

Constraint Programming for Diversity Pattern Set Mining

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GT CAVIAR
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... is "the use of sophisticated data analysis tools to **discover** unknown, **valid patterns and relationships** in large datasets"

Data mining:

- Core of KDD
- Search for regularities from transaction databases
- Pattern domain : item-sets, sequences, graphs, etc.
- examples including pattern mining, clustering, association rules, etc.

	g_1	g_2	g_3	g_4
s_1	x			x
s_2	x	x	x	
s_3		x		x
s_4	x	x	x	
s_5	x	x		x

frequent pattern : g_1g_2

association rule : $g_1g_2 \rightarrow g_3$



Problem Definition

Given the objects in $\mathcal{O}$ described with the Boolean attributes in $\mathcal{I}$, listing every itemset having a frequency above a given threshold $\theta \in \mathbb{N}$.

Input:

	a_1	a_2	$\dots$	a_n
o_1	$d_{1,1}$	$d_{1,2}$	$\dots$	$d_{1,n}$
o_2	$d_{2,1}$	$d_{2,2}$	$\dots$	$d_{2,n}$
$\vdots$	$\vdots$	$\vdots$	$\ddots$	$\vdots$
o_m	$d_{m,1}$	$d_{m,2}$	$\dots$	$d_{m,n}$

and a minimal frequency $\theta \in \mathbb{N}$.

where $d_{i,j} \in \{\text{true}, \text{false}\}$



Problem Definition

Given the objects in $\mathcal{O}$ described with the Boolean attributes in $\mathcal{I}$, listing every itemset having a frequency above a given threshold $\theta \in \mathbb{N}$.

Output: every $X \subseteq \mathcal{I}$ such that there are at least θ objects having all attributes in X .



Pattern flooding: Mining frequent itemsets

hepatitis - freq=30.0% - minsup=42

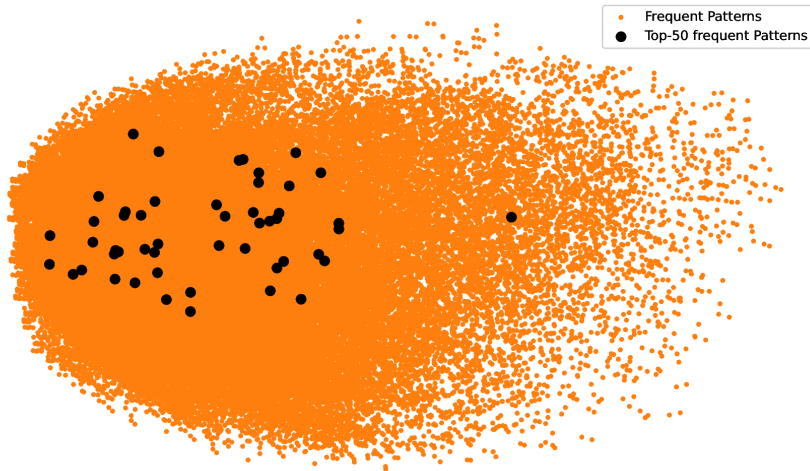
● Frequent Patterns





Pattern flooding: Mining top- k frequent itemsets

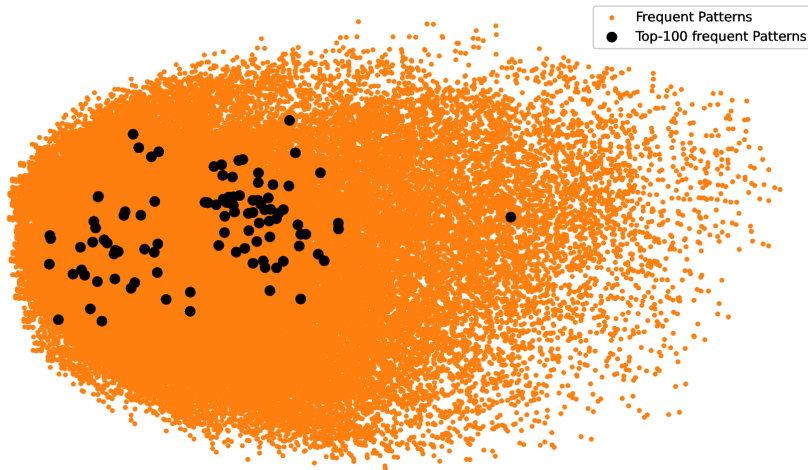
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Pattern flooding: Mining top- k frequent itemsets

