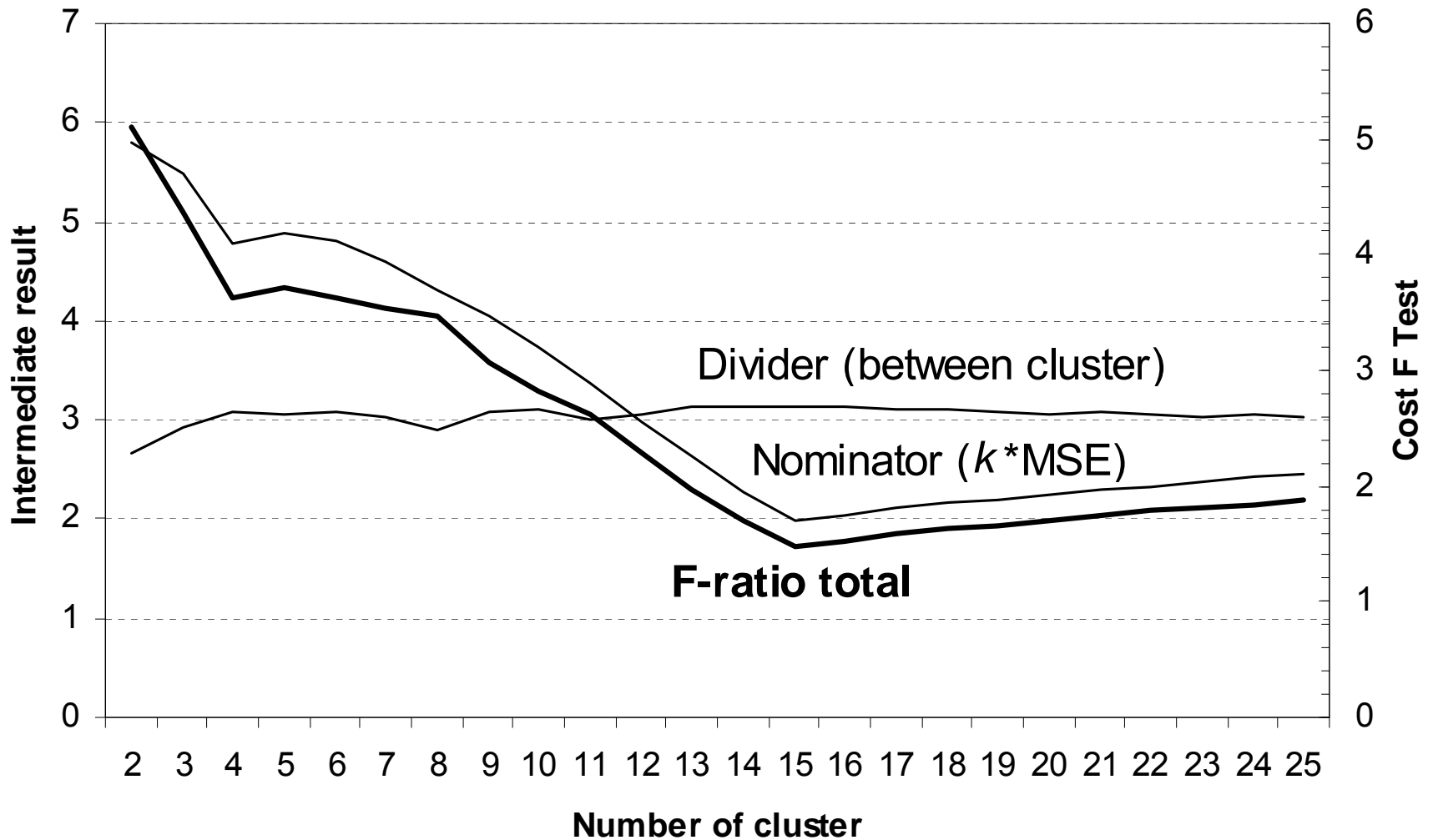
(d = dimensions; N = size of data; k = number of clusters)

SSW = Sum of squares **w**ithin the clusters (=TSE)

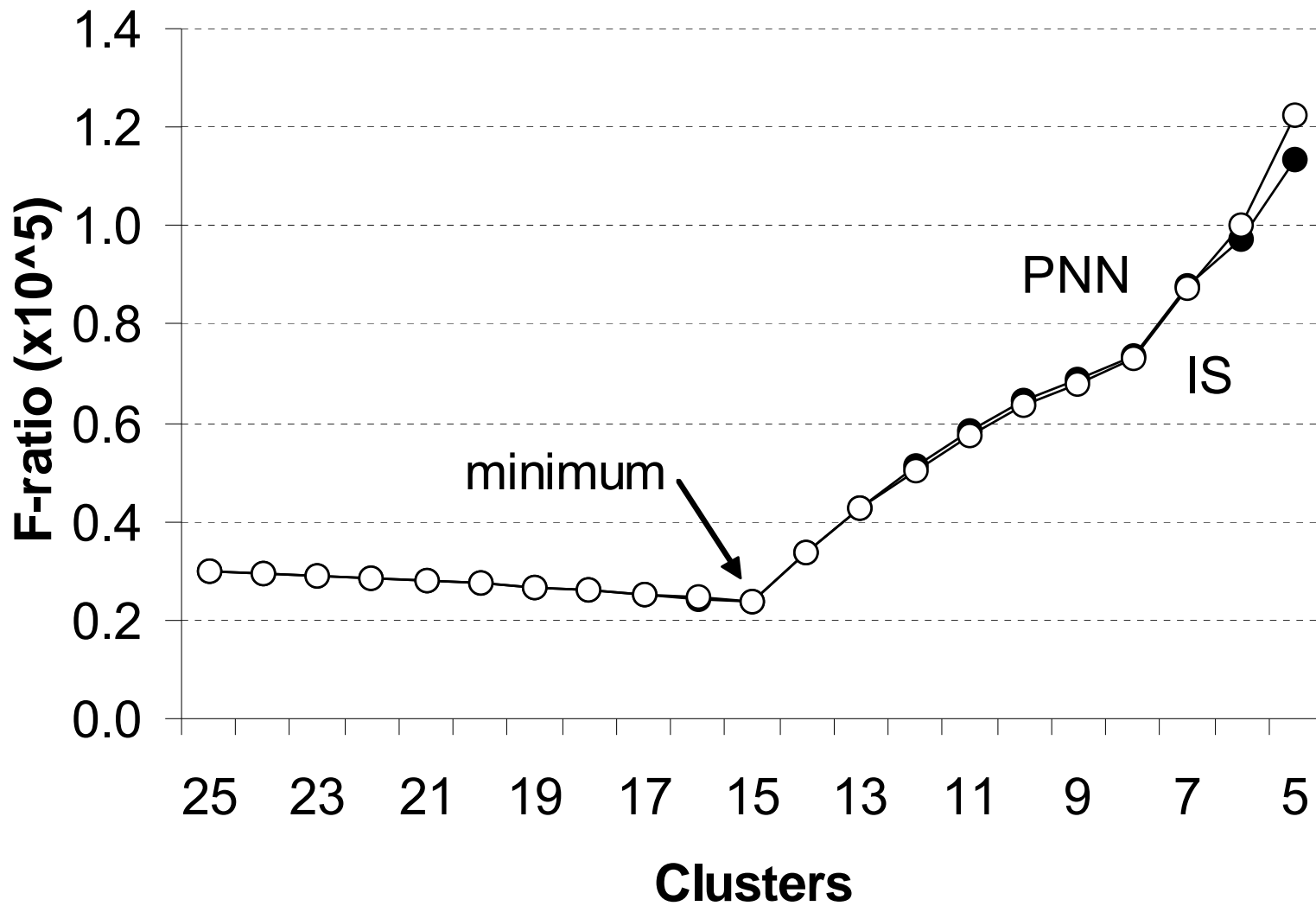
SSB = Sum of squares **b**etween the clusters

Calculation of WB-index

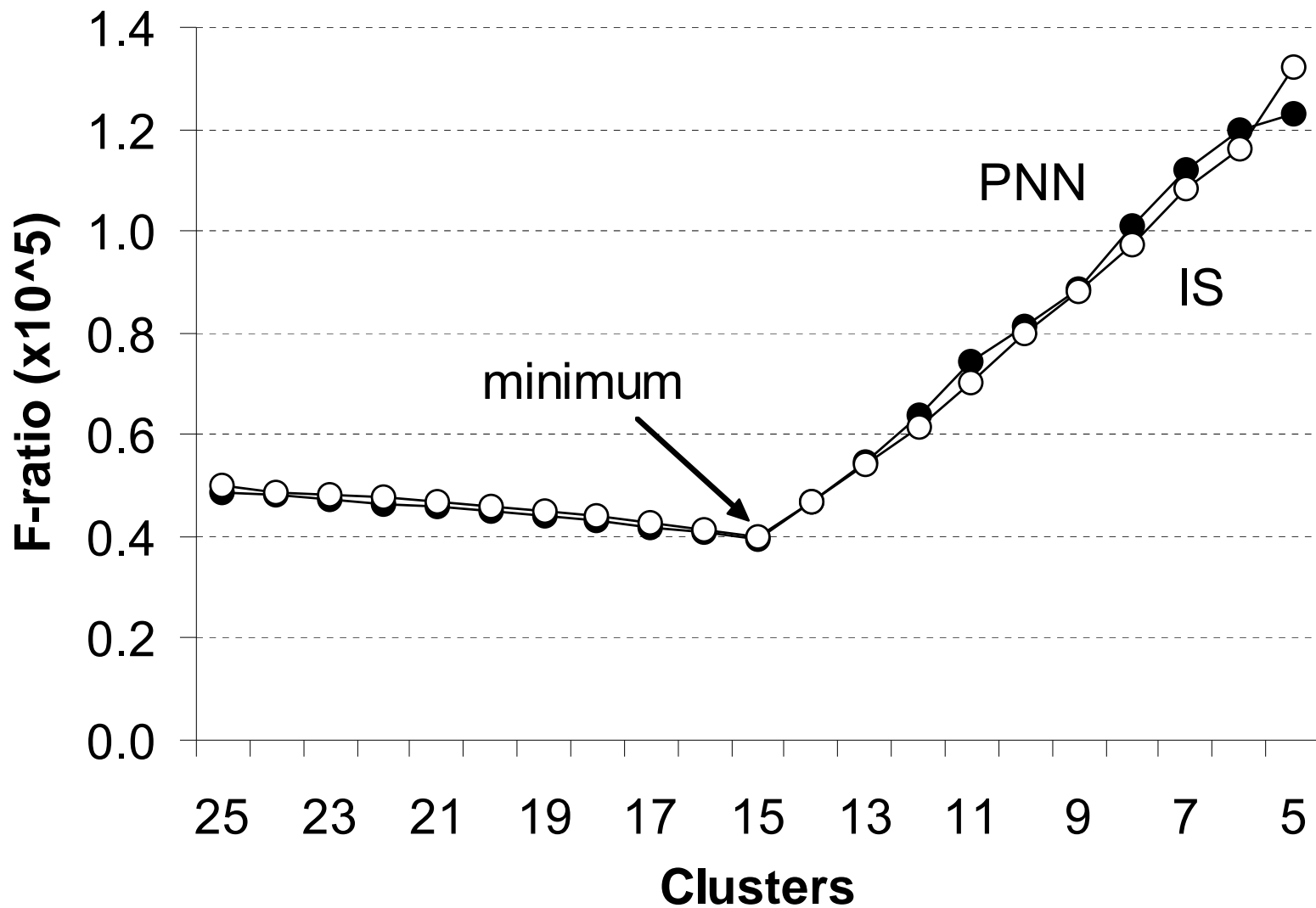
(called also F-ratio / F-test)