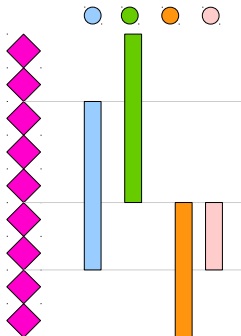
hepatitis - freq=30.0% - minsup=42





Redundancy

- Too many patterns, unmanageable and **diversity** not necessary assured
- Find a set of patterns that is:
 - small
 - non-redundant
- Several approaches for mining non-redundant patterns :
 - Mikis (Knobbe et al., ECML-PKDD'06)
 - Piker (Bringmann et al., KIS 2009)
 - Flexics (Dzyuba et al., DMKD 2017),
 - CFTP (Boley et al., KDD 2012)



⇒ Exploiting CP for mining diverse set of patterns



Constraint programming

A generic framework for solving combinatorial problems

- A **declarative** description of the problem by a triplet (X, D, C) where
 - $X = \{x_1, \dots, x_n\}$ is finite set of **variables**
 - $D = \{D_1, \dots, D_n\}$ is finite set of **domains** (a.k.a possible values) of variables
 - $C = \{c_1, \dots, c_e\}$ is a set of **constraints** restricting the values of variables x_i
- **Resolution** = Enumeration + Filtering

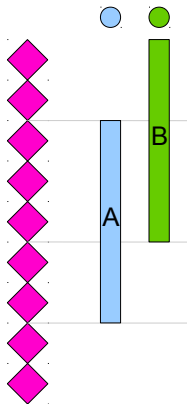
solution $\equiv$ assignments on X satisfying all constraints of C



Measuring redundancy

Exploiting similarity measure to compare pairs of patterns :

- $|A \cap B| / \min(|A|, |B|)$ (Overlap)
- $|A \cup B| - |A \cap B|$ (Hamming distance)
- $|A \cap B| / |A| \cdot |B|$ (Cosine similarity)
- $|A \cap B| / |A \cup B|$ (Jaccard similarity)





Definition (Hebrard et al., AAAI 2005)

Let $k \geq 2$ be an integer, and $\mathcal{S}$ be a set of solutions.

- The $\text{MAXDIVERSEKSET}(k)$ problem is the problem of finding a subset $\tilde{S}$ of $\mathcal{S}$ of size k such that for all $S \subseteq \mathcal{S}$, $|S| = k$,

$$\min_{\substack{s_1, s_2 \in \tilde{S} \\ s_1 \neq s_2}} J_V(s_1, s_2) \leq \min_{\substack{s_1, s_2 \in S \\ s_1 \neq s_2}} J_V(s_1, s_2).$$

- Given a set S of previously found solutions (a history), the $\text{MOSTDISTANT}(S)$ problem is the problem of finding a solution $\tilde{s} \in \mathcal{S}$ such that for all $s \in S$,

$$\min_{s' \in S} J_V(\tilde{s}, s') \leq \min_{s' \in S} J_V(s, s').$$

- $\text{MAXDIVERSEKSET}(k)$ can be seen as finding a set of k solutions minimizing the pairwise Jaccard values.
- MOSTDISTANT aims at finding the most distant solution from the history of solution ; can be seen as a greedy algorithm solving MAXDIVERSEKSET problem.



First approximation of the MOSTDISTANT problem (Hien et al., ECML-PKDD 2020 & Constraints 2024)

Definition (Maximum Diversity Constraint)

Let $\mathcal{H}$ be a history of patterns, j_{max} a bound on the maximum allowed Jaccard, and P an itemset. The maximum diversity constraint ensures that P is diverse w.r.t. $\mathcal{H}$ and J_{max} .

$$\text{div}(P, \mathcal{H}, j_{max}) \Leftrightarrow \forall H \in \mathcal{H}, J_V(P, H) \leq J_{max}$$