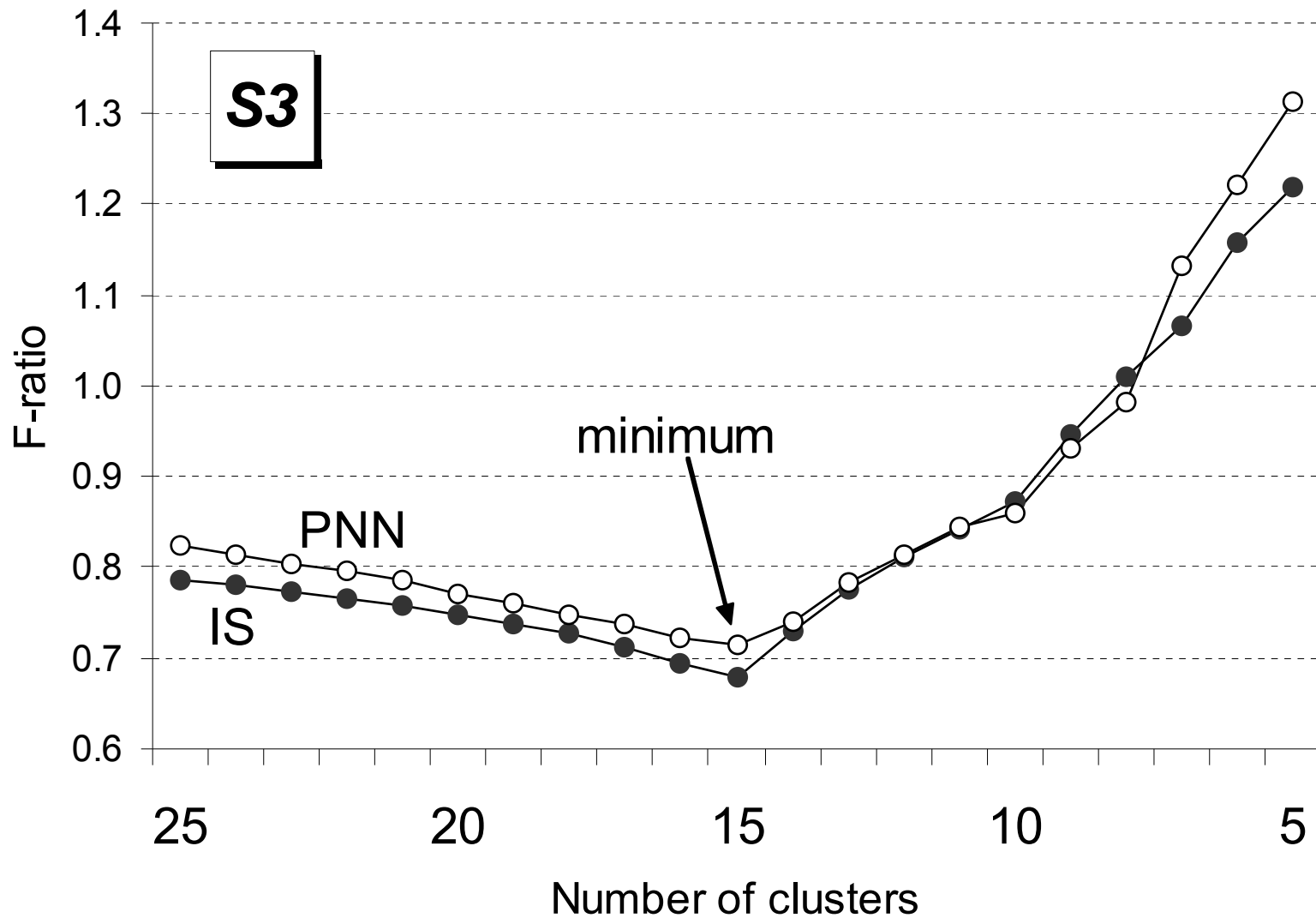
Dataset S1



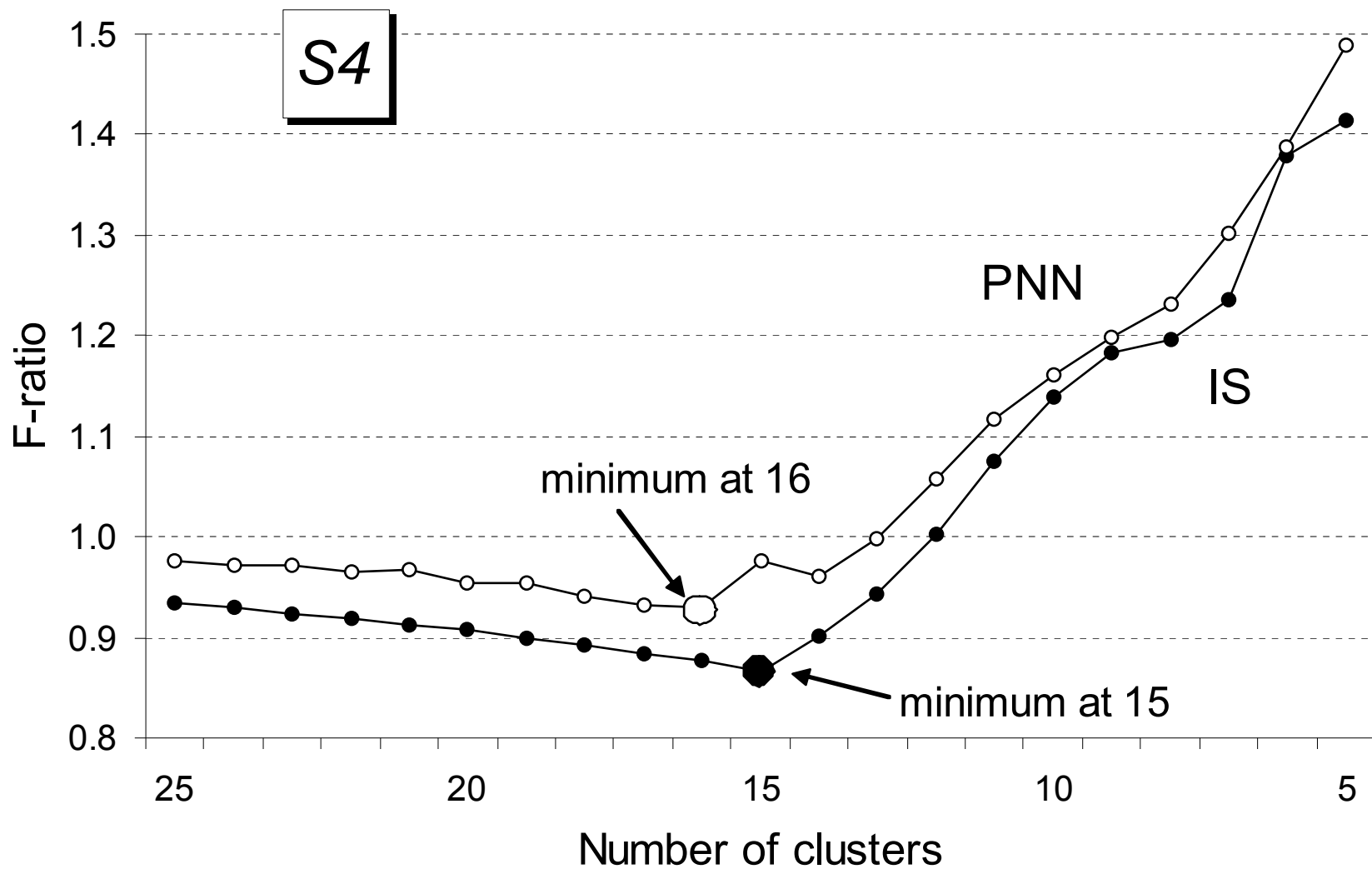
Dataset S2



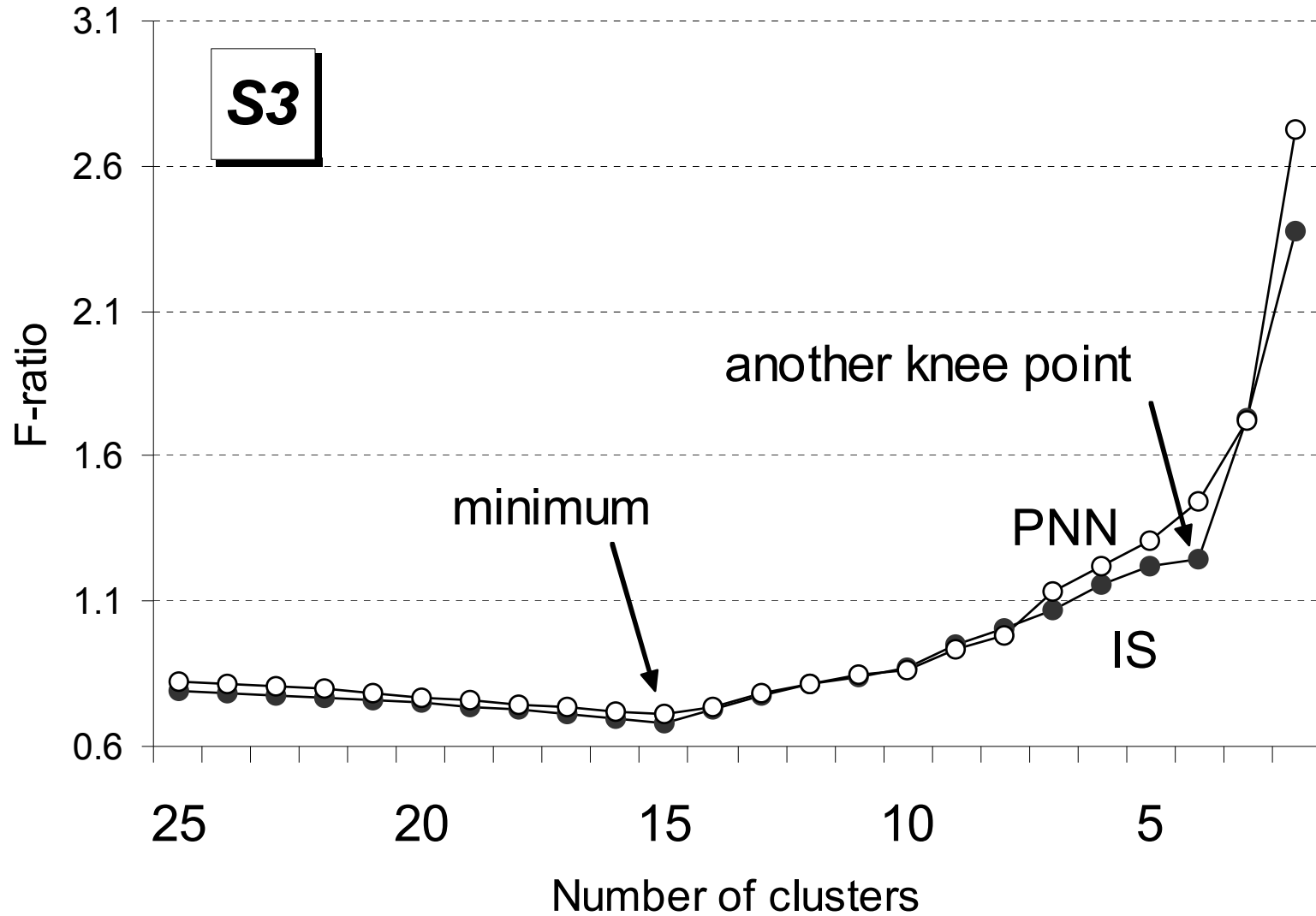
Dataset S3



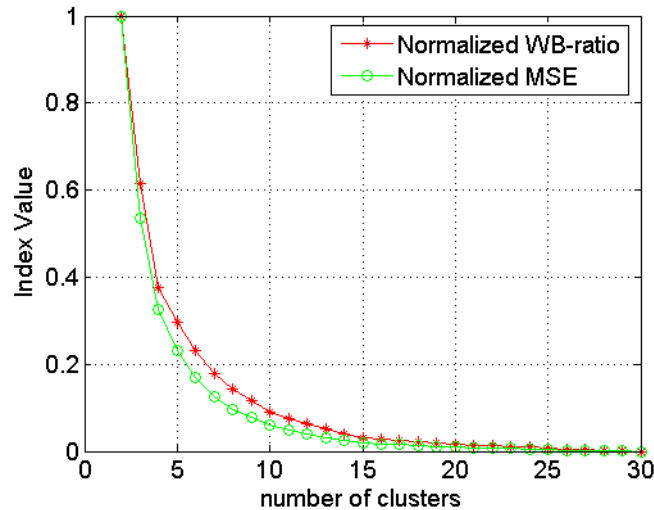
Dataset S4



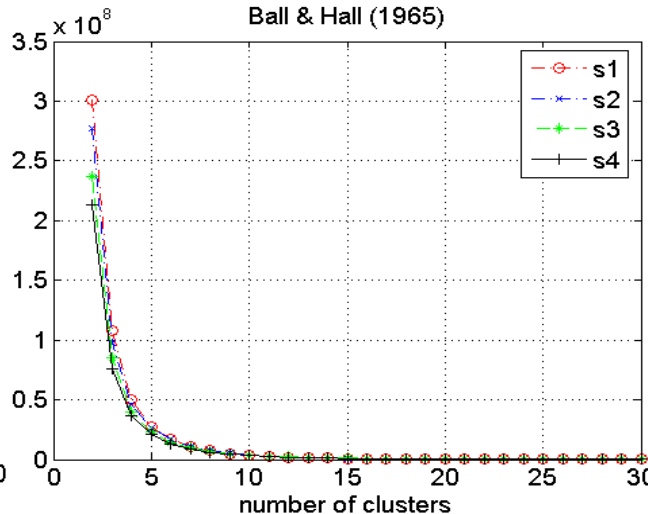
Extension for S3



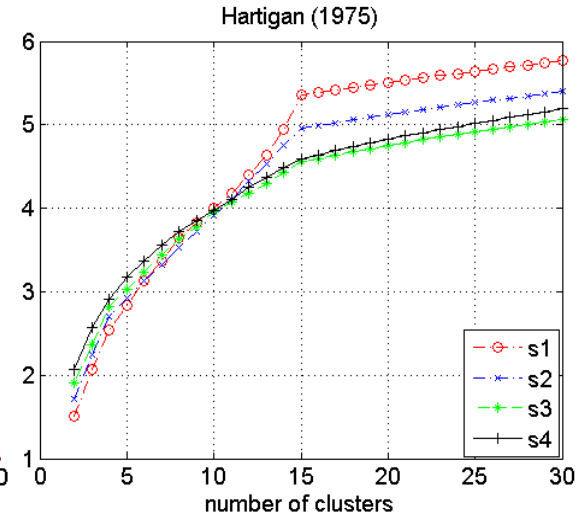
Sum-of-square based index



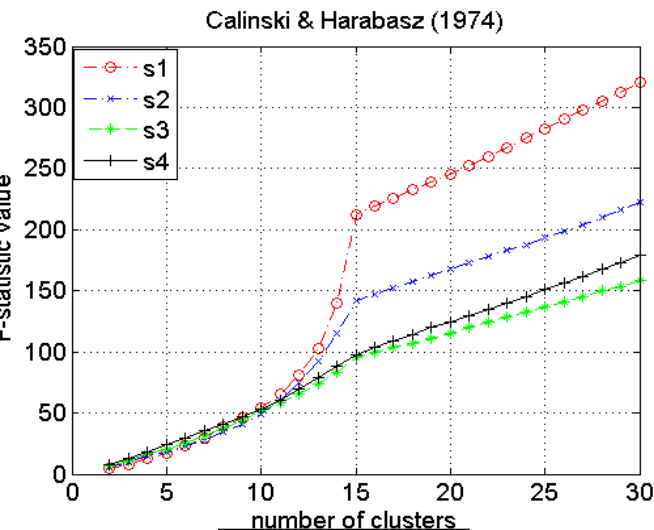
SSW / SSB & MSE



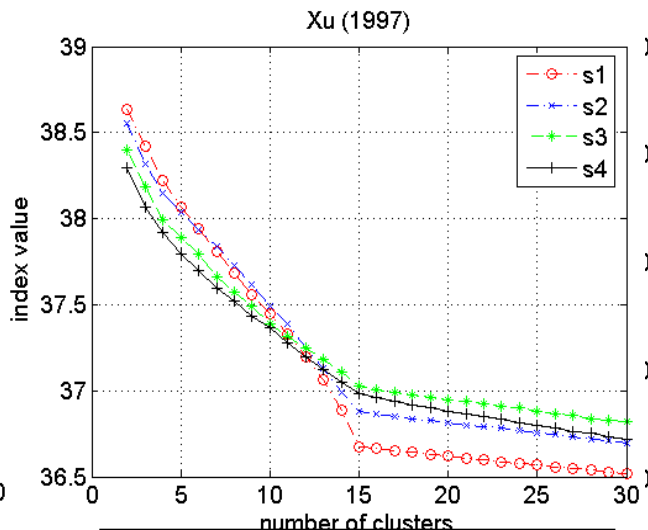
SSW / m



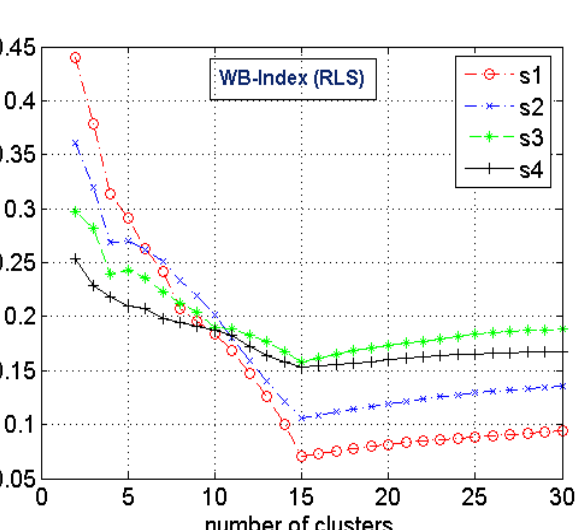
$\log(SSB/SSW)$



$\frac{SSB / m - 1}{SSW / n - m}$



$d \log(\sqrt{SSW / (dn^2)}) + \log(m)$



$m^* SSW/SSB$

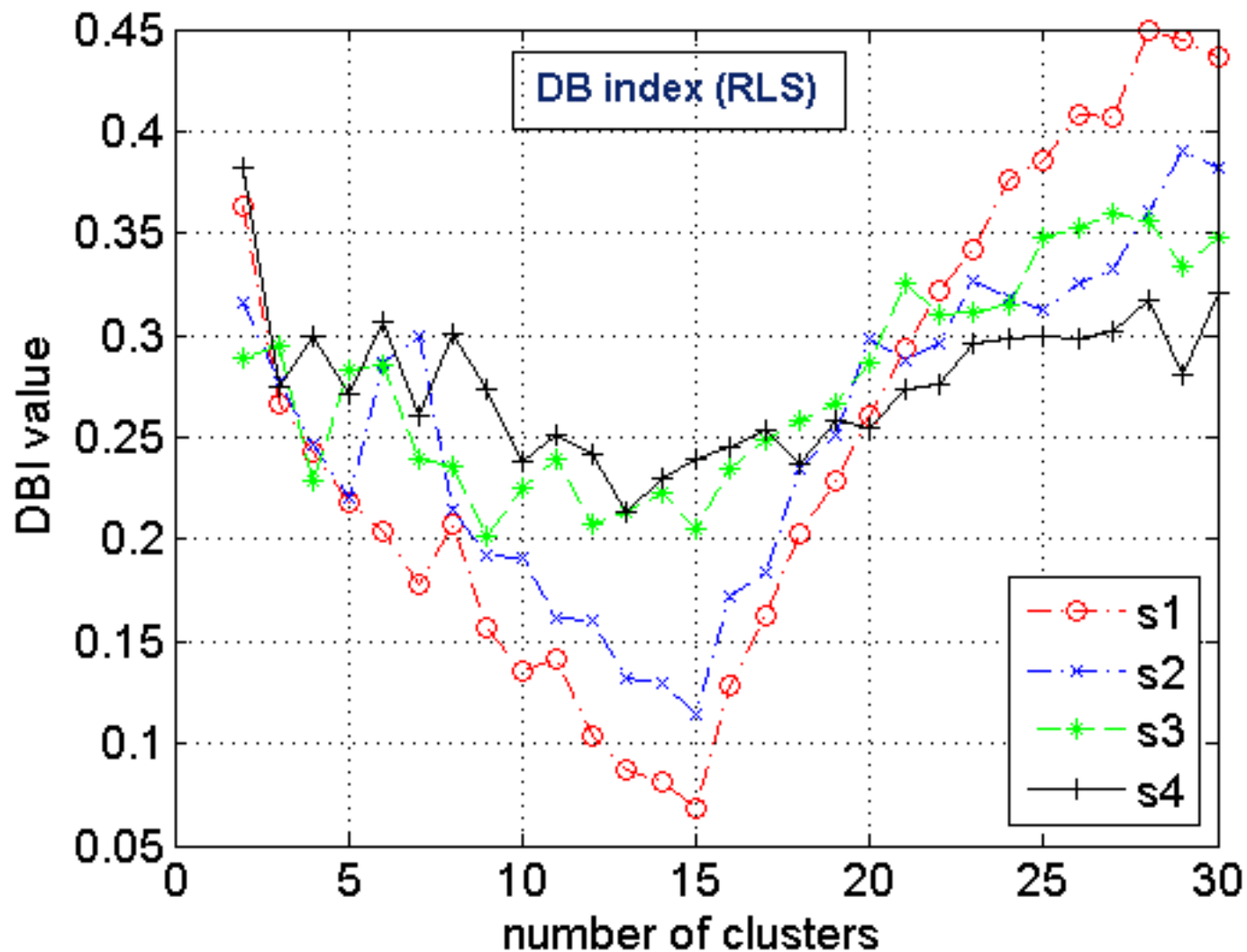
Davies-Bouldin index (DBI)

- Minimize intra cluster variance
- Maximize the distance between clusters
- Cost function weighted sum of the two:

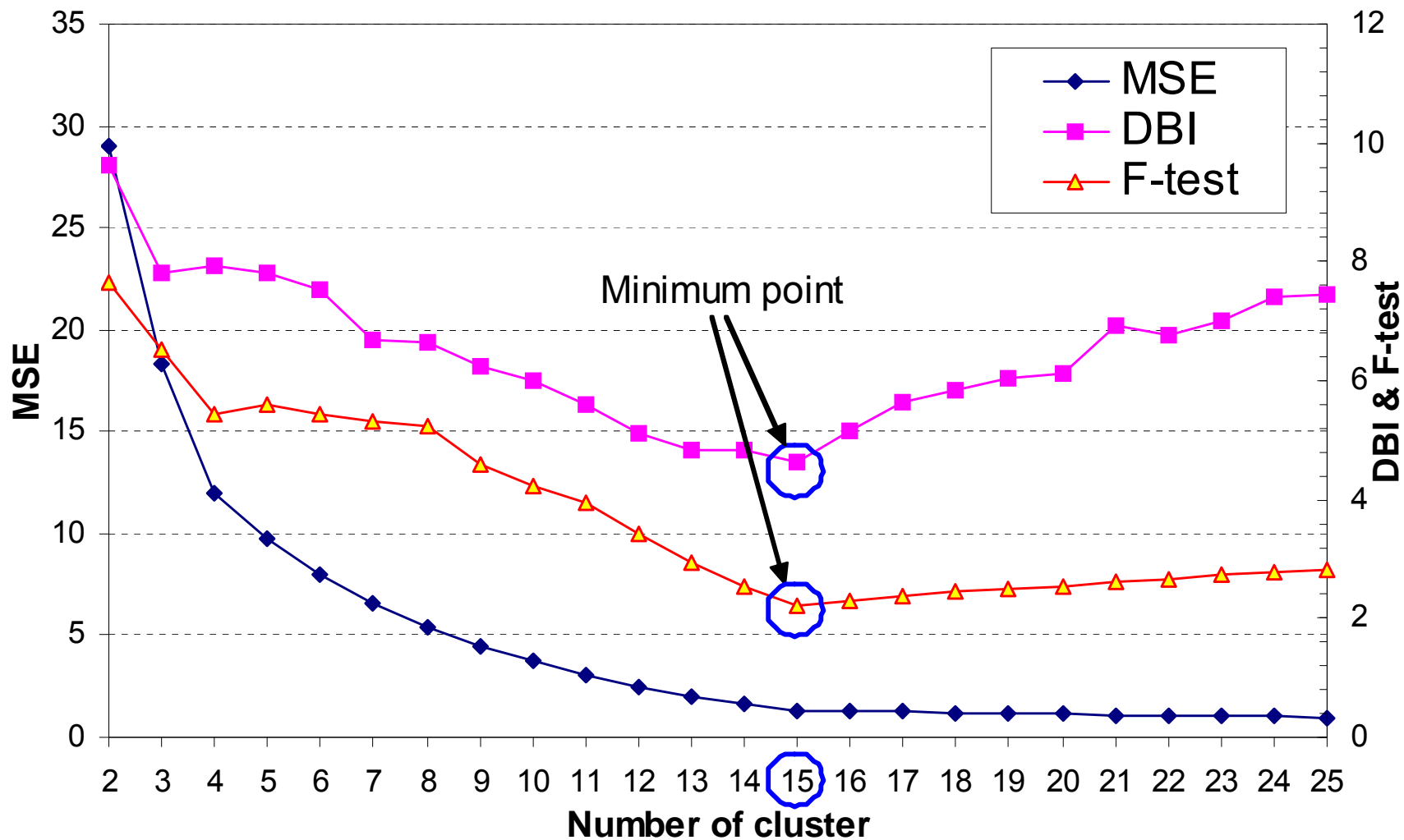
$$R_{j,k} = \frac{MAE_j + MAE_k}{d(c_j, c_k)}$$

$$DBI = \frac{1}{M} \sum_{j=1}^M \max_{j \neq k} R_{j,k}$$

Davies-Bouldin index (DBI)



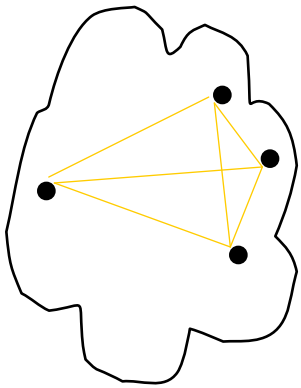
Measured values for S2



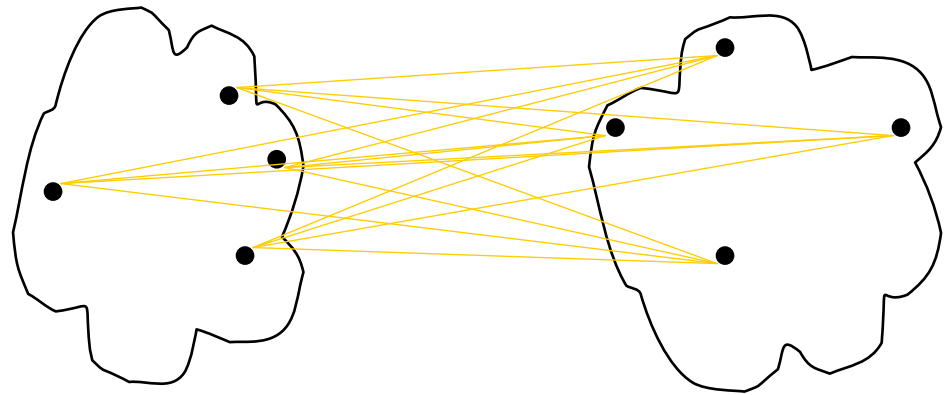
Silhouette coefficient

[Kaufman&Rousseeuw, 1990]

- **Cohesion:** measures how close objects are in a cluster
- **Separation:** measure how separated the clusters are



cohesion



separation

Silhouette coefficient

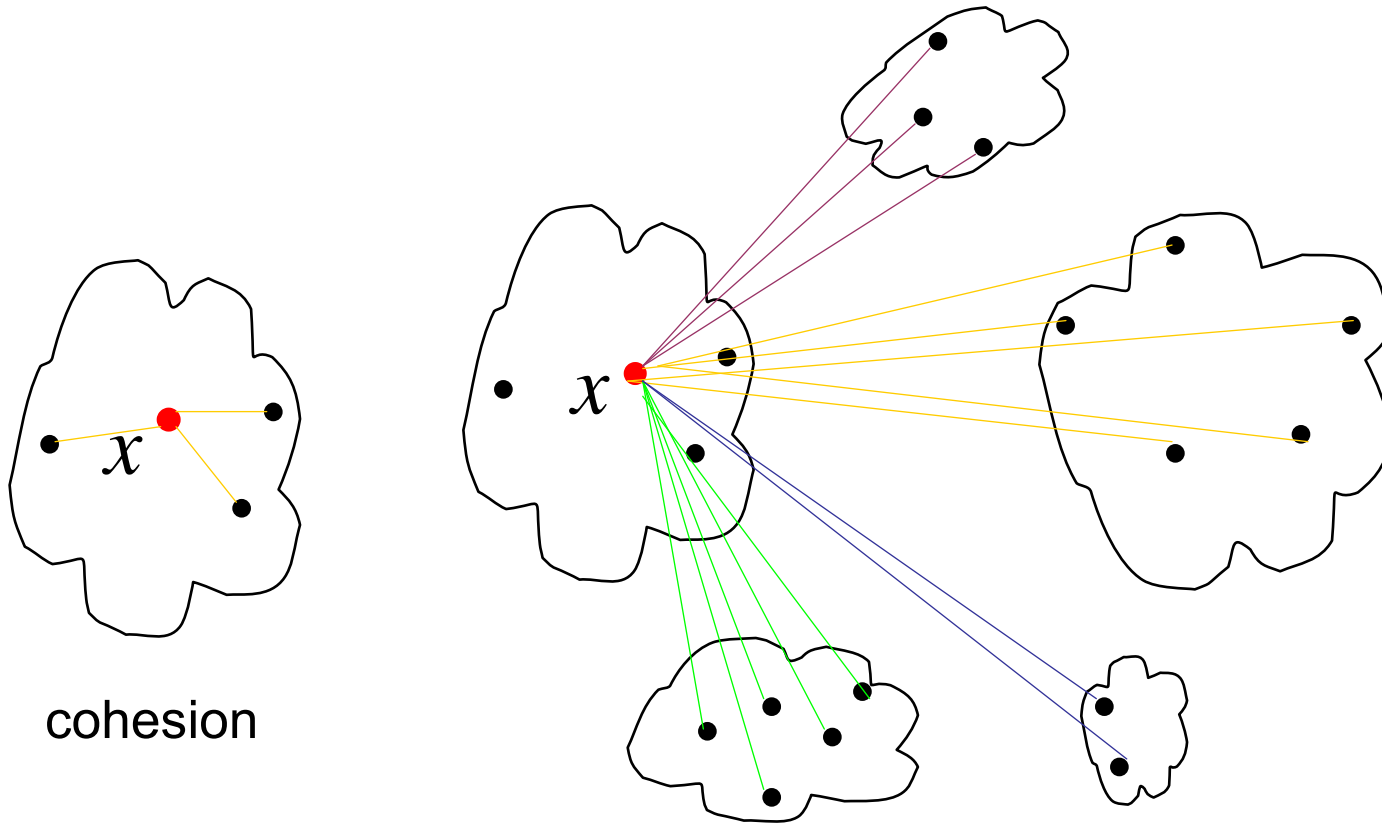
- *Cohesion* $a(x)$: average distance of x to all other vectors in the same cluster.
- *Separation* $b(x)$: average distance of x to the vectors in other clusters. Find the minimum among the clusters.
- *silhouette* $s(x)$:

$$s(x) = \frac{b(x) - a(x)}{\max\{a(x), b(x)\}}$$

- $s(x) = [-1, +1]$: -1=bad, 0=indifferent, 1=good
- Silhouette coefficient (SC):

$$SC = \frac{1}{N} \sum_{i=1}^N s(x)$$

Silhouette coefficient (SC)



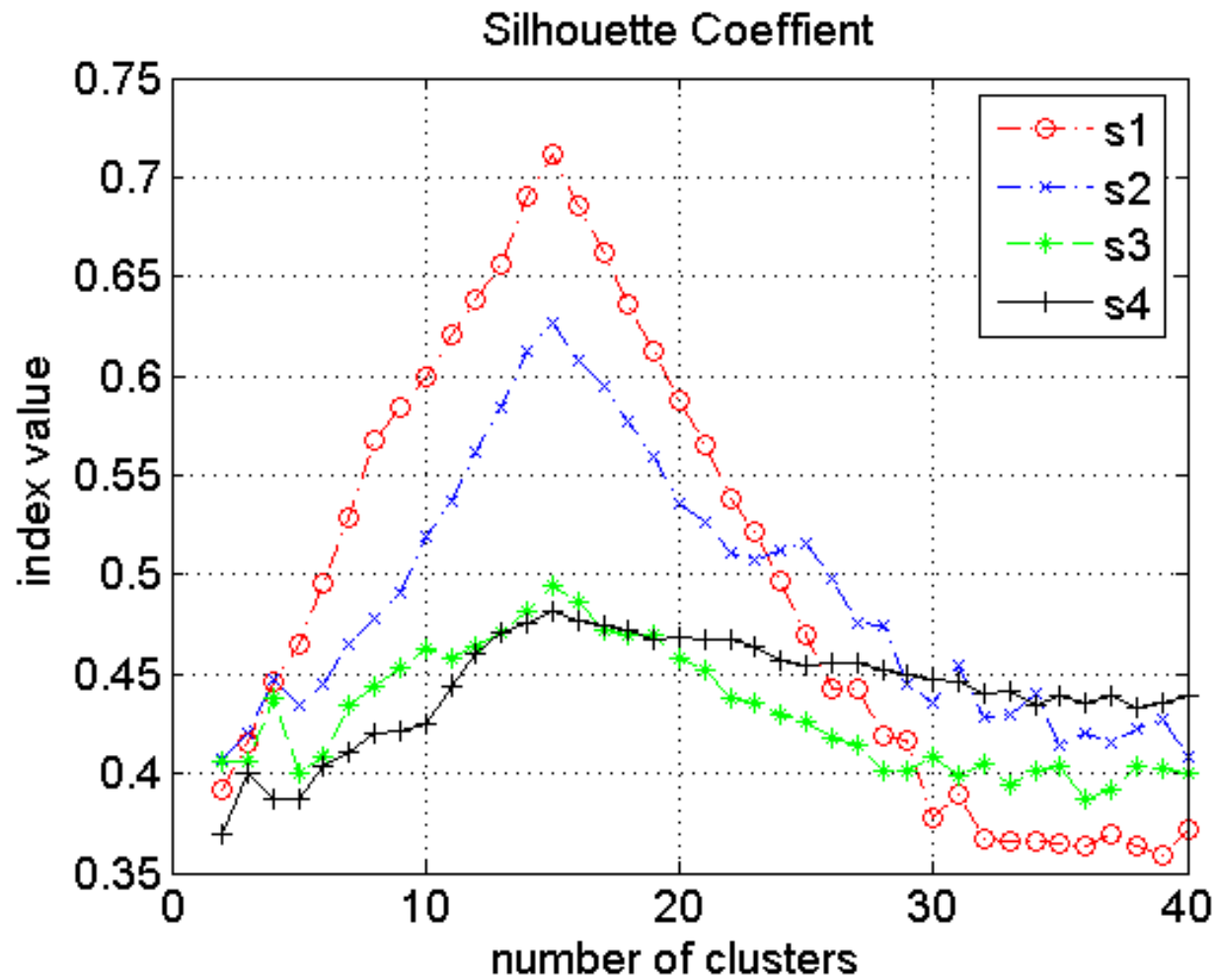
cohesion

$a(x)$: average distance
in the cluster

separation

$b(x)$: average distances to
others clusters, find minimal

Performance of SC



Bayesian information criterion (BIC)

Formula for GMM

$$BIC = L(\theta) - \frac{1}{2} m \log n$$

$L(\theta)$ -- log-likelihood function of all models;

n -- size of data set;

m -- number of clusters

Under spherical Gaussian assumption, we get :

Formula of BIC in partitioning-based clustering

$$BIC = \sum_{i=1}^m \left(n_i \log n_i - n_i \log n - \frac{n_i * d}{2} \log(2\pi) - \frac{n_i}{2} \log \Sigma_i - \frac{n_i - m}{2} \right) - \frac{1}{2} m \log n$$

d -- dimension of the data set

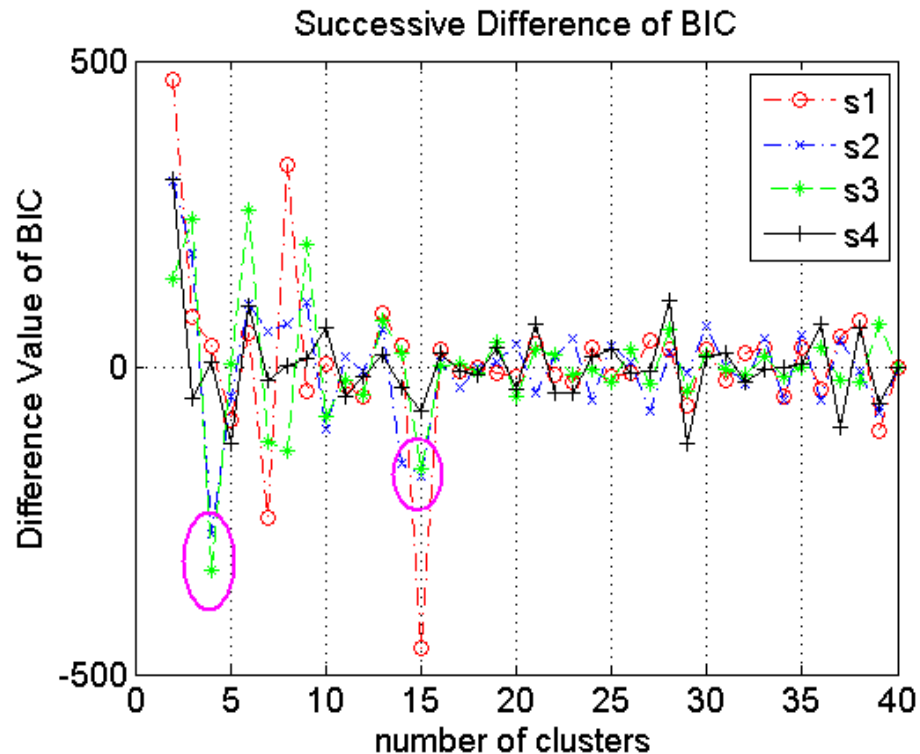
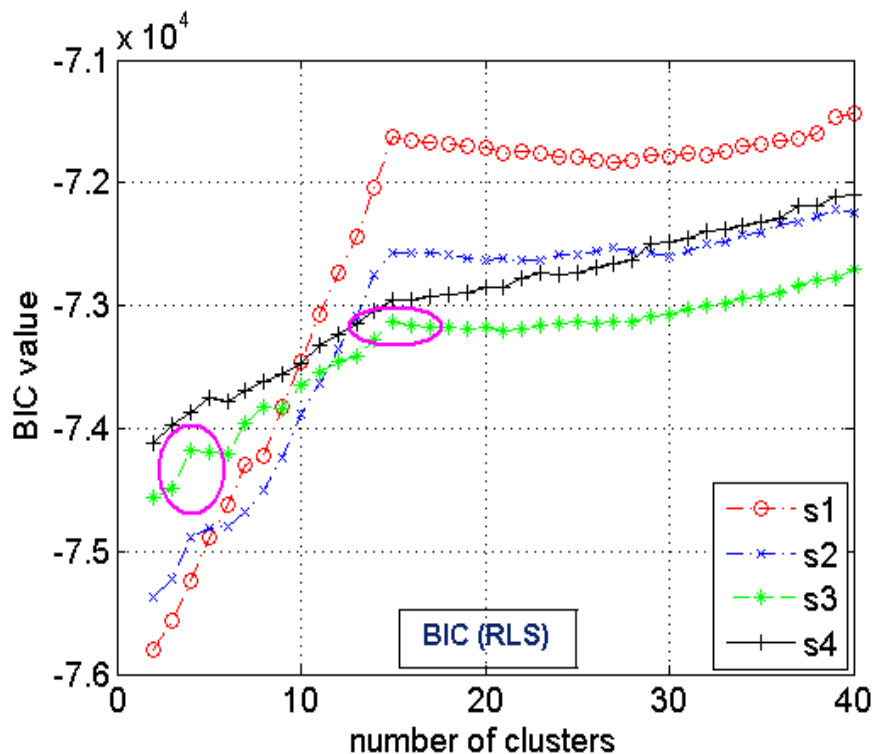
n_i -- size of the i^{th} cluster

Σ_i -- covariance of i^{th} cluster

Knee Point Detection on BIC

Original BIC = $F(m)$

$SD(m) = F(m-1) + F(m+1) - 2 \cdot F(m)$



Internal indexes

Table B.1: Formulas for internal indexes

Name	Formula
SSW	$SSW = \frac{1}{N} \sum_{i=1}^N \ x_i - C_{p_i}\ ^2$
SSB	$SSB = \frac{2}{M(M-1)} \sum_{i=1}^M \sum_{j=1, j \neq i}^M \ C_i - C_j\ ^2$
Calinski-Harabasz index	$CH = \frac{SSB/(M-1)}{SSW/(N-M)}$
Hartigan	$H_M = \left(\frac{SSW_M}{SSW_{M+1}} - 1 \right) (N - M - 1)$ or : $H_M = \log(SSB_M / SSW_M)$
Krzanowski-Lai index	$diff_M = (M-1)^{2/D} SSW_{M-1} - M^{2/D} SSW_M$ $KL_M = diff_M / diff_{M+1} $
Ball&Hall	$BH_M = SSW_M / M$
Xu-index	$Xu = D \log(\sqrt{SSW_M / (DN^2)}) + \log M$
Dunn's index	$Dunn = \sum_{i=1}^M \frac{\max(\ x_j - C_i\ ^2)_{j \in C_i}}{d_{ij}}$
Davies&Bouldin index	$R_{ij} = \frac{S_i + S_j}{d_{ij}}, i \neq j$ where : $d_{ij} = \ C_i - C_j\ ^2, S_i = \frac{1}{n_i} \sum_{j=1}^{n_i} \ x_j - C_i\ ^2$ and, $R_i = \max_{j=1, \dots, M} R_{ij}, i = 1, \dots, M$ $DBI = \frac{1}{M} \sum_{i=1}^M R_i$

Internal indexes

Silhouette Coefficients	$a(x_i) = \frac{1}{n_m - 1} \sum_{j=1, j \neq i}^{n_m} \ x_i - x_j\ _{x_i, x_j \in C_m}^2$ $b(x_i) = \min_t \left\{ \frac{1}{n_t} \sum_{j \in C_t} \ x_i - x_j\ ^2 \right\}_{x_i \notin C_t}$ $s(x_i) = \frac{b(x_i) - a(x_i)}{\max(a(x_i), b(x_i))}$ $SC = \frac{1}{N} \sum_{i=1}^N s(x_i)$ $b(x_i) = \min_{t \neq m} \left\{ \sum_{j \in C_t} \ C_t - C_m\ ^2 \right\}_{x_i \notin C_t} (SC'2008)$
RMSSTD	$RMSSTD = \frac{\sum_{d=1, \dots, D} \sum_{k=1, \dots, M}^{n_{kd}} (x_i - \bar{x}^d)^2}{\sum_{d=1, \dots, D} (n_{kd} - 1)}$
R-square	$RS = \frac{SST - SSW}{SST} = \frac{\sum_{d=1, \dots, D} \sum_{i=1}^{n_d} (x_i - \bar{x}^d)^2 - \sum_{k=1, \dots, M} \sum_{i=1}^{n_{kd}} (x_i - \bar{x}^d)^2}{\sum_{d=1, \dots, D} \sum_{i=1}^{n_d} (x_i - \bar{x}^d)^2}$
Bayesian Information Criterion	$BIC = L * N - \frac{1}{2} M(D + 1) \sum_{i=1}^M \log(n_i)$
Xie-Beni	$XB = \frac{\sum_{i=1}^N \sum_{k=1}^M u_{ik}^2 \ x_i - C_k\ ^2}{N \min_{t/s} \{ \ C_t - C_s\ ^2 \}}$
Partition Coefficient	$PC = \sum_{i=1}^N \sum_{k=1}^M u_{ik}^2 / N$
Partition Entropy	$PE = - \left(\sum_{i=1}^N \sum_{k=1}^M u_{ik} \log(u_{ik}) \right) / N$

Soft partitions

Comparison of the indexes K-means

Data Set																
ValidityCri	R15	A7	s1	s2	s3	s4	D31	birch1	Iris	wine	Control	Image	wdbc	yeast	bridge	zernike
Ball&Hall (knee min)	15	25	30	57	58	63	53	108	10	11	21	40	22	37	62	38
Calinski-Harabsz (max)	24	4	15	20	6	16(2)	39	198	2	2	2	2	2	2	2	2
Hartigan (diff knee min)	15	25	30	57	58	49	53	108	10	10	20	40	21	28	62	38
Wb-index (min)	24	28	15	20	19	20	39	198	2	2	2	3	2	3	2	2
Krzanowski-Lai index (mi)	18	19	15	12	63	50	25	86	9	2	2	23	2	29	33	39
Xu-index (min)	24	28	15	20	17	17	39	117	12	13	24	48	23	38	64	44
Dunn (max)	5	23	4	15	10	15	53		12	2	19	3	3	2	38	36
DBI (min)	24	4	15	13	4	13	23	93	2	2	2	2	2	23	2	2
SC (max)	17	3	15	17	13	17	34	103	2	2	2	2	2	2	2	6
SCI * (max)	NA	4	15	17	17	17	NA	95	2	2	2	2	2	2	NA	NA
XieBeni (min)	3	4	15	4	4	4	4	4	2	2	2	2	2	2	2	5
ORI_IBIC (first max)	NA	4	NA	20	4	5	NA(4)	26	NA(4)	NA	2	NA(3)	NA	NA	NA	NA
Angle-based	NA	4	8	15	4	3	3	199	8	NA	2	20	NA	NA	NA	NA
DiffBIC	NA	4	11	15	15	5	NA	4	NA	NA	NA	NA	NA	NA	NA	NA
# of clusters	15	7	15	15	15	15	31	100	3	3	6	7	2	10	NA	10

Comparison of the indexes

Random Swap

Data Set	Random Swap															
	R15	A7	s1	s2	s3	s4	D31	birch1	Iris	wine	Control	Image	wdbc	yeast	bridge	zernike
ValidityCri																
Ball&Hall (knee)	24	26	69	68	66	68	51	99	9	11	21	45	18	8	62	42
Calinski-Harabsz (max)	15	25	15	15	2	15	32	199	2	2	2	2	2	4	2	2
Hartigan (max knee)	23	26	69	68	66	68	51	99	9	11	21	40	18	8	62	36
Wb-index (min)	15	25	15	15	15	15	32	199	2	2	2	2	2	4	2	4
Krzanowski-Lai index (mi)	15	14	15	15	15	15	16	84	10	2	9	20	2	19	35	2or36
Xu-index (min)	15	25	15	15	15	15	32	102	12	13	24	48	23	38	64	44
Dunn (max)	8	25	15	15	63	20	31		2	2	19	33	21	36	51	26
DBI (min)	8	4	15	16	15	15	30	99	2	2	2	2	2	5	2	2
SC (max)	15	4	15	15	15	15	30	102	2	2	2	2	2	4	2	6
SCI * (max)	8	4	15	15	15	15	31	102	2	2	2	2	2	8	2	21
XieBeni (min)	15	3	15	15	4or15	15	30	99	2	10	6	2	2	4	2	25
ORI_IBIC (max knee)	8	4	15	15	15	15	31	4	5	2	NA	2	2	2	2	2
Angle-based	8	4	15	4	4	5	11	4	11	12	14	23	10	25	47	36
DiffBIC	8	4	14	15	15	5	12	4	2	2	NA	1	1	1	1	1
# of clusters	15	7	15	15	15	15	31	100	3	3	6	7	2	10	NA	10

Part III:

**Stochastic complexity for
binary data**

Stochastic complexity

- Principle of minimum description length (MDL): find clustering C that can be used for describing the data with minimum information.
- Data = Clustering + description of data.
- Clustering defined by the centroids.
- Data defined by:
 - which cluster (partition index)
 - where in cluster (difference from centroid)

Solution for binary data

$$SC = \sum_{j=1}^M n_j \sum_{i=1}^d h\left(\frac{n_{ij}}{n_j}\right) + \sum_{j=1}^M -n_j \log\left(\frac{n_j}{N}\right) + \frac{d}{2} \sum_{j=1}^M \log \max(1, n_j)$$

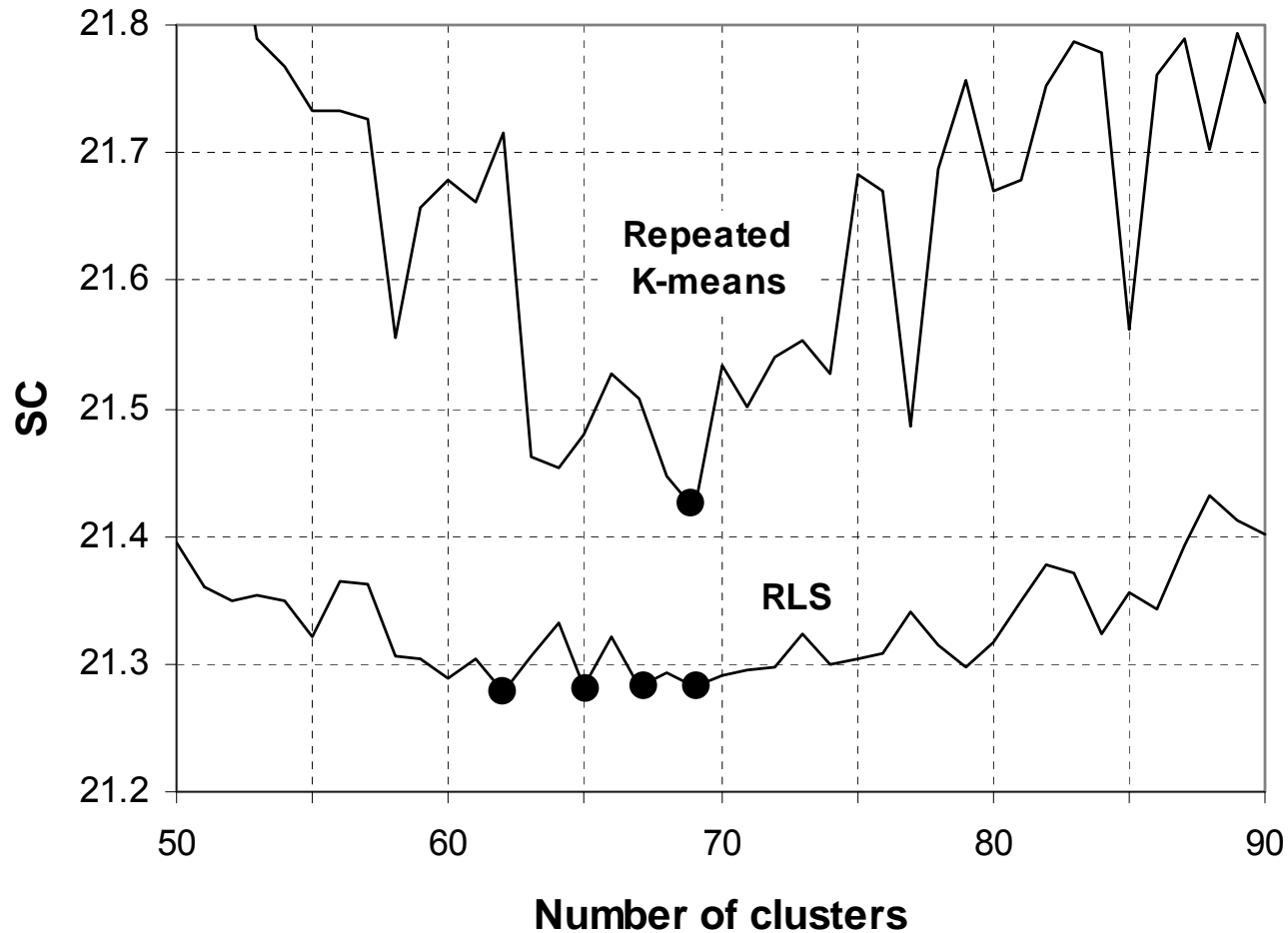
where

$$h(p) = -p \log(p) - (1-p) \log(1-p)$$

This can be simplified to:

$$SC \approx \sum_{j=1}^M n_j \sum_{i=1}^d h\left(\frac{n_{ij}}{n_j}\right) + \sum_{j=1}^M -n_j \log n_j + \frac{d}{2} \sum_{j=1}^M \log \max(1, n_j)$$

Number of clusters by stochastic complexity (SC)

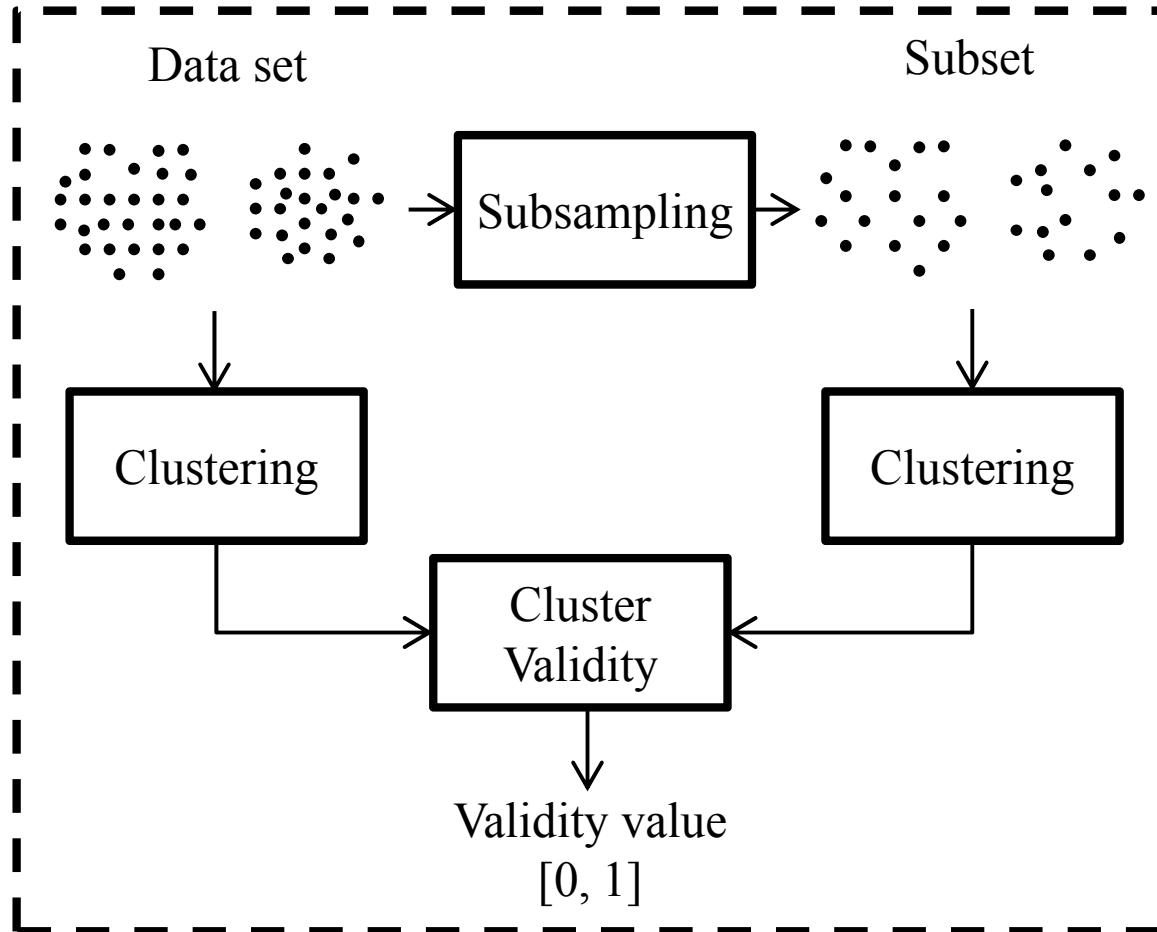


Part IV:

Stability-based approach

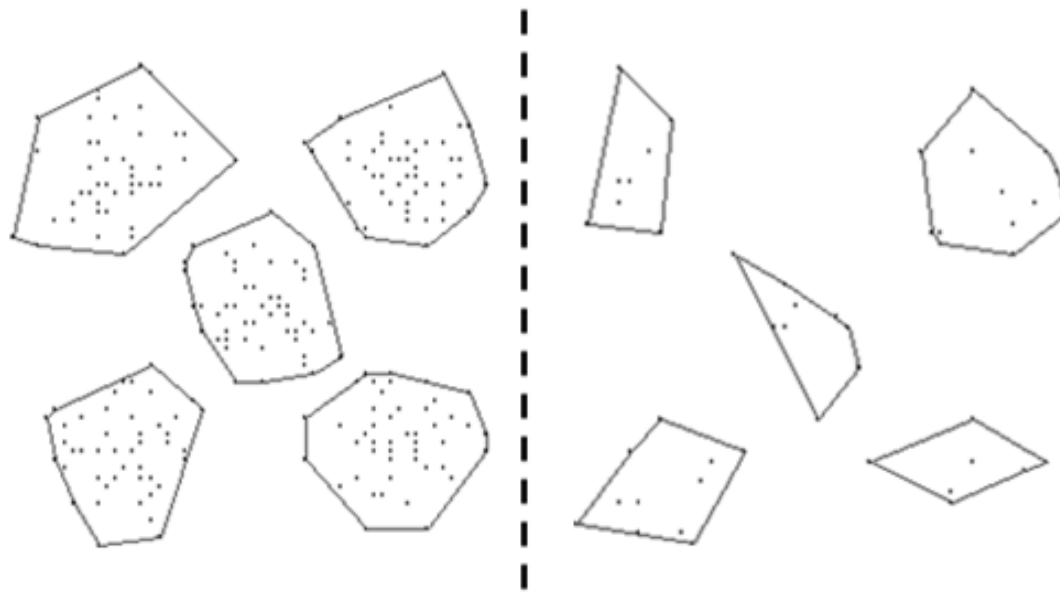
Cross-validation

Compare clustering of full data against sub-sample



Cross-validation: Correct

Correct number of clusters: $k=5$

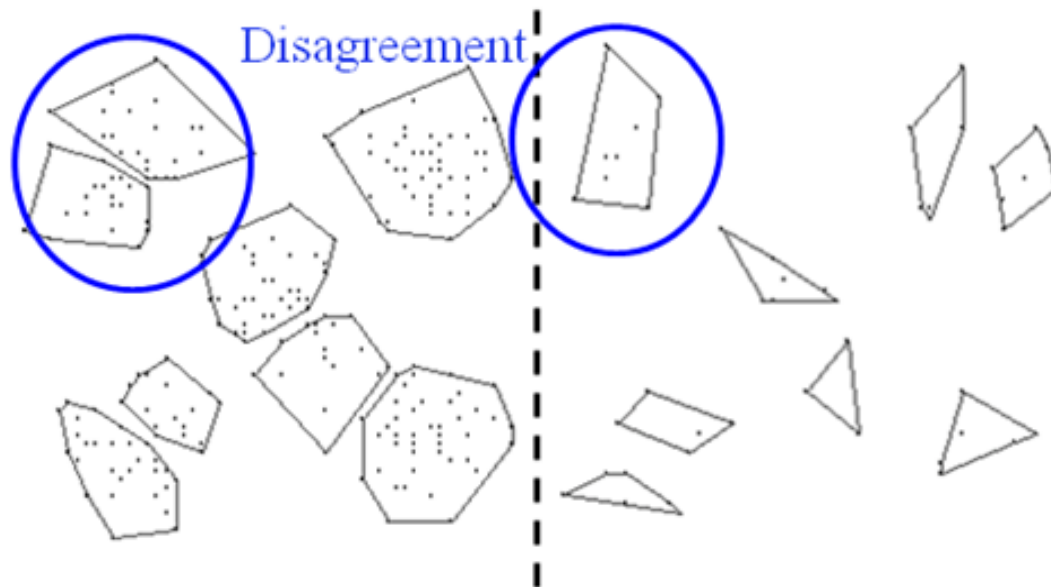


Same results

Cross-validation

Incorrect

Incorrect number of clusters: $k=8$



Different results

Stability approach in general

1. Add randomness
2. Cross-validation strategy
3. Solve the clustering
4. Compare clustering

Adding randomness

- Three choices:
 1. Subsample
 2. Add noise
 3. Randomize the algorithm
- What subsample size?
- How to model noise and how much?
- Use k-means?