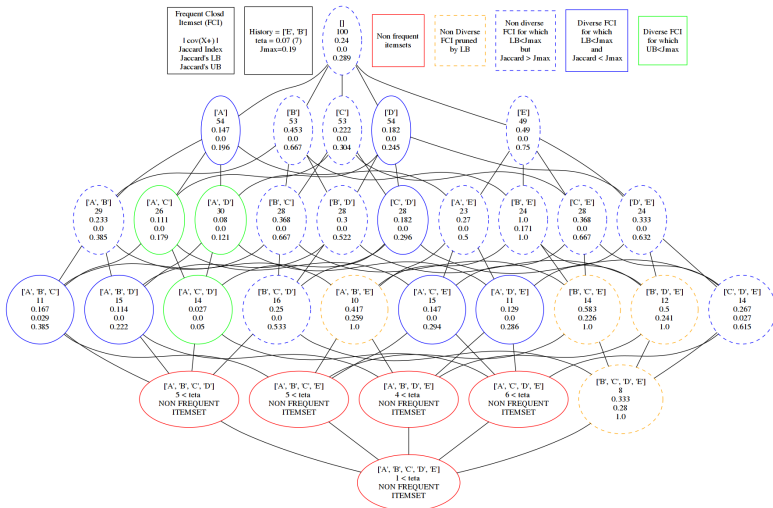
Idea: Push the Jaccard constraint during pattern discovery to prune non-diverse patterns.

Task : Given a history $\mathcal{H}$ of k pairwise diverse patterns (initially empty), the task is to mine new patterns P such that $\text{div}(P, \mathcal{H}, j_{max})$ is satisfied.



Anti-monotonicity of the maximum Jaccard constraint

The anti-monotonicity does not hold for the Maximum Jaccard constraint



- For $\mathcal{H} = \{BE\}$ and $J_{max} = 0.19$, $J_{acc}(AE, \mathcal{H}) = 0.27 \geq J_{max}$ whereas $J_{acc}(ACE, \mathcal{H}) = 0.147 \leq J_{max}$.



Relaxations of the Jaccard constraint

(i) **A lower bound** LB_J , which allows to prune non-diverse patterns

$$LB_J(H, P) = \begin{cases} \frac{\theta - |\mathcal{V}_H^{pr}(P)|}{|\mathcal{V}_D(H)| + |\mathcal{V}_H^{pr}(P)|} & \text{if } (|\mathcal{V}_H^{pr}(P)| < \theta) \\ 0 & \text{else} \end{cases}$$

(ii) **An upper bound** UB_J to find patterns ensuring diversity

$$UB_J(H, P) = \frac{|\mathcal{V}_D(H) \cap \mathcal{V}_D(P)|}{|\mathcal{V}_H^{pr}(P)| + \max\{\theta, |\mathcal{V}_D(H) \cap \mathcal{V}_D(P)|\}}$$



Jaccard lower and upper bounds

- **Monotonicity of LB_J :** Let $H \in \mathcal{H}$ be an itemset. For any two patterns $Q \subseteq P$, the relationship $LB_J(P, H) \geq LB_J(Q, H)$ holds.
 ⇒ If $LB_J(P, H) > J_{max} \Rightarrow J_V(P, H) > J_{max} \Rightarrow P$ **is not diverse**
 (any super-itemset $P' \supseteq P$ **will not be diverse**)
- **Anti-monotonicity of UB_J :** Let $H \in \mathcal{H}$ be an itemset. For any two patterns $P \subseteq Q$, the relationship $UB_J(P, H) \geq UB_J(Q, H)$ holds.
 ⇒ If $UB_J(P, H) \leq J_{max} \Rightarrow J_V(P, H) \leq J_{max} \Rightarrow P$ **is diverse**
 (any super-itemset $P' \supseteq P$ **will also be diverse**)
- A new global constraint $CLOSED DIVERSITY_{\mathcal{D}, \theta}(X, \mathcal{H}, J_{max})$ for mining pairwise diverse patterns that exploits the two previous relaxations.



CLOSED DIVERSITY $_{\mathcal{D}, \theta}(X, \mathcal{H}, J_{max})$ (A. Hien et al., Constraints 2024)

- Use a vector X of Boolean variables $(X_1, \dots, X_{|I|})$ for representing item sets
- X^+ will denote the set of present items,
- CLOSED DIVERSITY $_{\mathcal{D}, \theta}(X, \mathcal{H}, J_{max})$ holds if and only if :
 - (1) $freq(X^+) \geq \theta$ and X^+ is closed $\implies$ CLOSED PATTERNS
 - (2) X^+ is diverse, $\forall H \in \mathcal{H}, LB_J(X^+, H) \leq J_{max}$.