Sub-sample size

- Too large (80%): same clustering always
- Too small (5%): may break cluster structure
- Recommended 20-40%

Spiral dataset



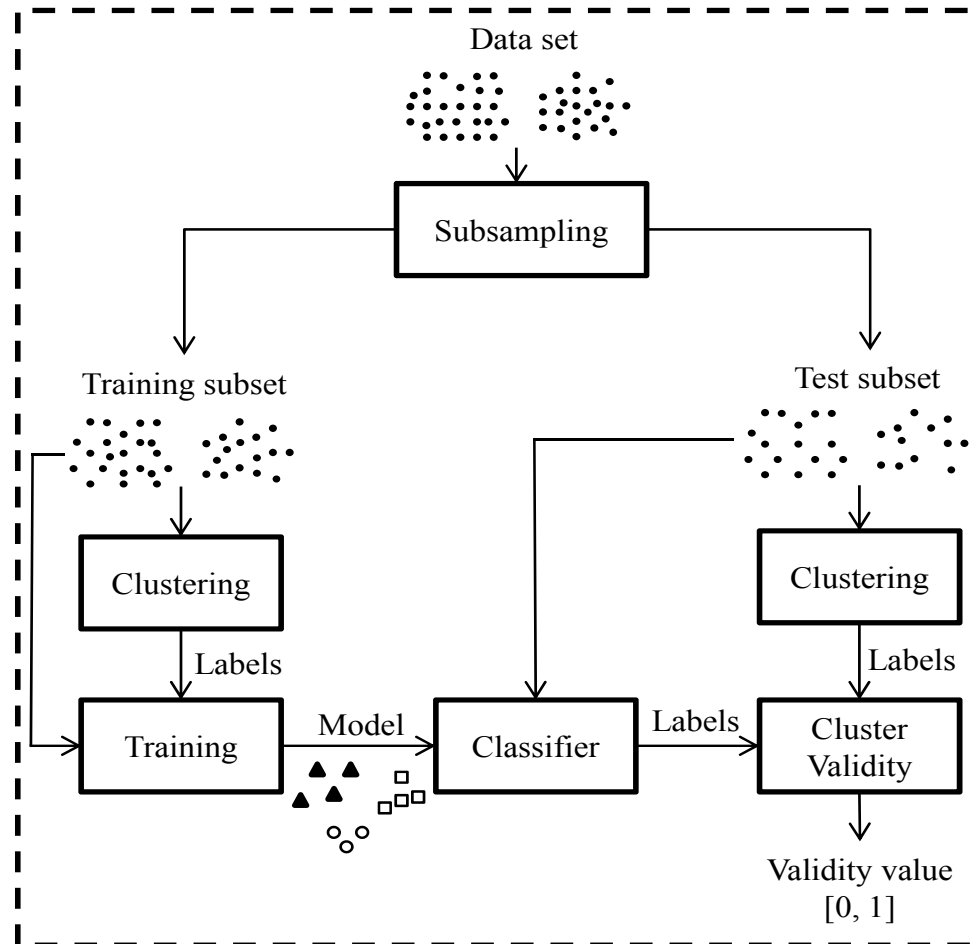
60% subsample



20% subsample



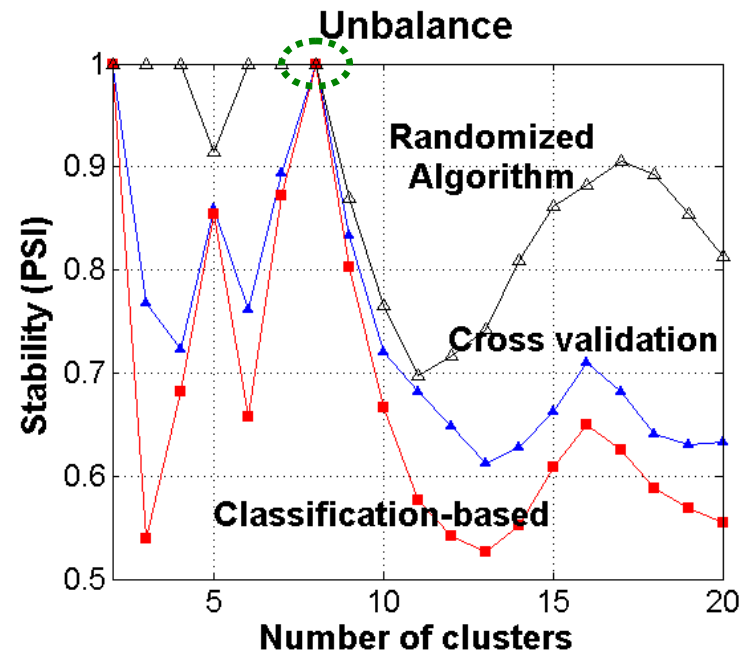
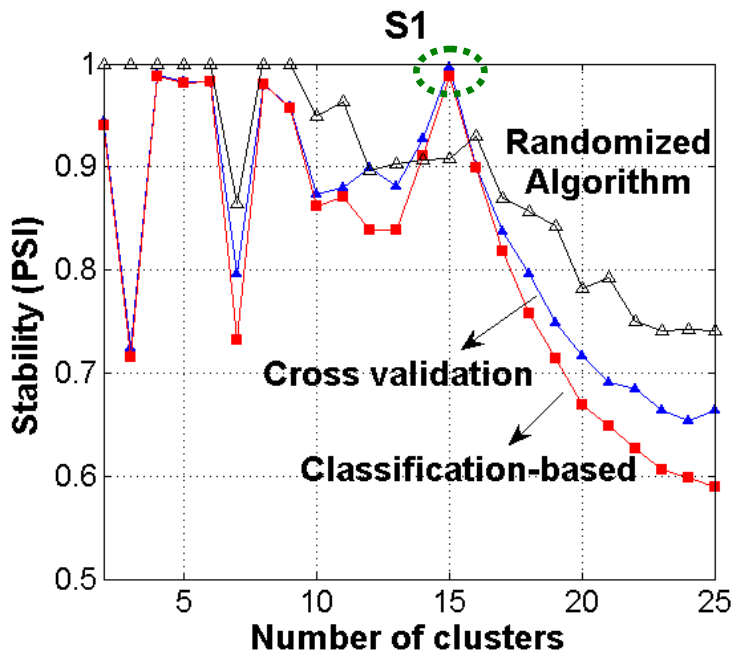
Classification approach



Does not really add anything more.
Just makes process more complex.

Comparison of three approaches

- Cross-validation works ok
- Classification also ok
- Randomizing algorithm fails



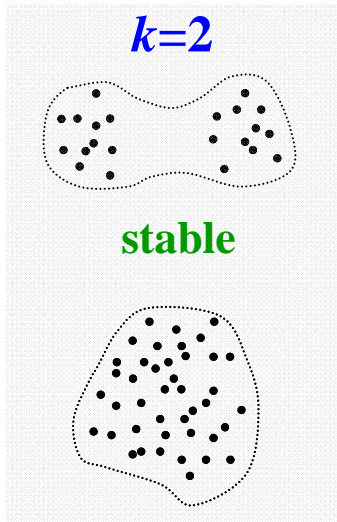
Problem

Stability can also come from other reasons:

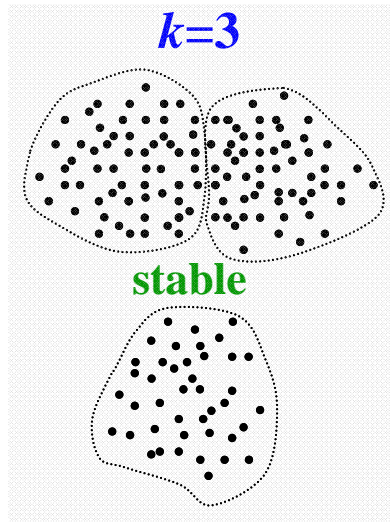
- Different cluster sizes
- Wrong cluster model

Happens when $k < k^*$

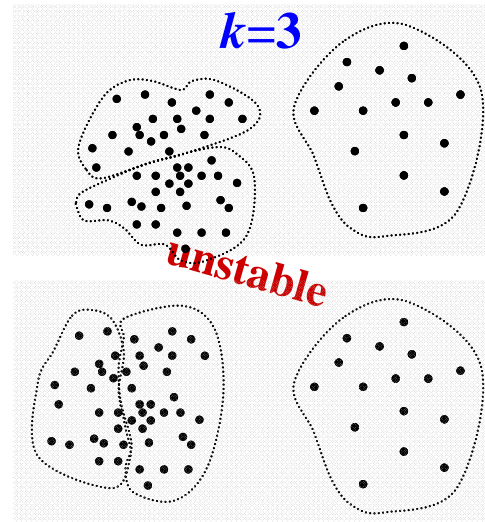
Too few clusters
different size



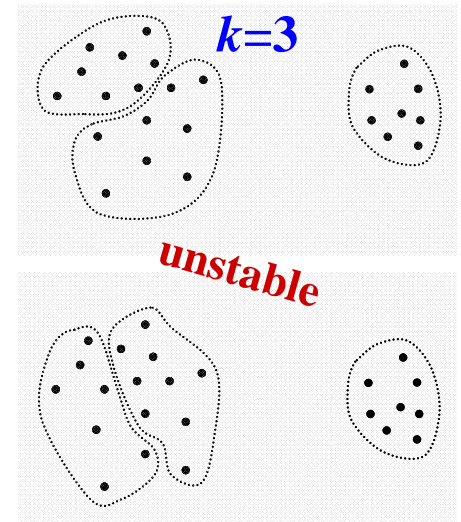
Too many clusters
wrong model



Too many clusters
different density

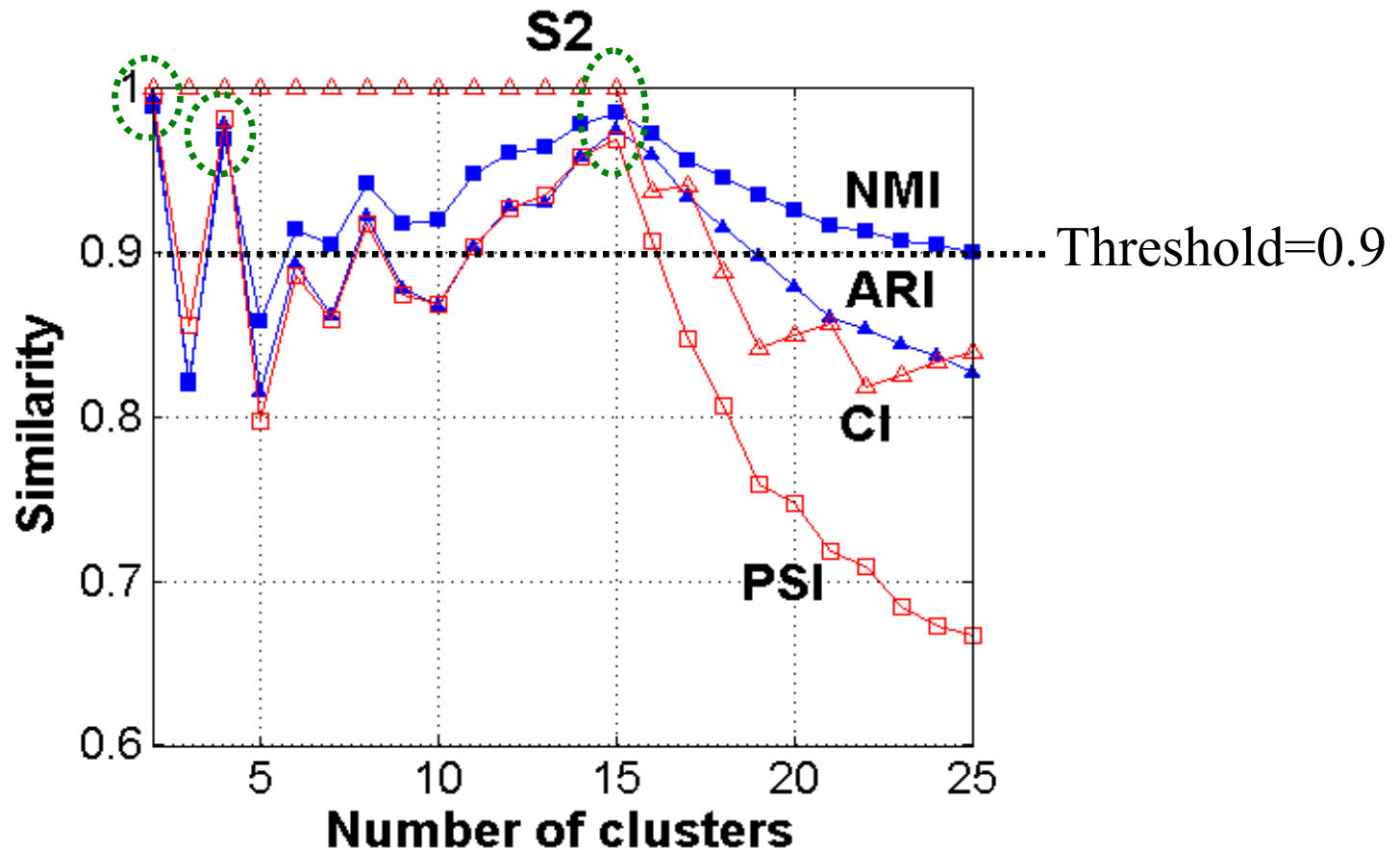


Too many clusters
different size



Solution

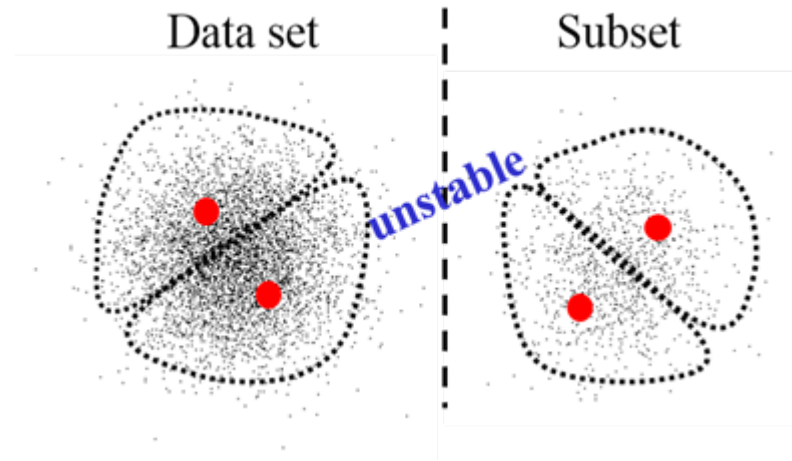
Instead of selecting k with maximum stability, select last k with stable result.



Effect of cluster shapes

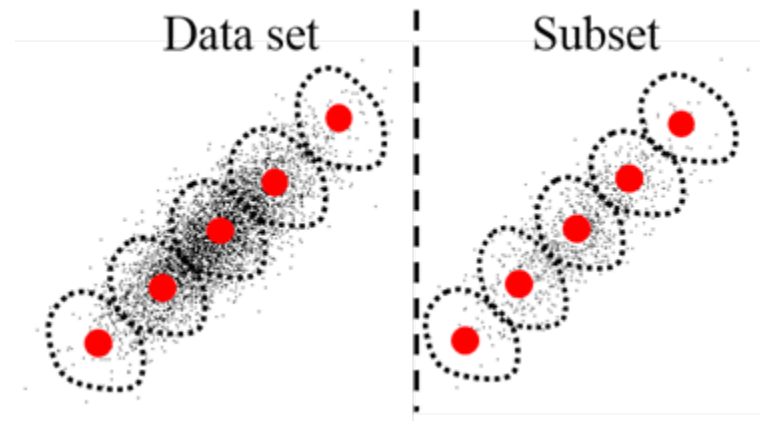
Correct model:

- Works ok.



Wrong model:

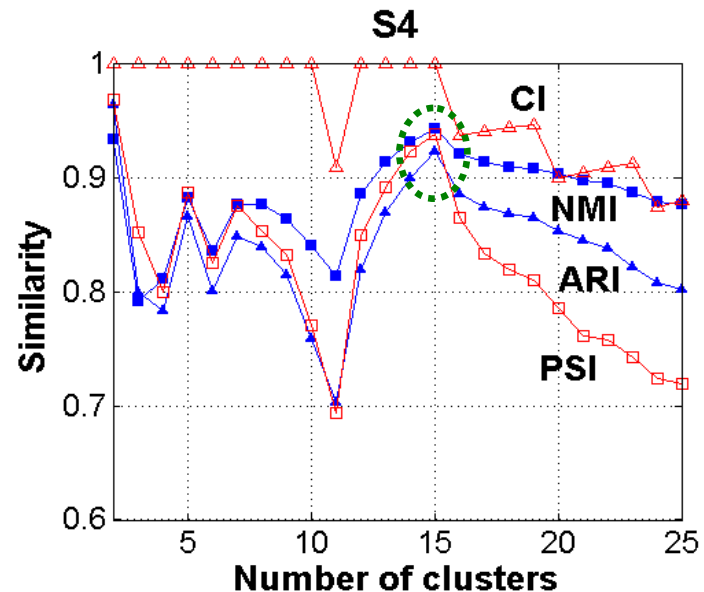
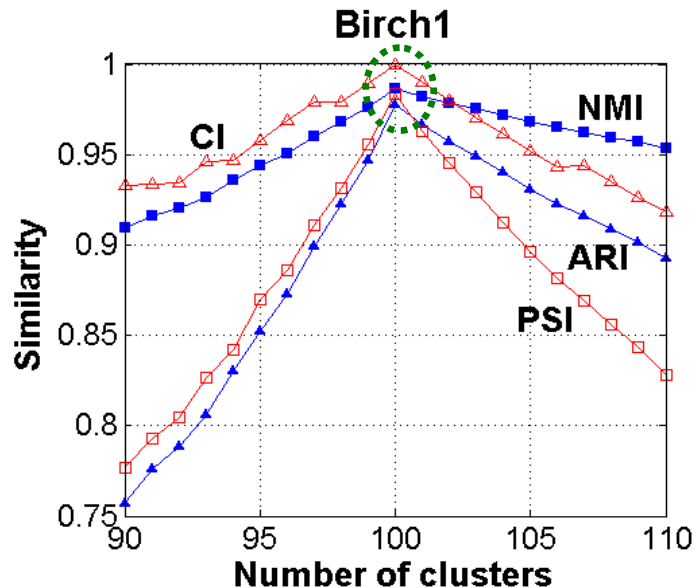
- Elliptical cluster
Minimizing TSE would
find 5 spherical clusters



Which external index?

Does not matter much

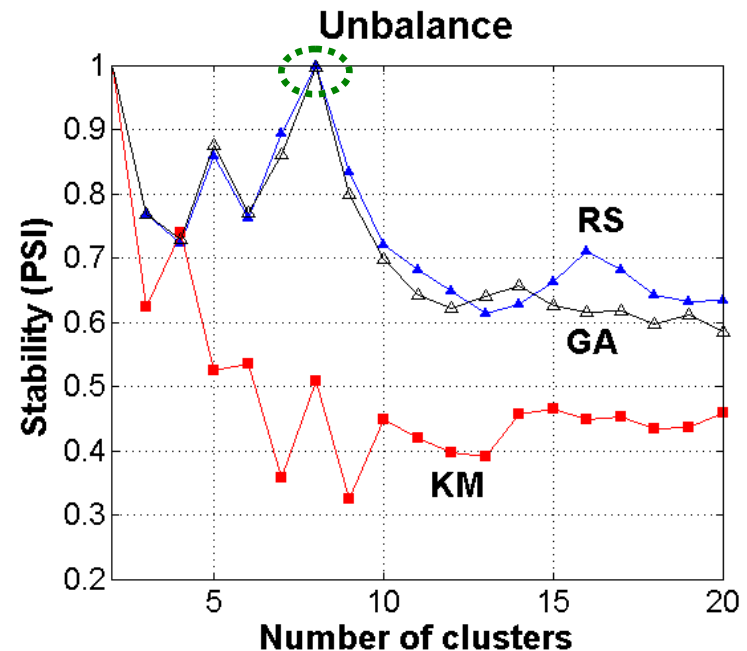
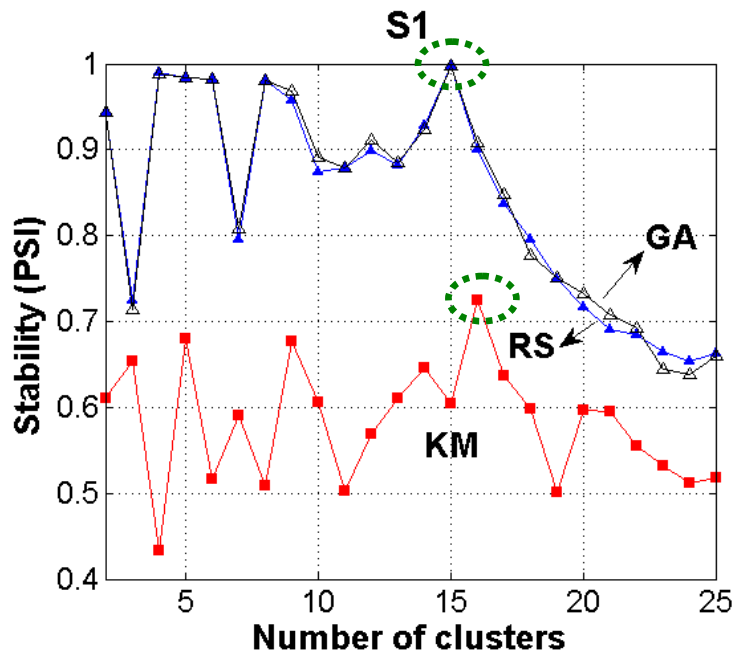
- This is not: RI
- These all ok: ARI, NMI, PSI, NVD, CSI
- CI cares only allocation: sometimes too rough.



Does algorithm matter?

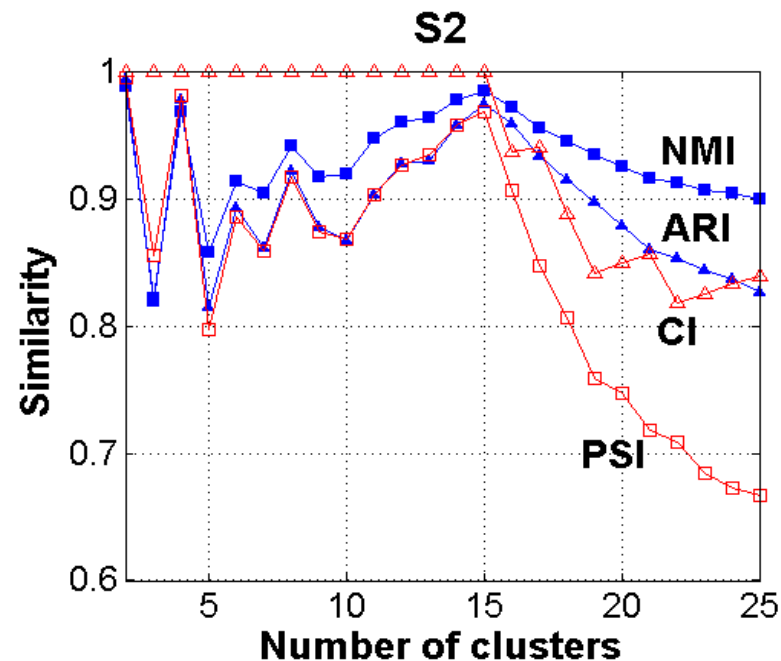
Yes it does.

- Ok: Random Swap (RS) and Genetic Algorithm (GA)
- Not: K-means (KM)



Summary

- The choice of the cross-validation strategy not critical
- Last stable clustering instead of global maximum
- The choice of external index is not critical
- Good clustering algorithm required (RS or GA)



Part V:

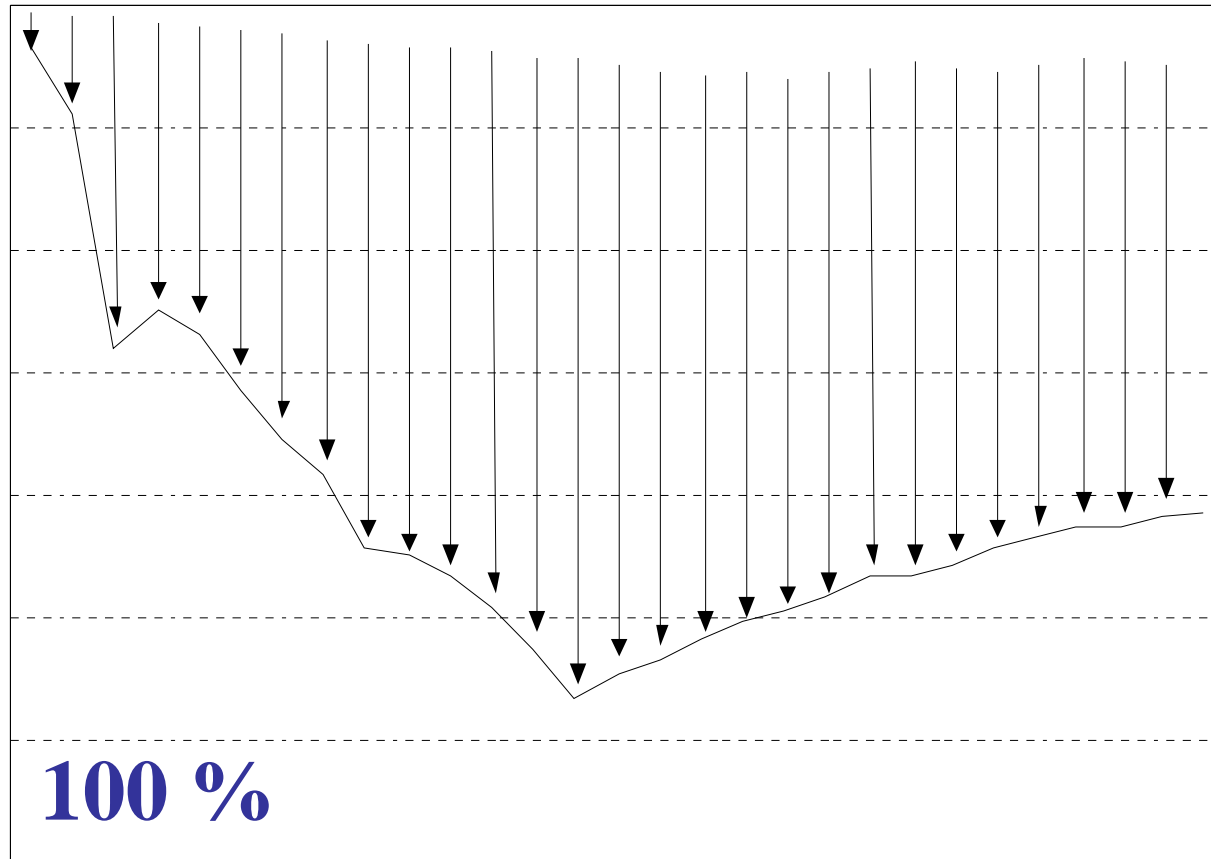
Efficient implementation

Strategies for efficient search

- **Brute force:** solve clustering for all possible number of clusters.
- **Stepwise:** as in brute force but start using previous solution and iterate less.
- **Criterion-guided search:** Integrate cost function directly into the optimization function.

Brute force search strategy

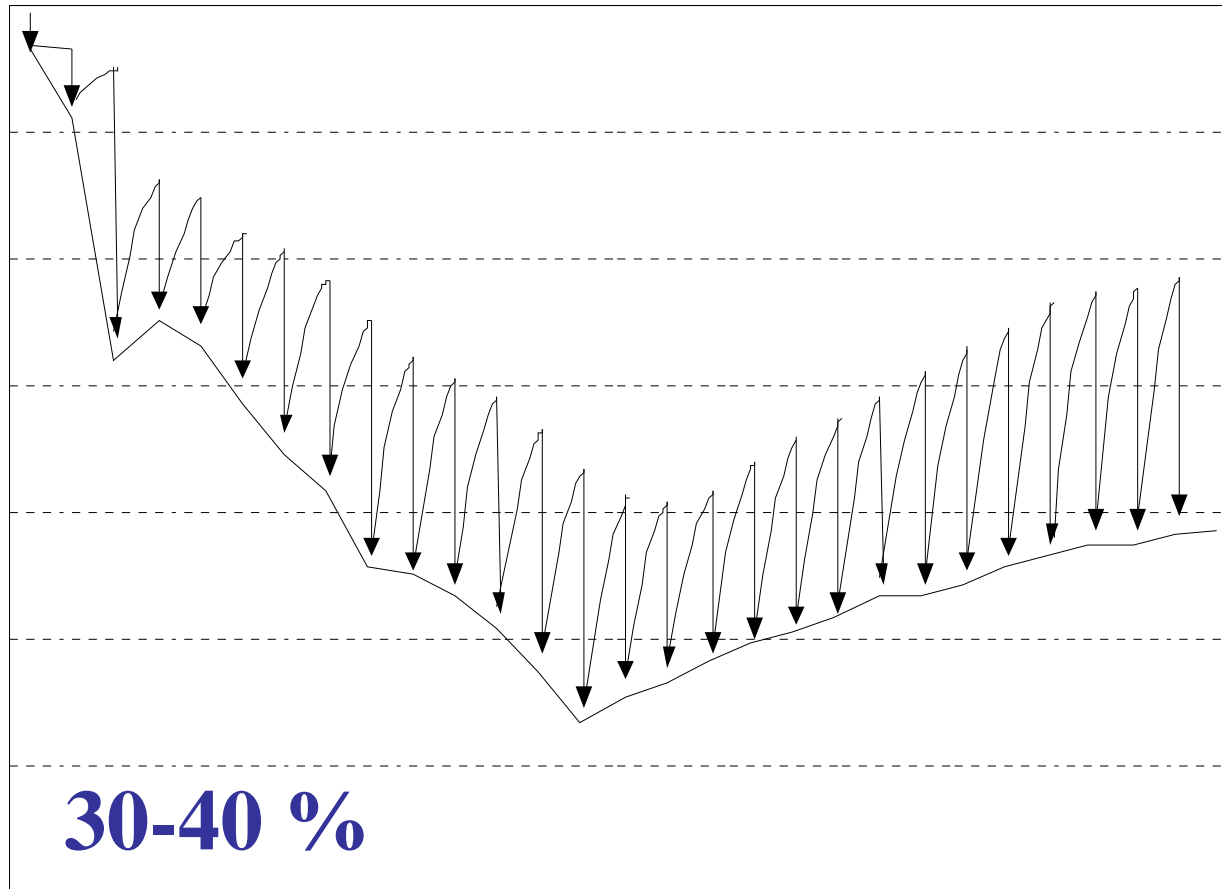
Search for each separately



Number of clusters

Stepwise search strategy

Start from the previous result



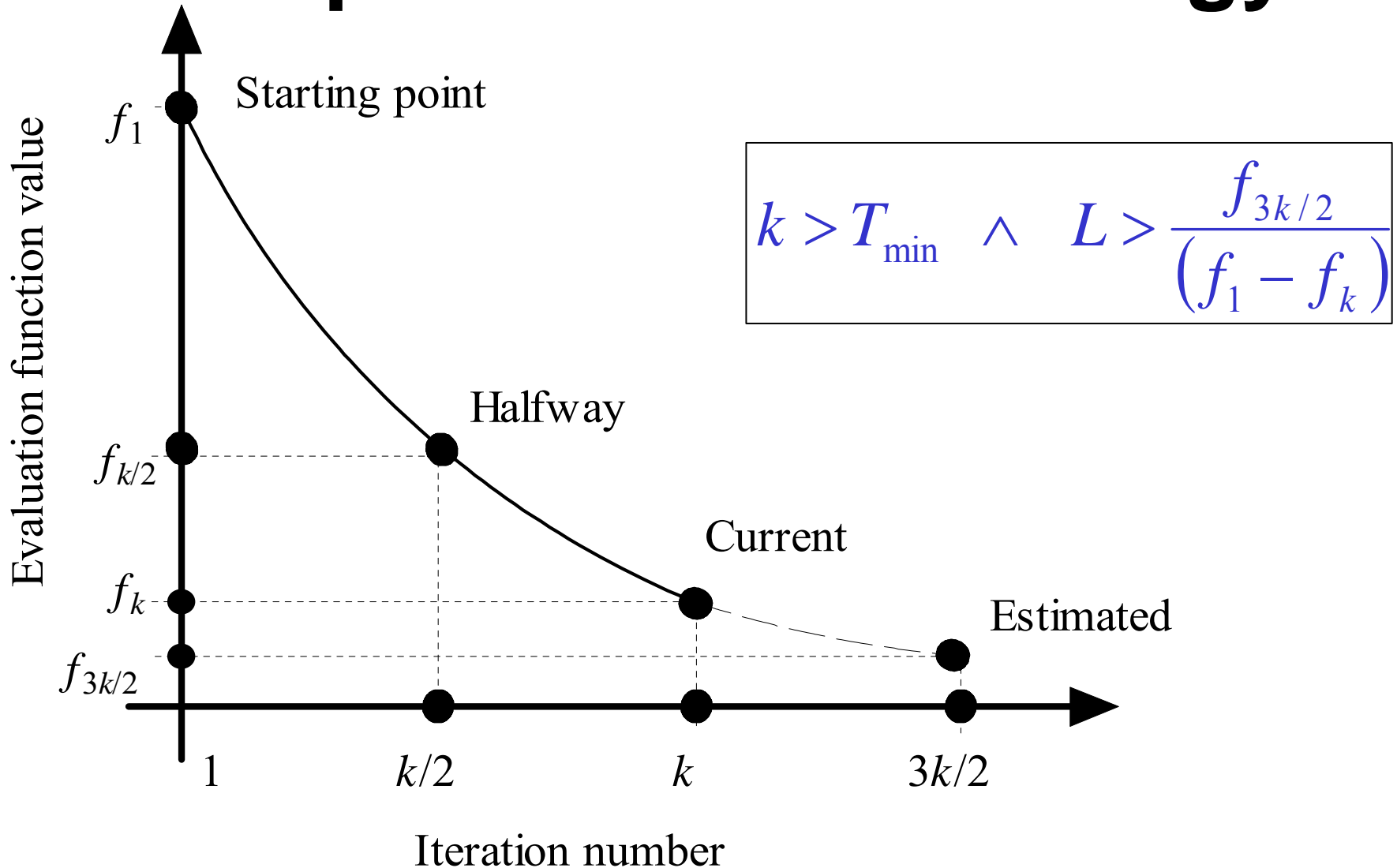
Number of clusters

Criterion guided search

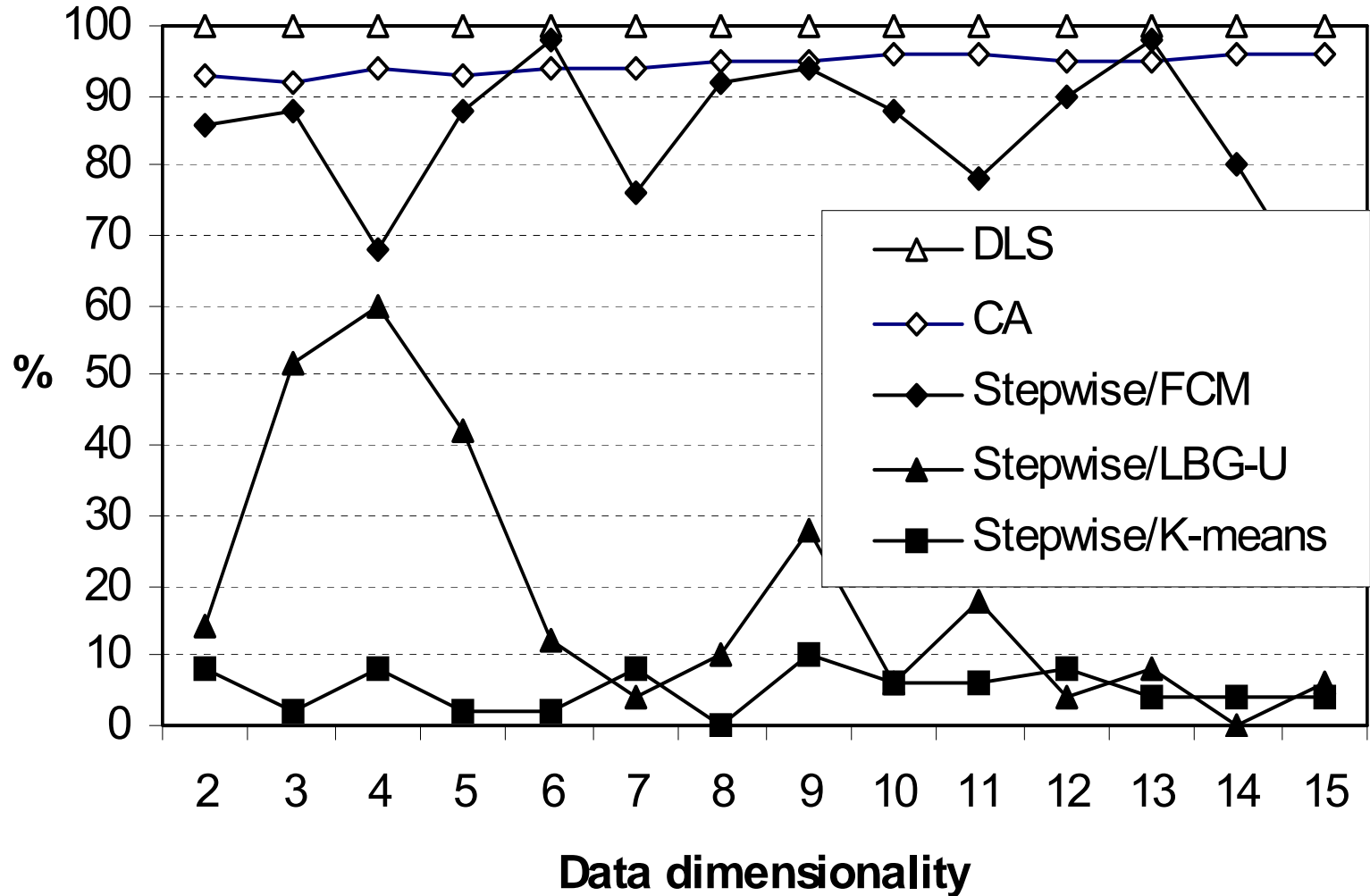
Integrate with the cost function!



Stopping criterion for stepwise search strategy



Comparison of search strategies



Open questions

Iterative algorithm (K-means or Random Swap) with criterion-guided search

... or ...

Hierarchical algorithm ???

*Potential topic for
MSc or PhD thesis
!!!*

Part VI:
External indexes

Pair-counting measures

The number of pairs that are in:

Same class **both** in P and G .

$$a = \frac{1}{2} \sum_{i=1}^K \sum_{j=1}^{K'} n_{ij} (n_{ij} - 1)$$

Same class in P but different in G .

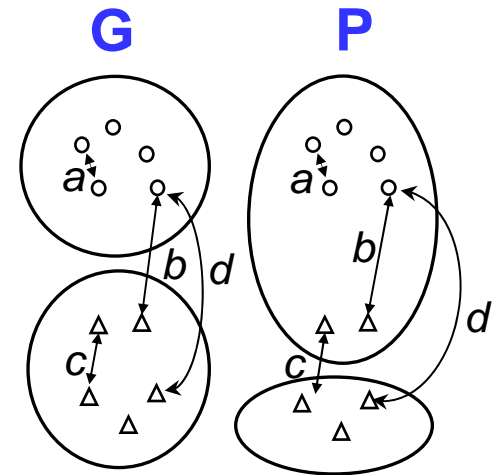
$$b = \frac{1}{2} \left(\sum_{j=1}^{K'} m_j^2 - \sum_{i=1}^K \sum_{j=1}^{K'} n_{ij}^2 \right)$$

Different classes in P but same in G .

$$c = \frac{1}{2} \left(\sum_{i=1}^K n_i^2 - \sum_{i=1}^K \sum_{j=1}^{K'} n_{ij}^2 \right)$$

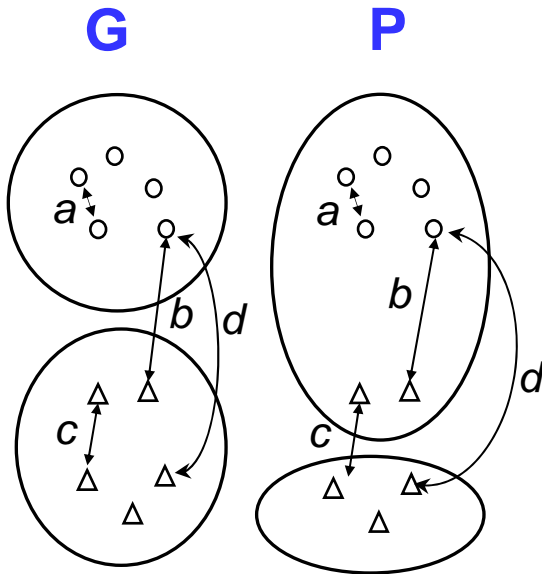
Different classes **both** in P and G .

$$d = \frac{1}{2} \left(N^2 + \sum_{i=1}^K \sum_{j=1}^{K'} n_{ij}^2 - \left(\sum_{i=1}^K n_i^2 + \sum_{j=1}^{K'} m_j^2 \right) \right)$$



Rand index

[Rand, 1971]



$$RI(P, G) = \frac{a + d}{a + b + c + d}$$

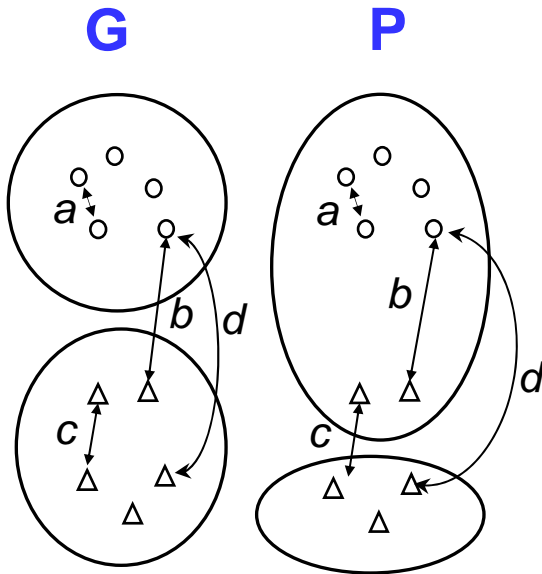
$$a = 20 \quad b = 24$$

$$d = 72 \quad c = 20$$

$$\text{Rand index} = (20 + 72) / (20 + 24 + 20 + 72) = 92 / 136 = \mathbf{0.68}$$