Two filtering rules : Let X_{Div} be the set of items filtered by (Rule #1)

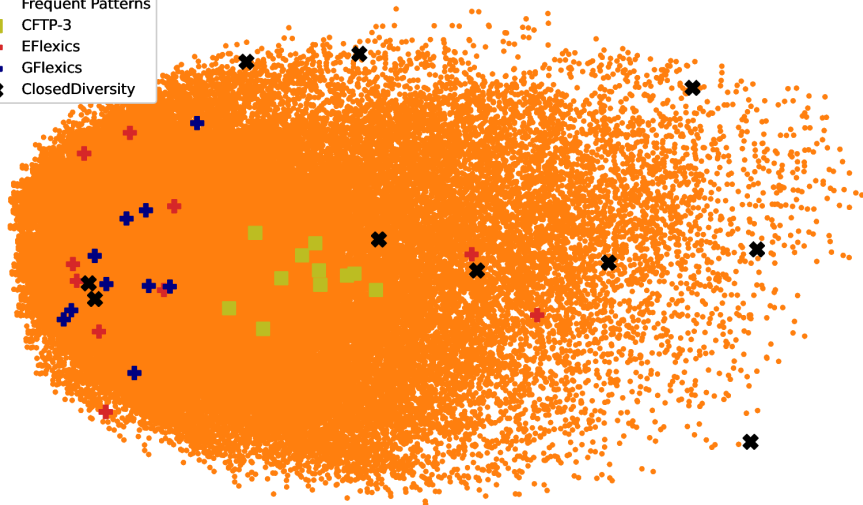
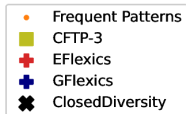
- 1 remove 1 from $dom(X_i)$ if $\exists H \in \mathcal{H}$ s.t. $LB_J(X^+ \cup \{i\}, H) > J_{max}$
- 2 remove 1 from $dom(X_i)$ if $\exists k \in X_{Div}$ s.t. $cover(X^+ \cup \{i\}) \subseteq cover(X^+ \cup \{k\})$, then $LB_J(X^+ \cup \{i\}, H) > LB_J(X^+ \cup \{k\}, H) > J_{max}$

Time complexity: $O(n \times (n \times m))$



Comparative Exploration of Extracted Pattern Landscapes

hepatitis - freq=30.0% - minsup=42





False positives

CLOSEDDIVERSITY ensures that $\forall H \in \mathcal{H}, LB_J(P, H) \leq J_{max}$

$$LB_J(P, H) \leq J_{max} : \begin{cases} J_{\mathcal{V}}(P, H) \leq J_{max} \\ J_{\mathcal{V}}(P, H) > J_{max} \end{cases}$$



False positives

CLOSEDDIVERSITY ensures that $\forall H \in \mathcal{H}, LB_J(P, H) \leq J_{max}$

$$LB_J(P, H) \leq J_{max} : \begin{cases} J_{\mathcal{V}}(P, H) \leq J_{max} \\ J_{\mathcal{V}}(P, H) > J_{max} \Rightarrow \text{false positive} \end{cases}$$



False positives

CLOSEDDIVERSITY ensures that $\forall H \in \mathcal{H}, LB_J(P, H) \leq J_{max}$

$$LB_J(P, H) \leq J_{max} : \begin{cases} J_V(P, H) \leq J_{max} \\ J_V(P, H) > J_{max} \Rightarrow \text{false positive} \end{cases}$$

Proposition : adding the Jaccard test before any update of $\mathcal{H}$

⇒ CLOSEDIV+JACCARD



- UCI datasets

- Comparisons:

- | | |
|----------------------|------------------------------------|
| (1) CLOSEDP | (6) PICKER |
| (2) CLOSEDP+JACCARD | (7) PATTERNSTEAM |
| (3) CLOSEDIV | (8) FLEXICS (EFLEXICS et GFLEXICS) |
| (4) CLOSEDIV+JACCARD | (9) GIBBS |
| (5) FULLCP | (10) CFTP |