Rand and Adjusted Rand index

[Hubert and Arabie, 1985]



$$ARI = \frac{RI - E(RI)}{1 - E(RI)}$$

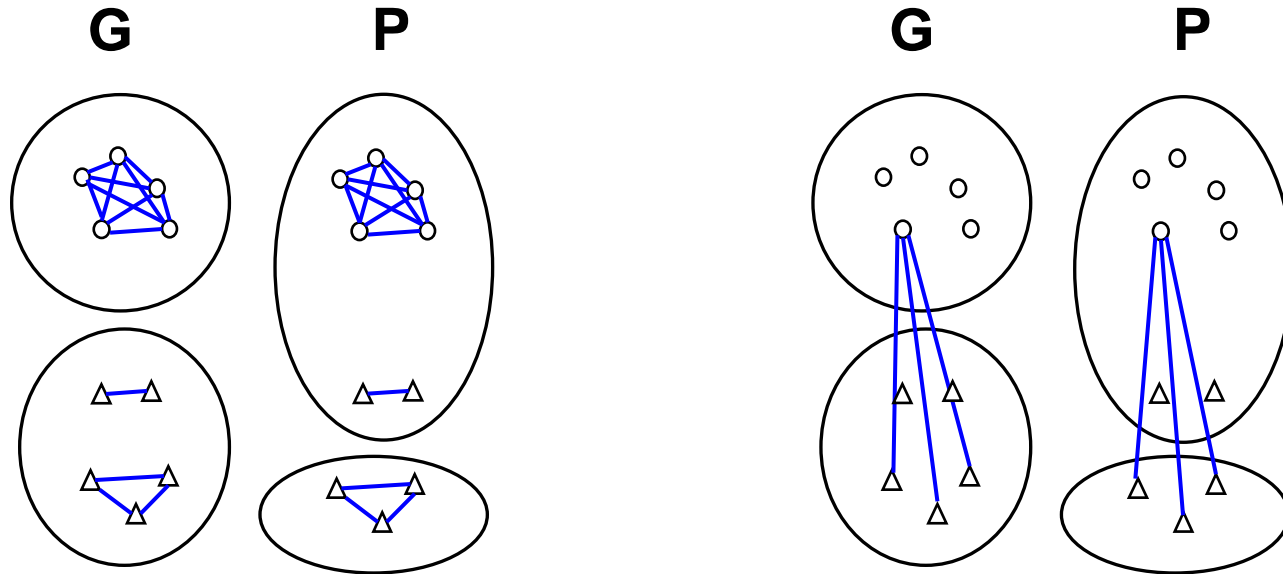
$$a = 20 \quad b = 24$$

$$d = 72 \quad c = 20$$

Adjusted Rand = (to be calculated) = **0.xx**

Rand statistics

Positive examples

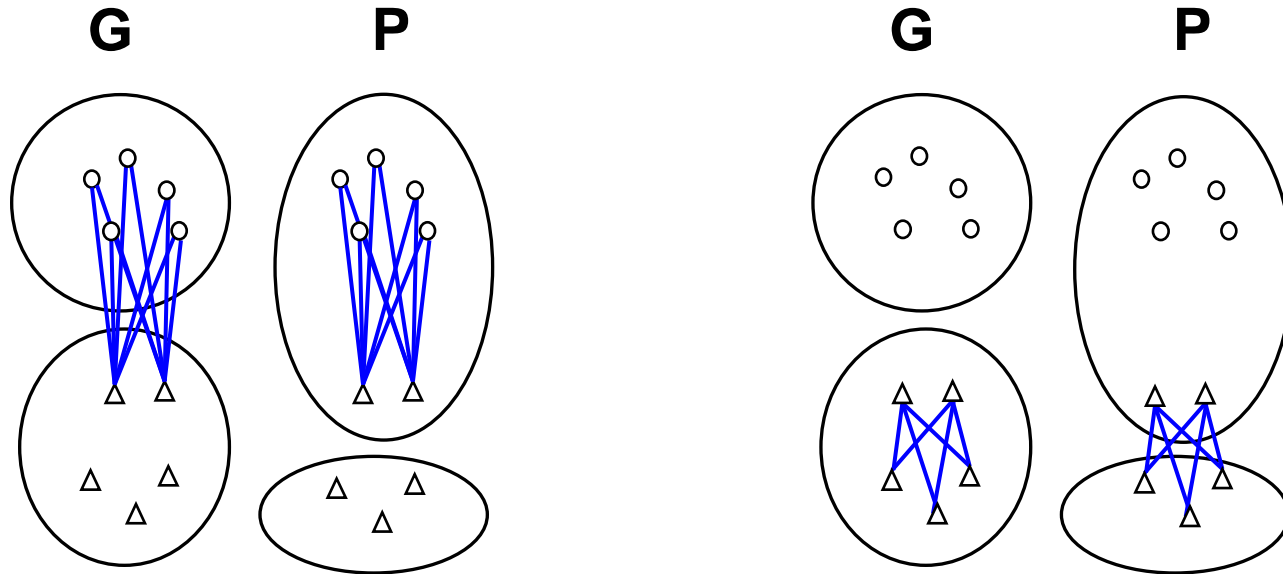


$$a = 20$$

$$d = 72$$

Rand statistics

Negative examples



$$b = 24$$

$$c = 20$$

External indexes

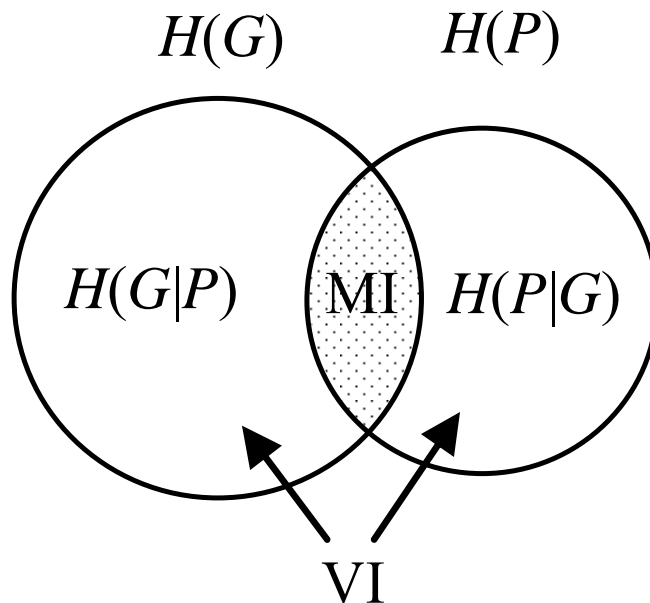
- Pair counting
- Information theoretic
- Set matching

Information-theoretic measures

- Based on the concept of entropy
- *Mutual Information* (MI): the shared informatio:

$$MI(P, G) = \sum_{i=1}^K \sum_{j=1}^{K'} p(P_i, G_j) \log \frac{p(P_i, G_j)}{p(P_i)p(G_j)}$$

- *Variation of Information* (VI) is complement of MI



Set-matching measures

Categories

- Point-level
- Cluster-level

Three problems

- How to measure the similarity of two clusters?
- How to pair clusters?
- How to calculate overall similarity?

Similarity of two clusters

Jaccard $J = \frac{|P_i \cap G_j|}{|P_i \cup G_j|}$

Sorensen-Dice $SD = \frac{2|P_i \cap G_j|}{|P_i| + |G_j|}$

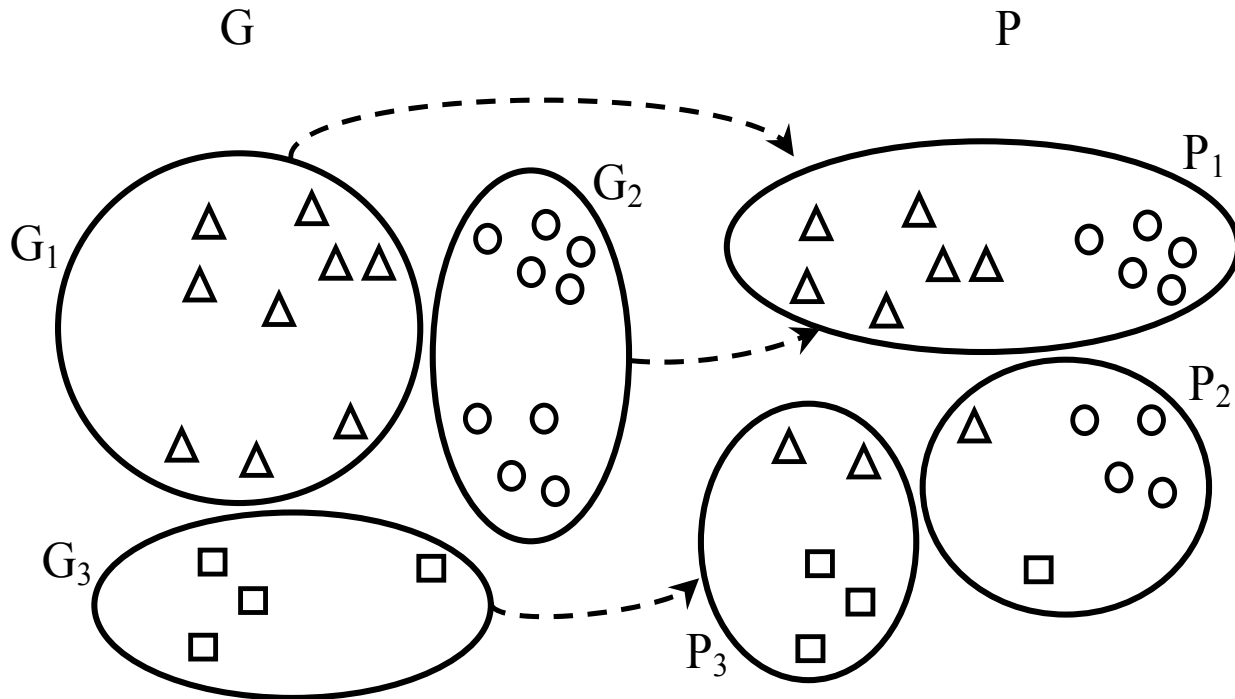
Braun-Banquet $BB = \frac{|P_i \cap G_j|}{\max(|P_i|, |G_j|)}$

The diagram illustrates three sets: P₂ (n₂=200), P₁ (n₁=250), and P₃ (n₃=1000). P₂ and P₁ are overlapping circles, and both are contained within the larger circle P₃.

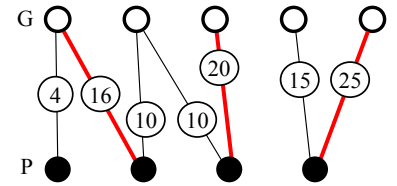
Measure:	P ₁ , P ₂	P ₁ , P ₃
Criterion H / NVD / CSI	200	250
J	0.80	0.25
SD	0.89	0.40
BB	0.80	0.25

Matching

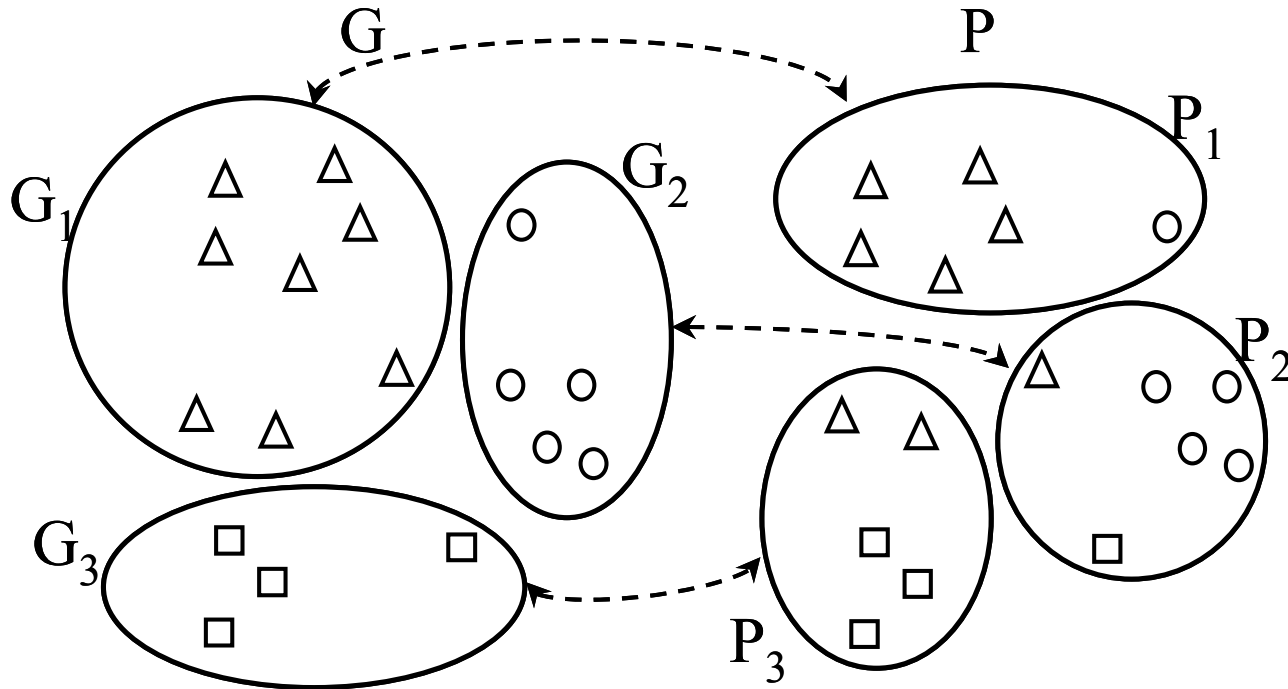
Every cluster is mapped to the cluster with maximum overlap



Pairing

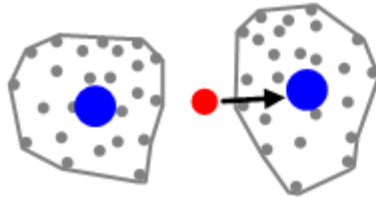
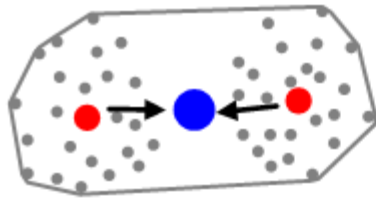


Optimal pairing by Hungarian algorithm
or greedy pairing



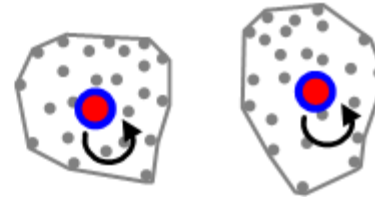
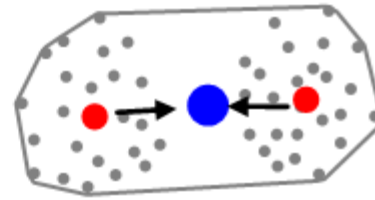
Matching vs. Pairing

3-vs-3 clusters



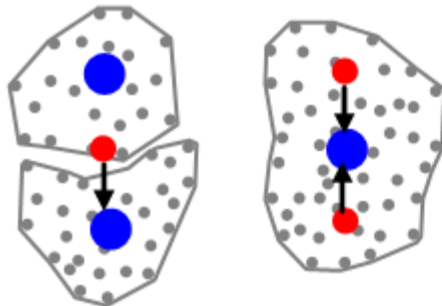
Matching=75%, Pairing=50%

3-vs-4 clusters



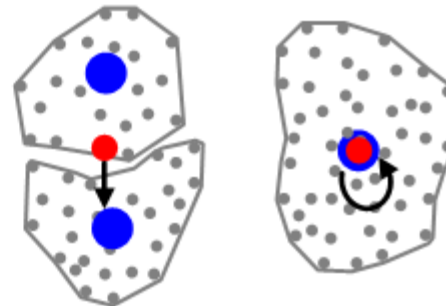
Matching=87%, Pairing=75%

3-vs-3 clusters



Matching=75%, Pairing=50%

3-vs-2 clusters



Matching=87%, Pairing=75%

Summary of matching

	Pairing/ Matching	Matching criterion	Algorithm
FM	Matching	SD	One-way
CH	Pairing	$ P_i \cap G_j $	Greedy
NVD	Matching	$ P_i \cap G_j $	Two-way
Purity	Matching	$ P_i \cap G_j $	One-way
PSI	Pairing	BB	Optimal
CI	Matching	Centroid distance	Two-way
CSI	Matching	Centroid distance	Two-way
CR	Pairing	Centroid distance	Greedy

Overall similarity

	Total summation	Range	Normalization
FM	similarity of matched clusters	$[0, 1]$	N
CH	Shared objects	$[0, 1]$	N
NVD	Shared objects in both directions	$[0, 1]$	$2N$
Purity	Shared objects in one direction	$[0, 1]$	N
PSI	Normalized similarity of paired clusters	$[0, 1]$	K
CI	Orphan clusters	$[0, K-1]$	-
CSI	Shared objects in both directions	$[0, 1]$	$2N$
CR	Unstable clusters	$[0, 1]$	K

Normalized Van Dongen

Closely related to Purity and CSI
(Assumed that matching is symmetric)

$$NVD = 1 - \frac{\sum_{i=1}^K n_{ij} + \sum_{j=1}^{K'} n_{ji}}{2N} = 1 - \frac{2 \sum_{i=1}^K n_{ij}}{2N} =$$

$$1 - \frac{\sum_{i=1}^K n_{ij}}{N} = CH = 1 - Purity = 1 - CSI$$

Pair Set Index (PSI)

M. Rezaei and P. Fränti, "Set matching measures for external cluster validity",
IEEE Trans. on Knowledge and Data Engineering, August 2016.

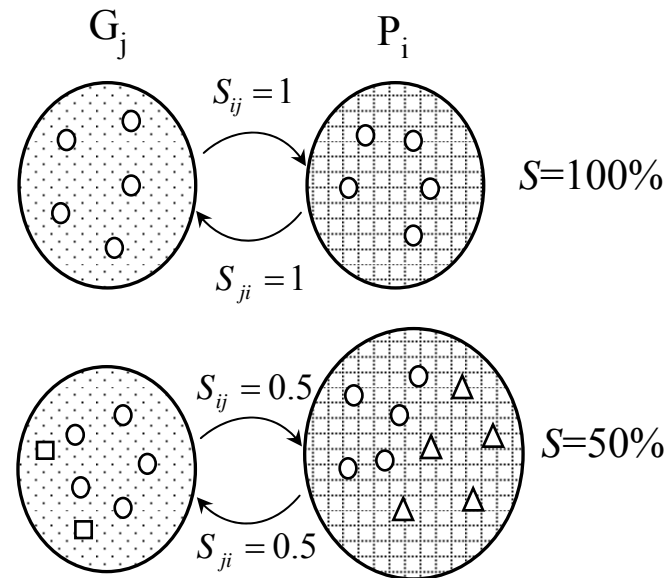
– Similarity of two clusters:

$$S_{ij} = \frac{n_{ij}}{\max(|P_i|, |G_j|)}$$

– Total similarity:

$$S_{PG} = \sum_i S_{ij}$$

– Pairing by Hungarian:



Pair Set Index (PSI)

Correction for chance

$$\text{Max}(S) = \min(K, K')$$

$$E(S) = \sum_{i=1}^{\min(K, K')} \frac{n_i \times (m_i / N)}{\max(n_i, m_i)}$$

size of clusters in $P : n_1 > n_2 > \dots > n_K$
size of clusters in $G : m_1 > m_2 > \dots > m_{K'}$

$$\text{Transformation : } \begin{cases} \text{Max}(S) \rightarrow 1 \\ E(S) \rightarrow 0 \end{cases}$$

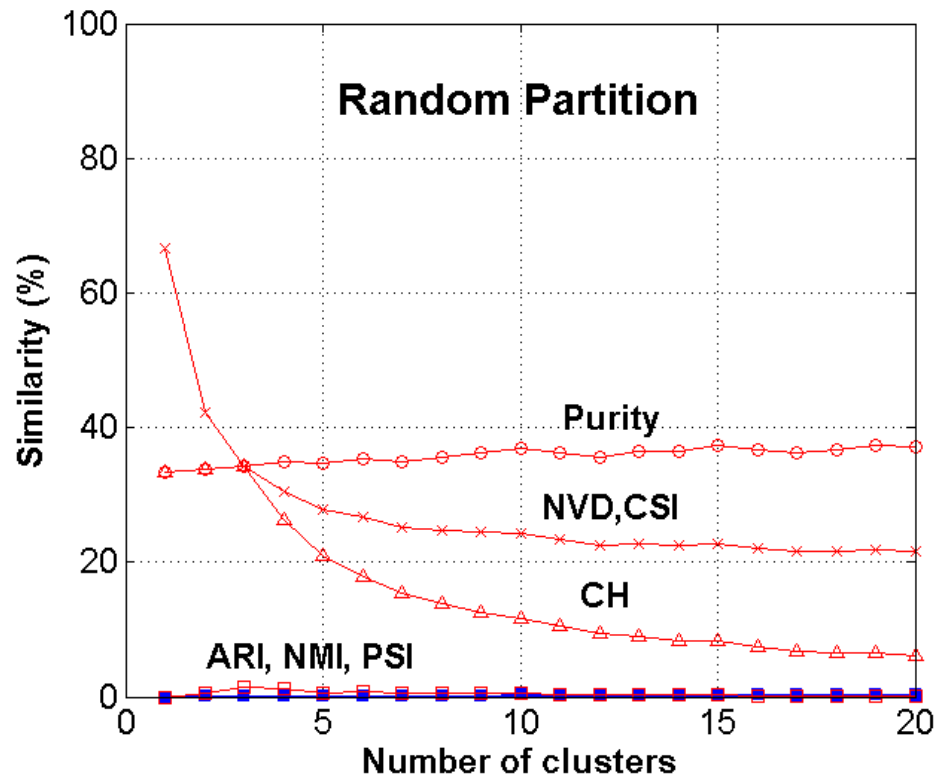
$$\text{PSI} = \begin{cases} \frac{S - E}{\max(K, K') - E} & S \geq E, \max(K, K') > 1 \\ 0 & S < E \\ 1 & K = K' = 1 \end{cases}$$

Properties of PSI

- Symmetric
- Normalized to number of clusters
- Normalized to size of clusters
- Adjusted
- Range in $[0, 1]$
- Number of clusters can be different

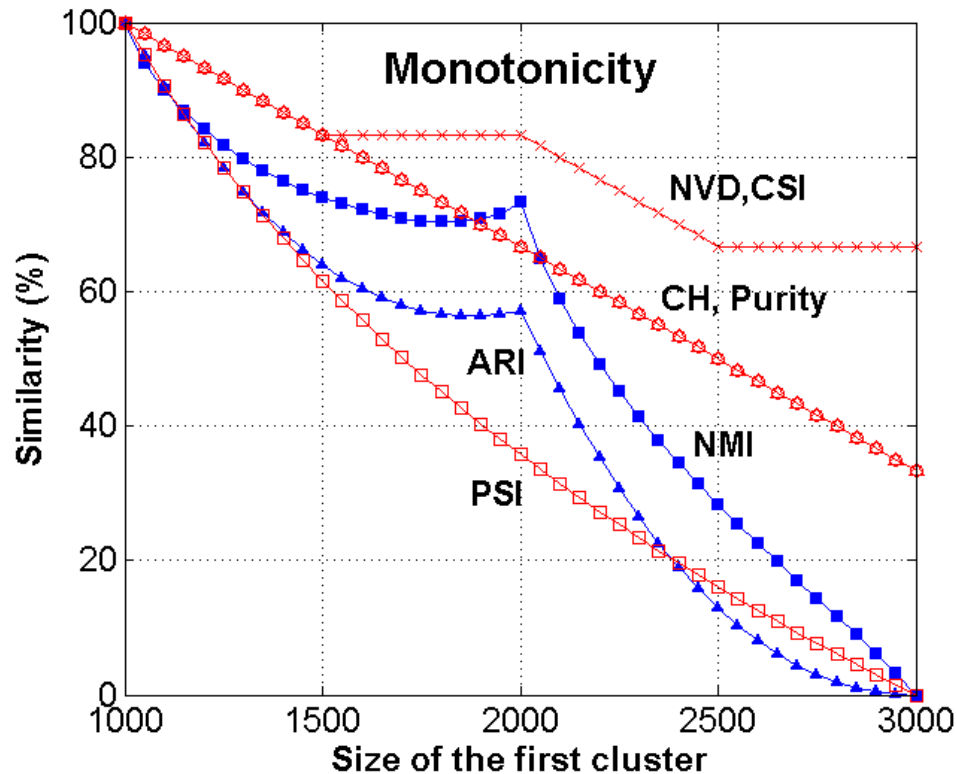
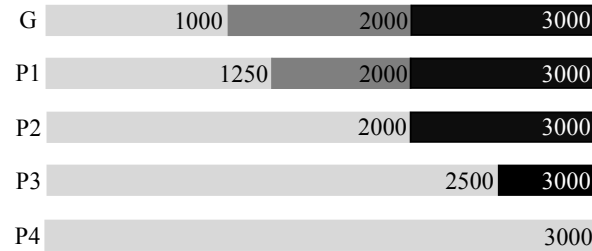
Random partitioning