- Implementation: Choco solver, timeout = 24 hours



Quality of diversification

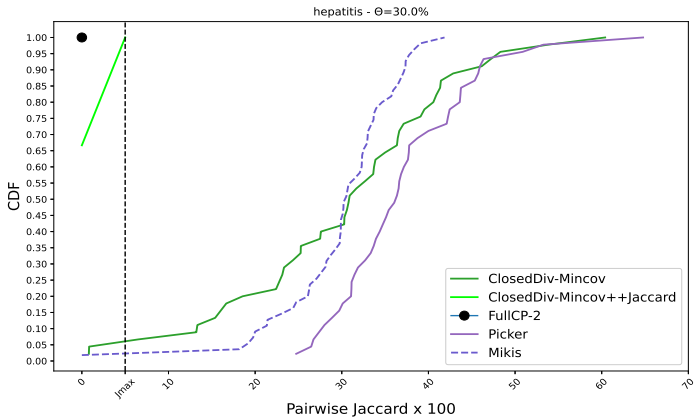
Let a set of patterns $\mathcal{H} = \{H_1, \dots, H_k\}$

$$CDF_{\mathcal{H}}(\tau) = \#\{(i, j) | J_{\mathcal{V}}(H_i, H_j) \leq \tau, 1 \leq i < j \leq k\} \cdot \frac{2}{k(k-1)}$$

► Cumulative distribution function on the Jaccard indices of each pair of patterns



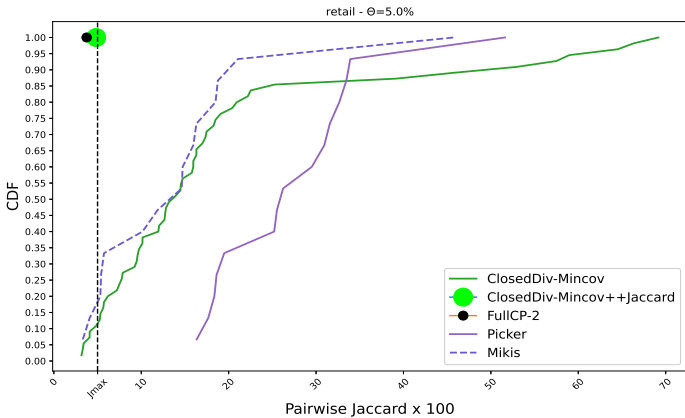
Quality of diversification : heuristic approaches



(a) HEPATITIS ($\theta = 30\%$)



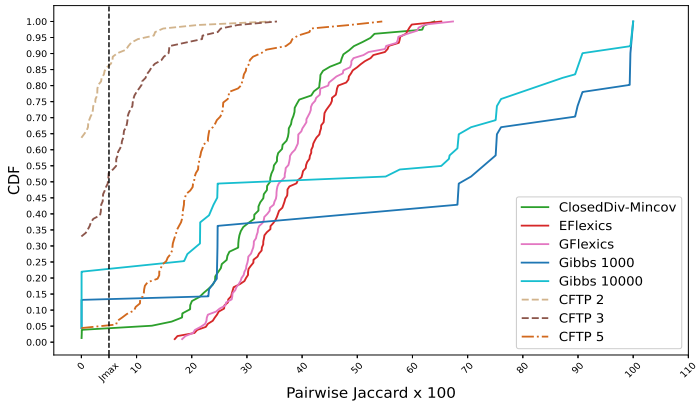
Quality of diversification : heuristic approaches



(b) RETAIL ($\theta = 5\%$)



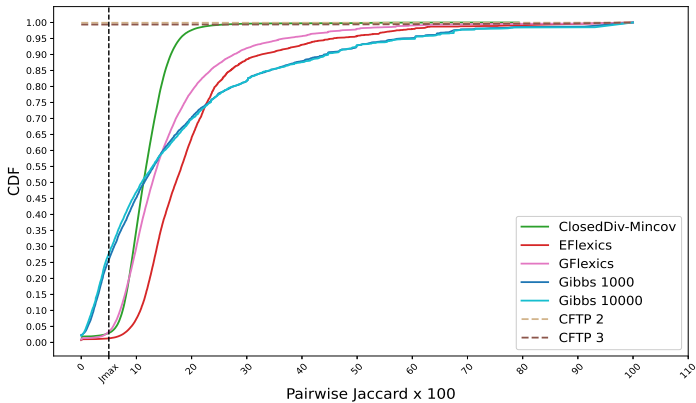
Quality of diversification : sampling approaches



(c) HEPATITIS ($\theta = 10\%$)



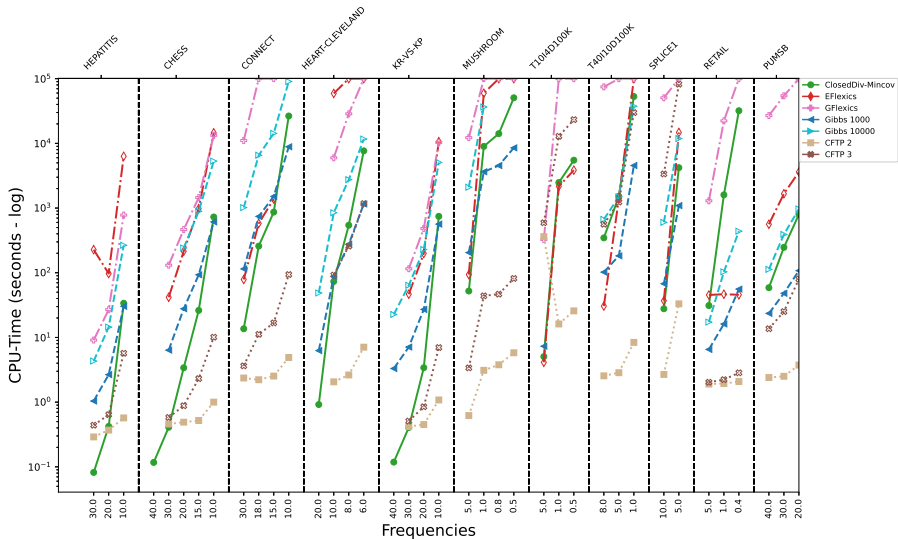
Quality of diversification : sampling approaches



(d) SPLICE1 ($\theta = 10\%$)

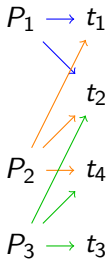


CPU times : sampling approaches

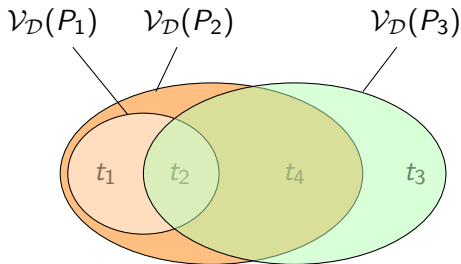




Jaccard index vs. Overlap



(e)



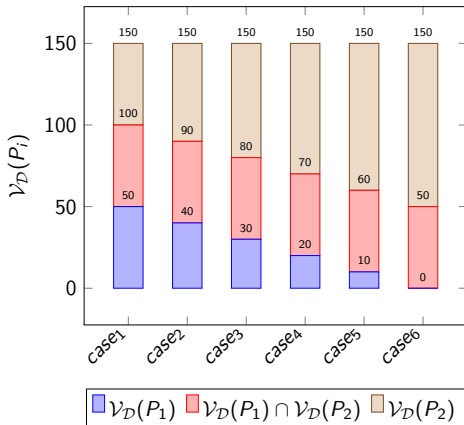
(f)

Pairwise comparison	Jaccard Index	Overlap Coefficient
P_1 vs. P_2	$\frac{ P_1 \cap P_2 }{ P_1 \cup P_2 } = \frac{2}{3} \approx 0.66$	$\frac{ P_1 \cap P_2 }{\min(P_1 , P_2)} = \frac{2}{2} = 1.0$
P_1 vs. P_3	$\frac{ P_1 \cap P_3 }{ P_1 \cup P_3 } = \frac{1}{4} = 0.25$	$\frac{ P_1 \cap P_3 }{\min(P_1 , P_3)} = \frac{1}{2} = 0.5$