Changing number of clusters in P from 1 to 20



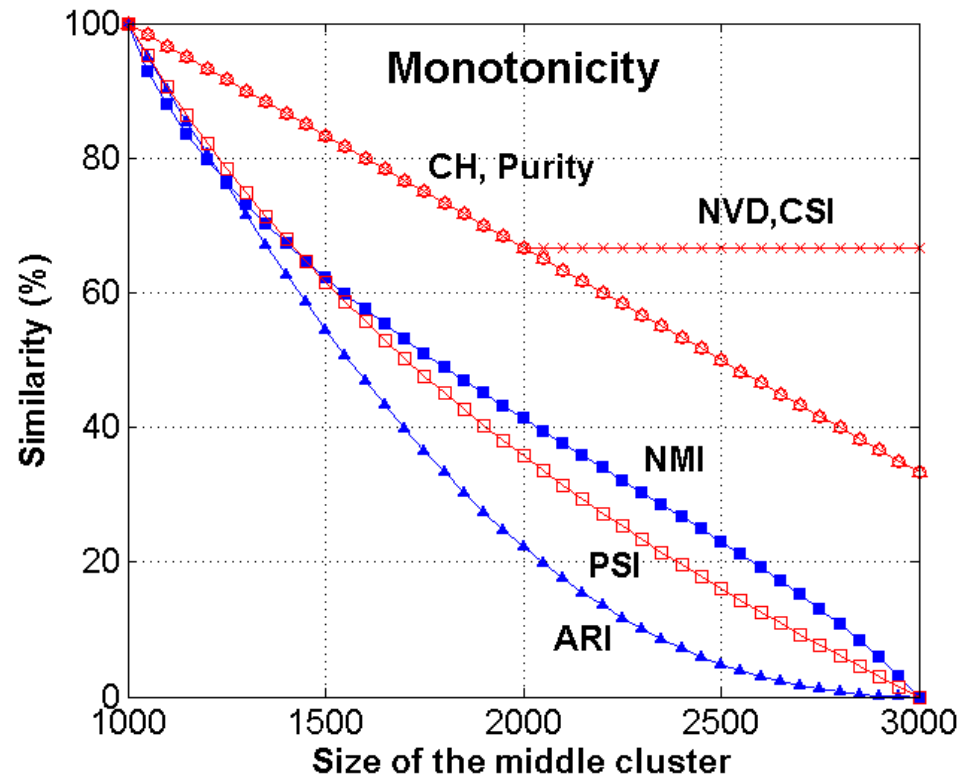
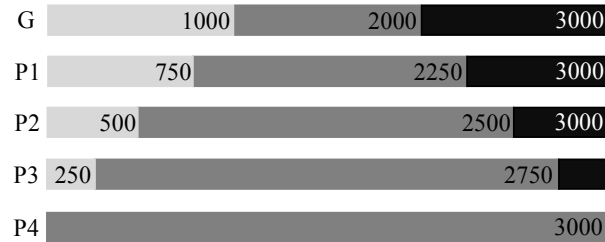
Monotonicity

Enlarging the first cluster



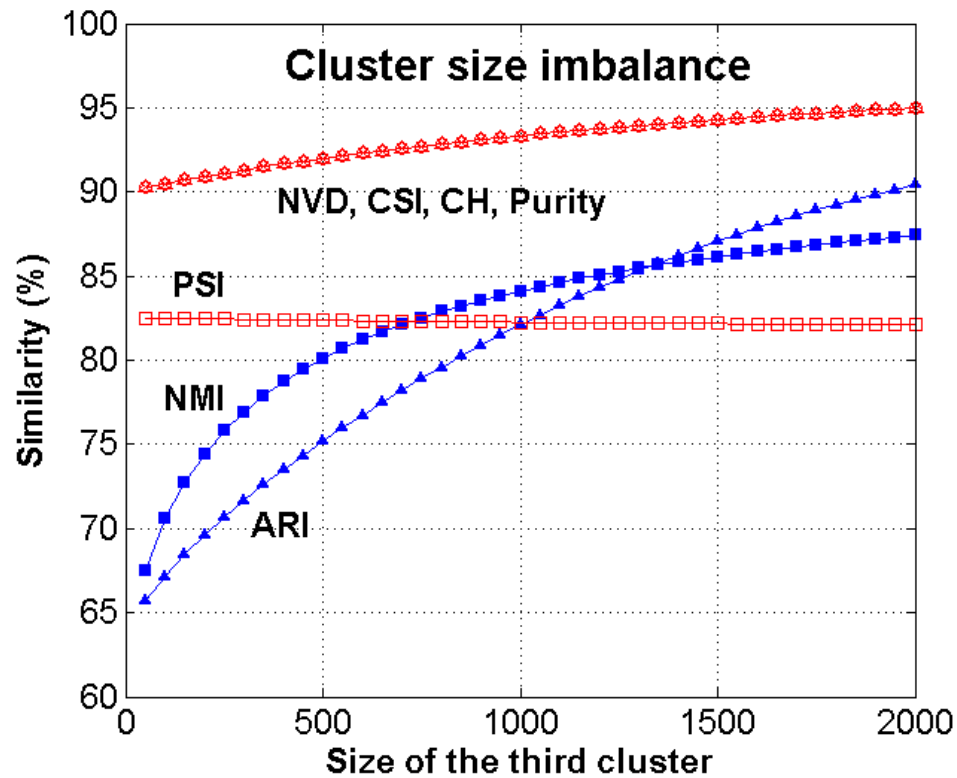
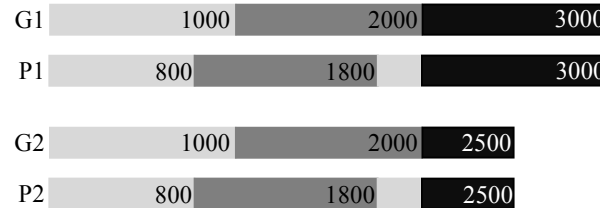
Monotonicity

Enlarging the second cluster



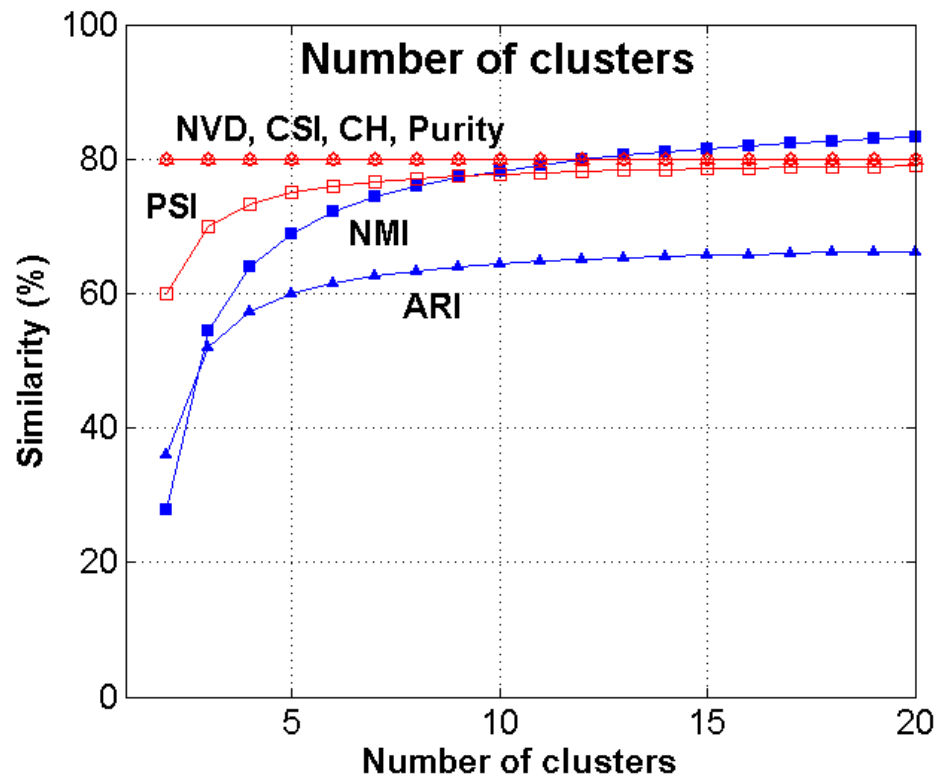
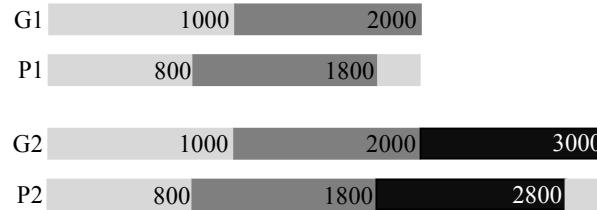
Cluster size imbalance

Same error in first two clusters



Number of clusters

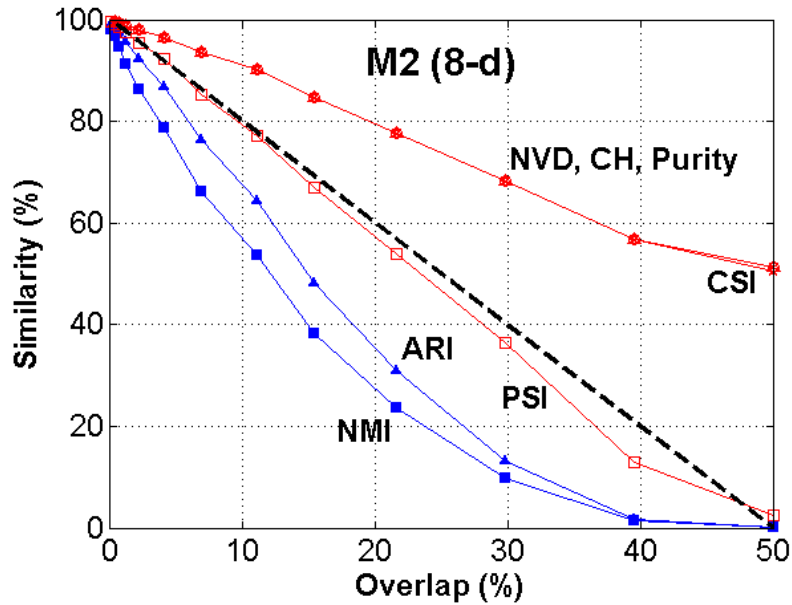
Always 200 errors; k varies



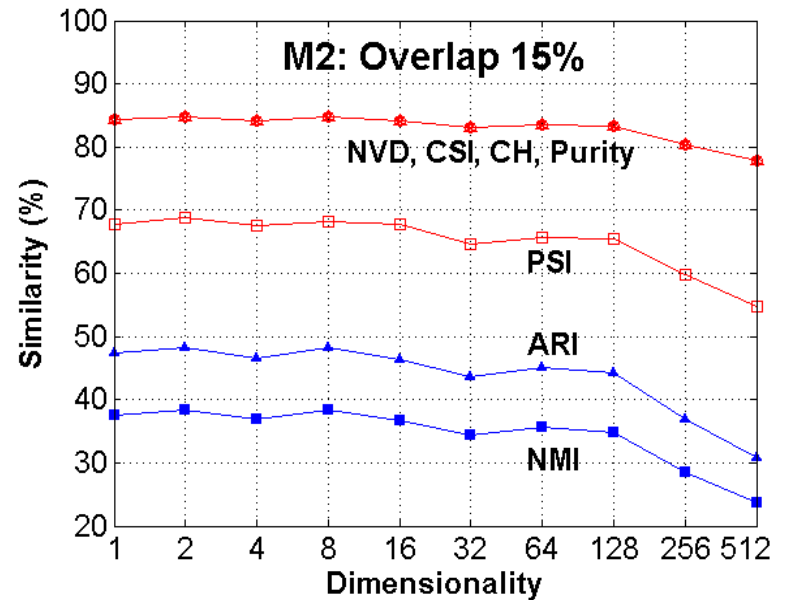
Overlap and dimensionality

Two clusters with varying overlap and dimensions

Overlap varies



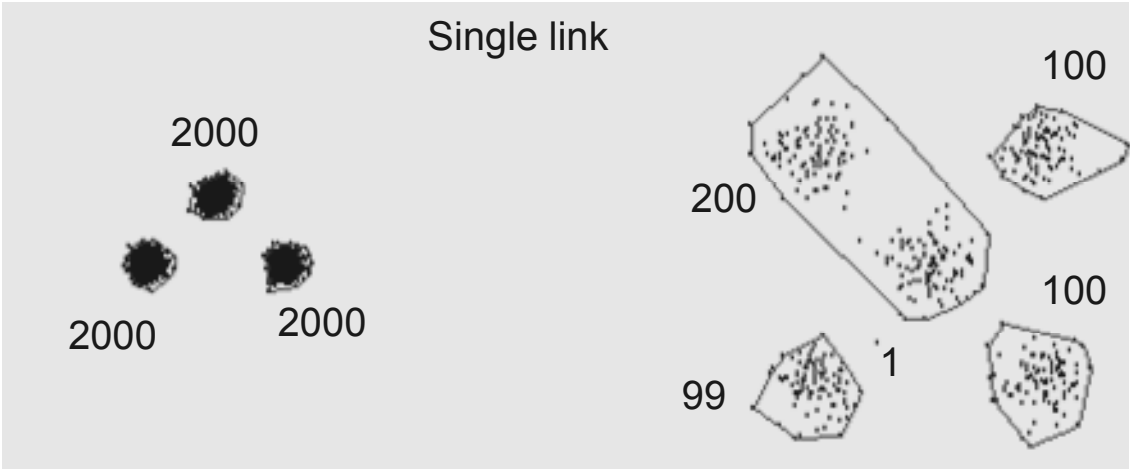
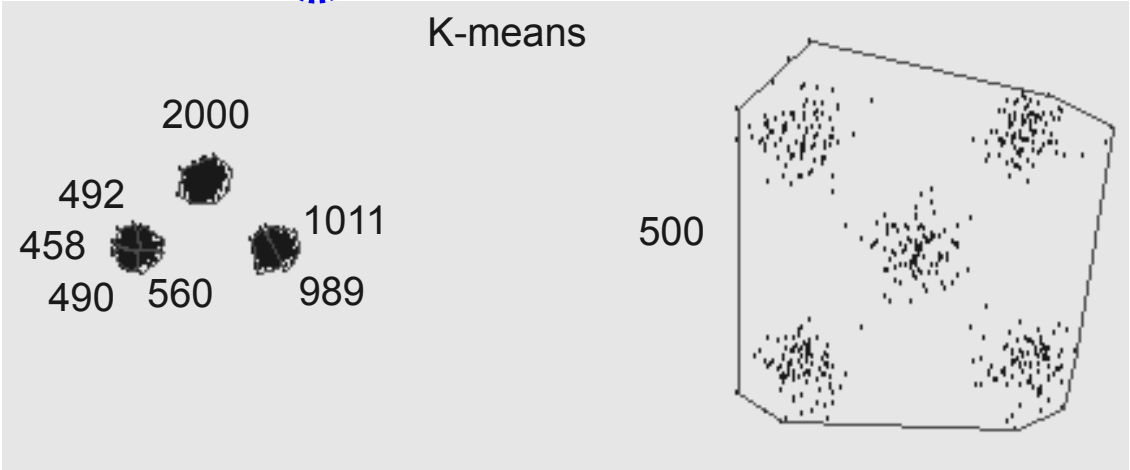
Dimensions



Unbalance

Algorithms	External indexes			
	ARI	NMI	NVD	PSI
RS	1.00	1.00	1.00	1.00
AC	1.00	1.00	1.00	1.00
SL	1.00	0.99	0.99	0.78
KM	0.66	0.77	0.78	0.18

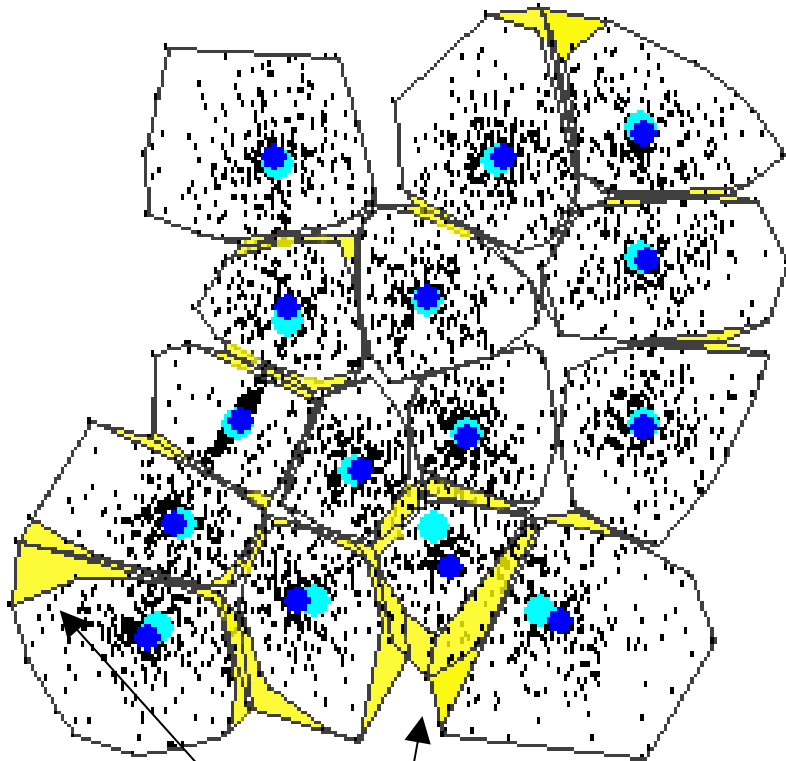
Unrealistic high



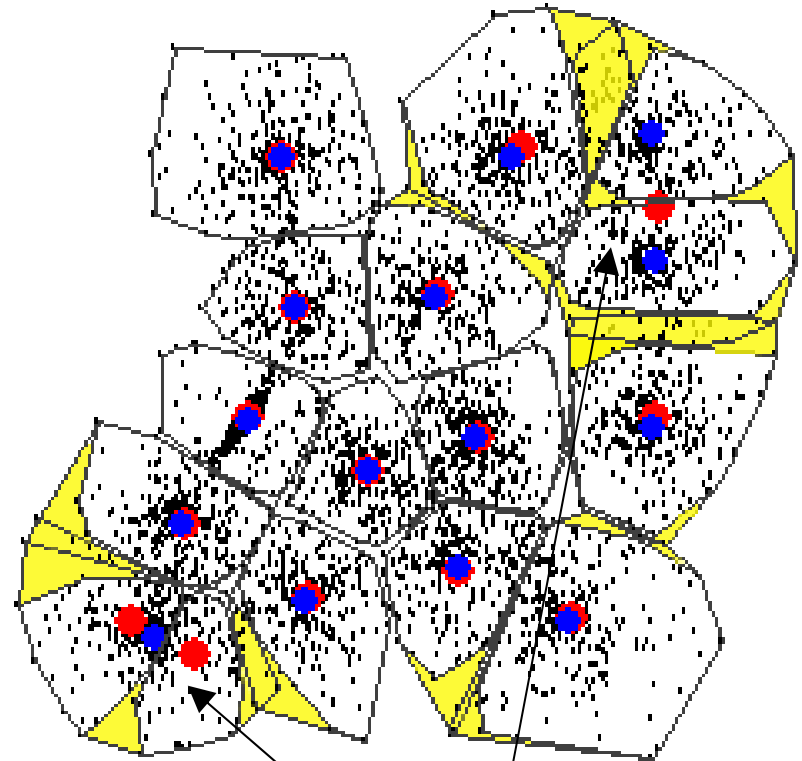
Part VII:

Cluster-level measure

Comparing partitions of centroids



Point-level differences



Cluster-level mismatches

Centroid index (CI)

[Fränti, Rezaei, Zhao, Pattern Recognition, 2014]

Given two sets of centroids C and C' ,
find nearest neighbor mappings ($C \rightarrow C'$):

$$q_i \leftarrow \arg \min_{1 \leq j \leq K_2} \|c_i - c'_j\|^2, \quad \forall i \in [1, K_1]$$

Detect prototypes with no mapping:

$$orphan(c'_j) = \begin{cases} 1, & q_i \neq j \quad \forall i \\ 0, & \text{otherwise} \end{cases}$$

Centroid index:

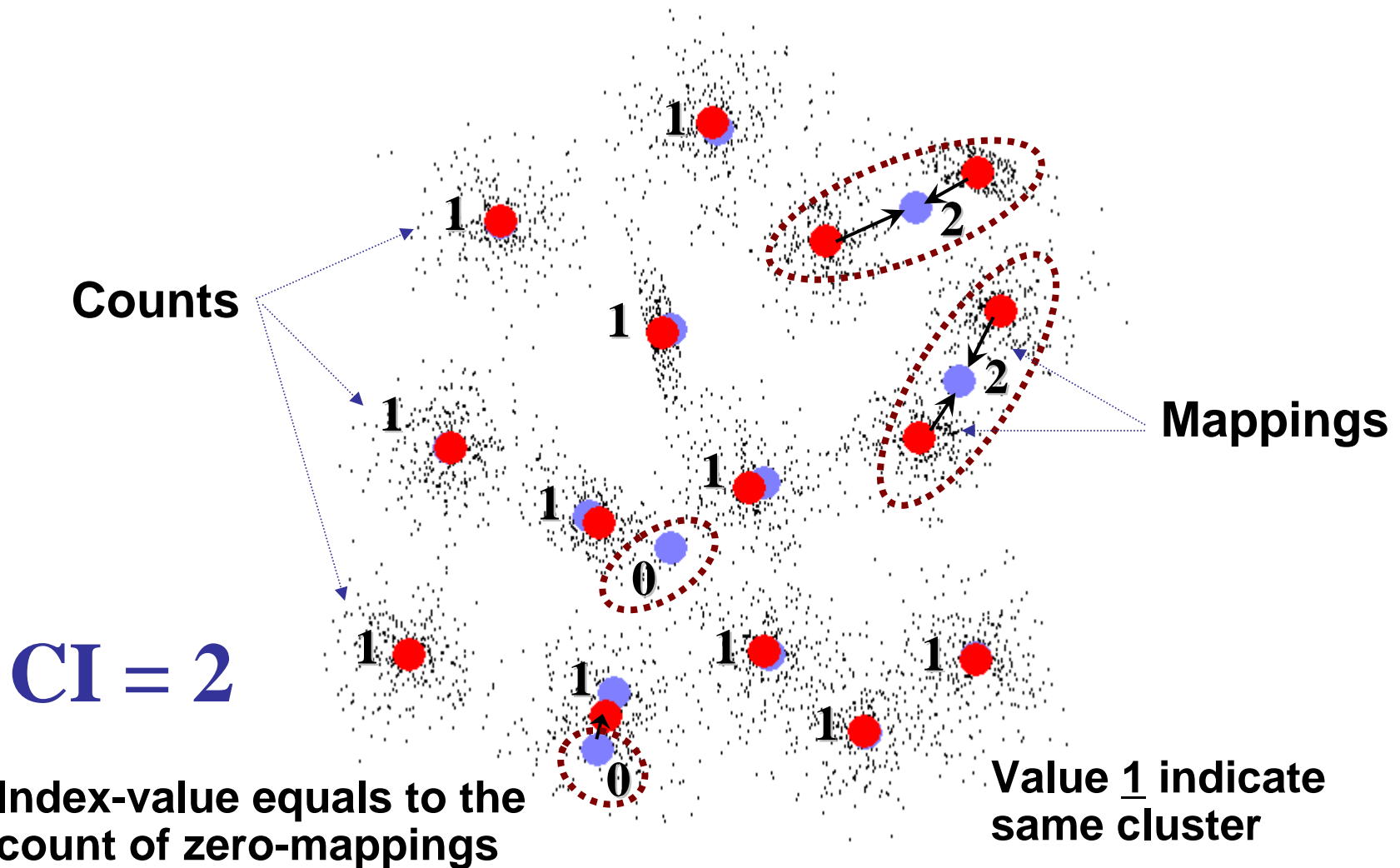
$$CI_1(C, C') = \sum_{j=1}^{K_2} orphan(c'_j)$$

Number of zero mappings!

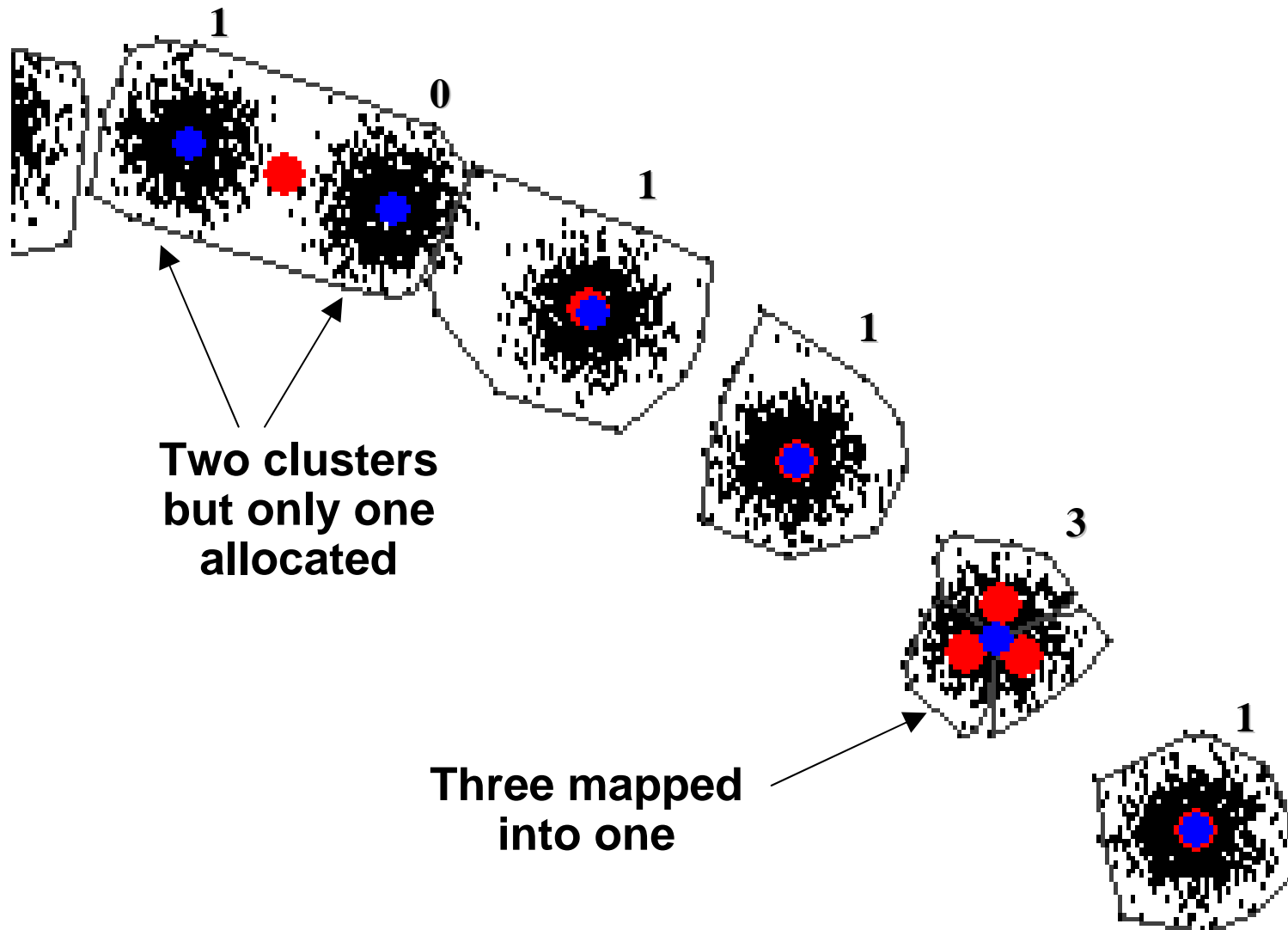


Example of centroid index

Data S_2

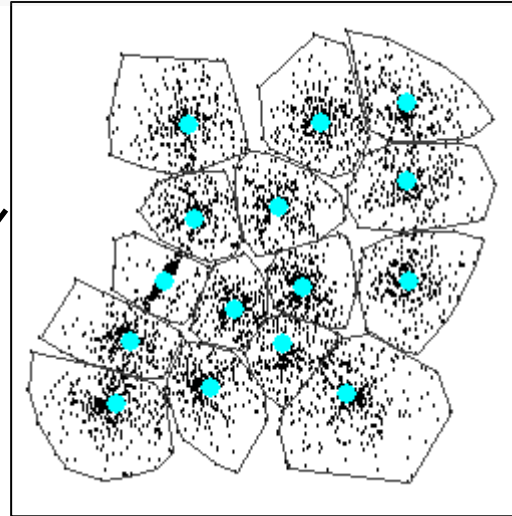


Example of the Centroid index



Adjusted Rand vs. Centroid index

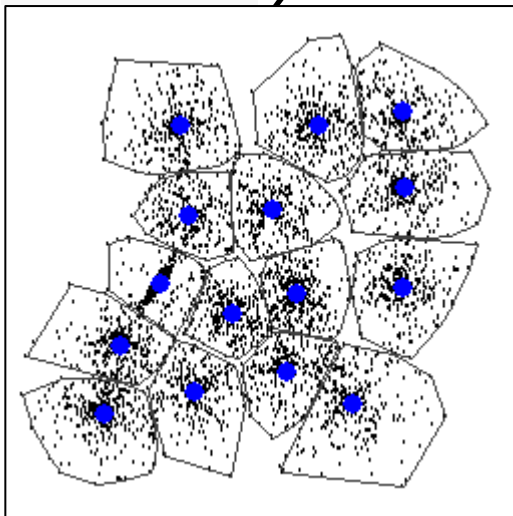
Merge-based (PNN)



ARI=0.91
CI=0

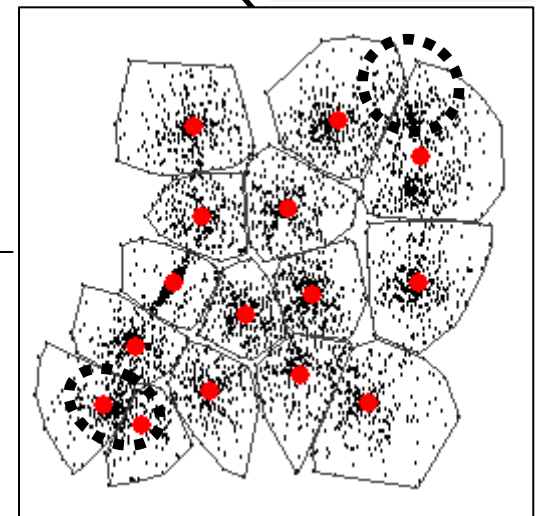
ARI=0.82
CI=1

Random
Swap



ARI=0.88
CI=1

K-means



Centroid index properties

- Mapping is not symmetric ($C \rightarrow C' \neq C' \rightarrow C$)

- Symmetric centroid index:

$$CI_2(C, C') = \max\{CI_1(C, C'), CI_1(C', C)\}$$

- Pointwise variant (Centroid Similarity Index):

– Matching clusters based on CI

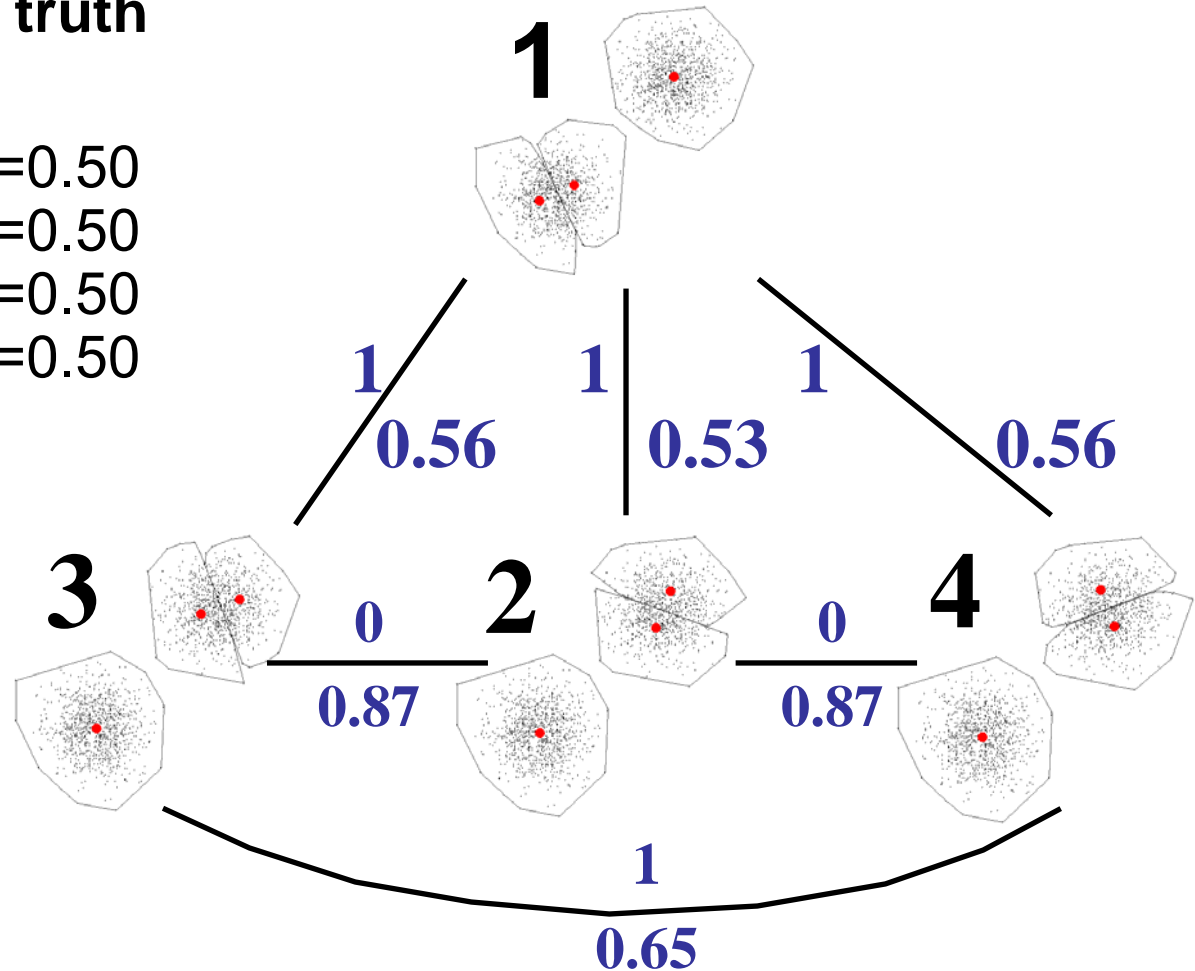
– Similarity of clusters

$$CSI = \frac{S_{12} + S_{21}}{2} \text{ where } S_{12} = \frac{\sum_{i=1}^{K_1} C_i \cap C_j}{N} \quad S_{21} = \frac{\sum_{j=1}^{K_2} C_j \cap C_i}{N}$$

Centroid index

Distance to ground truth
(2 clusters):

1 ↔ GT	CI=1	CSI=0.50
2 ↔ GT	CI=1	CSI=0.50
3 ↔ GT	CI=1	CSI=0.50
4 ↔ GT	CI=1	CSI=0.50



Mean Squared Errors

Data set	Clustering quality (MSE)							
	KM	RKM	KM++	XM	AC	RS	GKM	GA
<i>Bridge</i>	179.76	176.92	173.64	179.73	168.92	164.64	164.78	161.47
<i>House</i>	6.67	6.43	6.28	6.20	6.27	5.96	5.91	5.87
<i>Miss America</i>	5.95	5.83	5.52	5.92	5.36	5.28	5.21	5.10
<i>House</i>	3.61	3.28	2.50	3.57	2.62	2.83	-	2.44
<i>Birch</i> ₁	5.47	5.01	4.88	5.12	4.73	4.64	-	4.64
<i>Birch</i> ₂	7.47	5.65	3.07	6.29	2.28	2.28	-	2.28
<i>Birch</i> ₃	2.51	2.07	1.92	2.07	1.96	1.86	-	1.86
<i>S</i> ₁	19.71	8.92	8.92	8.92	8.93	8.92	8.92	8.92
<i>S</i> ₂	20.58	13.28	13.28	15.87	13.44	13.28	13.28	13.28
<i>S</i> ₃	19.57	16.89	16.89	16.89	17.70	16.89	16.89	16.89
<i>S</i> ₄	17.73	15.70	15.70	15.71	17.52	15.70	15.71	15.70

Adjusted Rand Index

Data set	Adjusted Rand Index (ARI)							
	KM	RKM	KM++	XM	AC	RS	GKM	GA
<i>Bridge</i>	0.38	0.40	0.39	0.37	0.43	0.52	0.50	1
<i>House</i>	0.40	0.40	0.44	0.47	0.43	0.53	0.53	1
<i>Miss America</i>	0.19	0.19	0.18	0.20	0.20	0.20	0.23	1
<i>House</i>	0.46	0.49	0.52	0.46	0.49	0.49	-	1
<i>Birch</i> ₁	0.85	0.93	0.98	0.91	0.96	1.00	-	1
<i>Birch</i> ₂	0.81	0.86	0.95	0.86	1	1	-	1
<i>Birch</i> ₃	0.74	0.82	0.87	0.82	0.86	0.91	-	1
<i>S</i> ₁	0.83	1.00	1.00	1.00	1.00	1.00	1.00	1.00
<i>S</i> ₂	0.80	0.99	0.99	0.89	0.98	0.99	0.99	0.99
<i>S</i> ₃	0.86	0.96	0.96	0.96	0.92	0.96	0.96	0.96
<i>S</i> ₄	0.82	0.93	0.93	0.94	0.77	0.93	0.93	0.93

Normalized Mutual information

Data set	Normalized Mutual Information (NMI)							
	KM	RKM	KM++	XM	AC	RS	GKM	GA
<i>Bridge</i>	0.77	0.78	0.78	0.77	0.80	0.83	0.82	1.00
<i>House</i>	0.80	0.80	0.81	0.82	0.81	0.83	0.84	1.00
<i>Miss America</i>	0.64	0.64	0.63	0.64	0.64	0.66	0.66	1.00
<i>House</i>	0.81	0.81	0.82	0.81	0.81	0.82	-	1.00
<i>Birch</i> ₁	0.95	0.97	0.99	0.96	0.98	1.00	-	1.00
<i>Birch</i> ₂	0.96	0.97	0.99	0.97	1.00	1.00	-	1.00
<i>Birch</i> ₃	0.90	0.94	0.94	0.93	0.93	0.96	-	1.00
<i>S</i> ₁	0.93	1.00	1.00	1.00	1.00	1.00	1.00	1.00
<i>S</i> ₂	0.90	0.99	0.99	0.95	0.99	0.93	0.99	0.99
<i>S</i> ₃	0.92	0.97	0.97	0.97	0.94	0.97	0.97	0.97
<i>S</i> ₄	0.88	0.94	0.94	0.95	0.85	0.94	0.94	0.94

Normalized Van Dongen

Data set	Normalized Van Dongen (NVD)							
	KM	RKM	KM++	XM	AC	RS	GKM	GA
<i>Bridge</i>	0.45	0.42	0.43	0.46	0.38	0.32	0.33	0.00
<i>House</i>	0.44	0.43	0.40	0.37	0.40	0.33	0.31	0.00
<i>Miss America</i>	0.60	0.60	0.61	0.59	0.57	0.55	0.53	0.00
<i>House</i>	0.40	0.37	0.34	0.39	0.39	0.34	-	0.00
<i>Birch</i> ₁	0.09	0.04	0.01	0.06	0.02	0.00	-	0.00
<i>Birch</i> ₂	0.12	0.08	0.03	0.09	0.00	0.00	-	0.00
<i>Birch</i> ₃	0.19	0.12	0.10	0.13	0.13	0.06	-	0.00
<i>S</i> ₁	0.09	0.00	0.00	0.00	0.00	0.00	0.00	0.00
<i>S</i> ₂	0.11	0.00	0.00	0.06	0.01	0.04	0.00	0.00
<i>S</i> ₃	0.08	0.02	0.02	0.02	0.05	0.00	0.00	0.02
<i>S</i> ₄	0.11	0.04	0.04	0.03	0.13	0.04	0.04	0.04

Centroid Index

Data set	C-Index (CI_2)							
	KM	RKM	KM++	XM	AC	RS	GKM	GA
<i>Bridge</i>	74	63	58	81	33	33	35	0
<i>House</i>	56	45	40	37	31	22	20	0
<i>Miss America</i>	88	91	67	88	38	43	36	0
<i>House</i>	43	39	22	47	26	23	---	0
<i>Birch</i> ₁	7	3	1	4	0	0	---	0
<i>Birch</i> ₂	18	11	4	12	0	0	---	0
<i>Birch</i> ₃	23	11	7	10	7	2	---	0
<i>S</i> ₁	2	0	0	0	0	0	0	0
<i>S</i> ₂	2	0	0	1	0	0	0	0
<i>S</i> ₃	1	0	0	0	0	0	0	0
<i>S</i> ₄	1	0	0	0	1	0	0	0

Centroid Similarity Index

Data set	Centroid Similarity Index (CSI)							
	KM	RKM	KM++	XM	AC	RS	GKM	GA
<i>Bridge</i>	0.47	0.51	0.49	0.45	0.57	0.62	0.63	1.00
<i>House</i>	0.49	0.50	0.54	0.57	0.55	0.63	0.66	1.00
<i>Miss America</i>	0.32	0.32	0.32	0.33	0.38	0.40	0.42	1.00
<i>House</i>	0.54	0.57	0.63	0.54	0.57	0.62	---	1.00
<i>Birch</i> ₁	0.87	0.94	0.98	0.93	0.99	1.00	---	1.00
<i>Birch</i> ₂	0.76	0.84	0.94	0.83	1.00	1.00	---	1.00
<i>Birch</i> ₃	0.71	0.82	0.87	0.81	0.86	0.93	---	1.00
<i>S</i> ₁	0.83	1.00	1.00	1.00	1.00	1.00	1.00	1.00
<i>S</i> ₂	0.82	1.00	1.00	0.91	1.00	1.00	1.00	1.00
<i>S</i> ₃	0.89	0.99	0.99	0.99	0.98	0.99	0.99	0.99
<i>S</i> ₄	0.87	0.98	0.98	0.99	0.85	0.98	0.98	0.98

High quality clustering

	Method	MSE
GKM	Global K-means	164.78
RS	Random swap (5k)	164.64
GA	Genetic algorithm	161.47
RS _{8M}	Random swap (8M)	161.02
GAIS-2002	GAIS	160.72
+ RS _{1M}	GAIS + RS (1M)	160.49
+ RS _{8M}	GAIS + RS (8M)	160.43
GAIS-2012	GAIS	160.68
+ RS _{1M}	GAIS + RS (1M)	160.45
+ RS _{8M}	GAIS + RS (8M)	160.39
+ PRS	GAIS + PRS	160.33
+ RS _{8M} +	GAIS + RS (8M) +	160.28

Centroid index values

Main algorithm:	RS _{8M}	GAIS 2002			GAIS 2012					
		×	×	RS _{1M}	RS _{8M}	×	RS _{1M}	RS _{8M}	×	RS _{8M}
+ Tuning 1	×	×	×	×	×	×	×	×	×	×
+ Tuning 2	×	×	×	×	×	×	×	×	×	×
RS _{8M}	---	19	19	19	23	24	24	23	22	
GAIS (2002)	23	---	0	0	14	15	15	14	16	
+ RS _{1M}	23	0	---	0	14	15	15	14	13	
+ RS _{8M}	23	0	0	---	14	15	15	14	13	
GAIS (2012)	25	17	18	18	---	1	1	1	1	
+ RS _{1M}	25	17	18	18	1	---	0	0	1	
+ RS _{8M}	25	17	18	18	1	0	---	0	1	
+ PRS	25	17	18	18	1	0	0	---	1	
+ RS _{8M} + PRS	24	17	18	18	1	1	1	1	---	

Summary of external indexes

(existing measures)

Table 1: External Cluster Validation Measures.

Measure	Notation	Definition	Range
1 Entropy	E	$-\sum_i p_i (\sum_j \frac{p_{ij}}{p_i} \log \frac{p_{ij}}{p_i})$	$[0, \log K']$
2 Purity	P	$\sum_i p_i (\max_j \frac{p_{ij}}{p_i})$	$(0,1)$
3 F-measure	F	$\sum_j p_j \max_i [2 \frac{p_{ij}}{p_i} \frac{p_{ij}}{p_j} / (\frac{p_{ij}}{p_i} + \frac{p_{ij}}{p_j})]$	$(0,1)$
4 Variation of Information	VI	$-\sum_i p_i \log p_i - \sum_j p_j \log p_j - 2 \sum_i \sum_j p_{ij} \log \frac{p_{ij}}{p_i p_j}$	$[0, 2 \log \max(K, K')]$
5 Mutual Information	MI	$\sum_i \sum_j p_{ij} \log \frac{p_{ij}}{p_i p_j}$	$(0, \log K')$
6 Rand statistic	R	$[\binom{n}{2} - \sum_i \binom{n_i}{2} - \sum_j \binom{n_j}{2} + 2 \sum_{ij} \binom{n_{ij}}{2}] / \binom{n}{2}$	$(0,1)$
7 Jaccard coefficient	J	$\sum_{ij} \binom{n_{ij}}{2} / [\sum_i \binom{n_i}{2} + \sum_j \binom{n_j}{2} - \sum_{ij} \binom{n_{ij}}{2}]$	$[0,1]$
8 Fowlkes and Mallows index	FM	$\frac{\sum_{ij} \binom{n_{ij}}{2}}{\sqrt{\sum_i \binom{n_i}{2} \sum_j \binom{n_j}{2}}}$	$[0,1]$
9 Hubert Γ statistic I	Γ	$\frac{\binom{n}{2} \sum_{ij} \binom{n_{ij}}{2} - \sum_i \binom{n_i}{2} \sum_j \binom{n_j}{2}}{\sqrt{\sum_i \binom{n_i}{2} \sum_j \binom{n_j}{2} [(\binom{n}{2} - \sum_i \binom{n_i}{2})][(\binom{n}{2} - \sum_j \binom{n_j}{2})]}}$	$(-1,1)$
10 Hubert Γ statistic II	Γ'	$[\binom{n}{2} - 2 \sum_i \binom{n_i}{2} - 2 \sum_j \binom{n_j}{2} + 4 \sum_{ij} \binom{n_{ij}}{2}] / \binom{n}{2}$	$[0,1]$
11 Minkowski score	MS	$\sqrt{\sum_i \binom{n_i}{2} + \sum_j \binom{n_j}{2} - 2 \sum_{ij} \binom{n_{ij}}{2}} / \sqrt{\sum_j \binom{n_j}{2}}$	$[0, +\infty)$
12 classification error	ε	$1 - \frac{1}{n} \max_{\sigma} \sum_j n_{\sigma(j),j}$	$[0,1]$
13 van Dongen criterion	VD	$(2n - \sum_i \max_j n_{ij} - \sum_j \max_i n_{ij}) / 2n$	$[0, 1)$
14 micro-average precision	MAP	$\sum_i p_i (\max_j \frac{p_{ij}}{p_i})$	$(0,1)$
15 Goodman-Kruskal coefficient	GK	$\sum_i p_i (1 - \max_j \frac{p_{ij}}{p_i})$	$[0,1)$
16 Mirkin metric	M	$\sum_i n_i^2 + \sum_j n_j^2 - 2 \sum_i \sum_j n_{ij}^2$	$[0, 2 \binom{n}{2})$

Note: $p_{ij} = n_{ij}/n$, $p_i = n_i/n$, $p_j = n_j/n$.

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