Table: Comparison of Similarity Measures



Sensitivity analysis of the Overlap coefficient

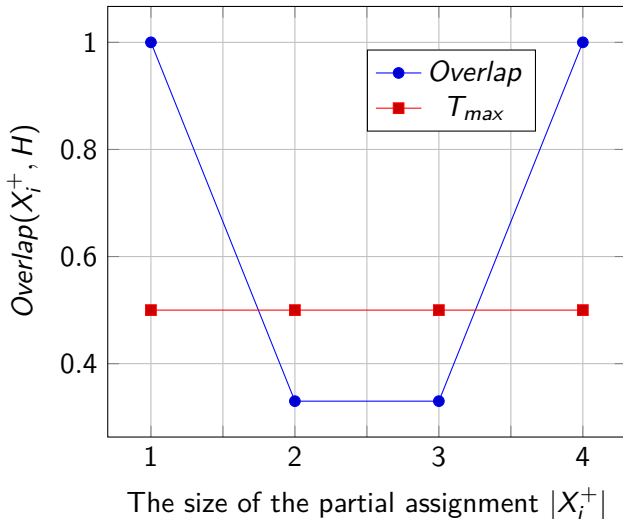


case#	$ V_D(P_1) $	$ V_D(P_2) $	(J)	(O)
case ₁	100	100	0.333	0.500
case ₂	110	90	0.333	0.556
case ₃	120	80	0.333	0.625
case ₄	130	70	0.333	0.714
case ₅	140	60	0.333	0.833
case ₆	150	50	0.333	1.0

By considering the proportion of the smaller set contained within the larger set, the Overlap coefficient offers a more informative and nuanced measure of set similarity.



Anti-monotonicity of the Overlap Coefficient



- For $\mathcal{H} = \{AE\}$, $X^+ = \{A\} \subseteq \{A, C\} \subseteq \{A, C, D\} \subseteq \{A, C, D, E\}$,
 $Overlap(A, \mathcal{H}) = 1 \geq T_{max}$ whereas $Overlap(AC, \mathcal{H}) = .33 \leq T_{max}$.



Relaxations of the overlap coefficient (M. Douad et al., APIN 2025)

(i) **A lower bound** which allows to prune non-diverse patterns

$$LB_{OC}(P, H) = \begin{cases} \frac{\theta - |\mathcal{V}_H^{pr}(P)|}{\min(|\mathcal{V}_D(P)|, |\mathcal{V}_D(H)|)} & \text{if } (\mathcal{V}_H^{pr}(P) < \theta) \\ 0 & \text{else} \end{cases}$$

(ii) **An upper bound** to find patterns ensuring diversity

$$UB_{OC}(P, H) = \min \left(\frac{|\mathcal{V}_D(P) \cap \mathcal{V}_D(H)|}{\theta}, 1 \right)$$



Overlap lower and upper bounds

- **Monotonicity of LB_{OC} :** Let $H \in \mathcal{H}$ be an itemset. For any two patterns $Q \subseteq P$, the relationship $LB_{OC}(P, H) \leq LB_{OC}(Q, H)$ holds.
 ⇒ If $LB_{OC}(P, H) > J_{max} \Rightarrow J_V(P, H) > J_{max} \Rightarrow P$ **is not diverse**
 (any super-itemset $P' \supseteq P$ **will not be diverse**)
- **Anti-monotonicity of UB_{OC} :** Let $H \in \mathcal{H}$ be an itemset. For any two patterns $P \subseteq Q$, the relationship $UB_{OC}(P, H) \geq UB_{OC}(Q, H)$ holds.
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 (any super-itemset $P' \supseteq P$ **will also be diverse**)
- A new global constraint $OverlapDiv_{\mathcal{D}, \theta}(X, \mathcal{H}, T_{max})$ for mining pairwise diverse patterns that exploits the two previous relaxations.



OverlapDiv $_{\mathcal{D},\theta}(X, \mathcal{H}, T_{max})$ (A. Douad et al., APIN 2025)

- Use a vector X of Boolean variables $(X_1, \dots, X_{|I|})$ for representing item sets
- X^+ will denote the set of present items,
- OverlapDiv $_{\mathcal{D},\theta}(X, \mathcal{H}, T_{max})$ holds if and only if :
 X^+ is diverse; $\forall H \in \mathcal{H}, LB_{OC}(X^+, H) \leq T_{max}$

Two filtering rules : Let X_{Div} be the set of items filtered by (Rule #1)

$$1 \notin Dom(X_i) \text{ iff } \begin{cases} \exists H \in \mathcal{H} \text{ s.t. } LB_{OC}(X^+ \cup \{i\}, H) > T_{max} & \text{(R1)} \\ \exists k \in X_{Div} \text{ s.t. } \mathcal{V}_{\mathcal{D}}(X^+ \cup \{i\}) \subseteq \mathcal{V}_{\mathcal{D}}(X^+ \cup \{k\}) & \text{(R2)} \end{cases}$$

Time complexity: $O(n \times (n \times m))$



A CP model for mining diverse closed and frequent itemsets

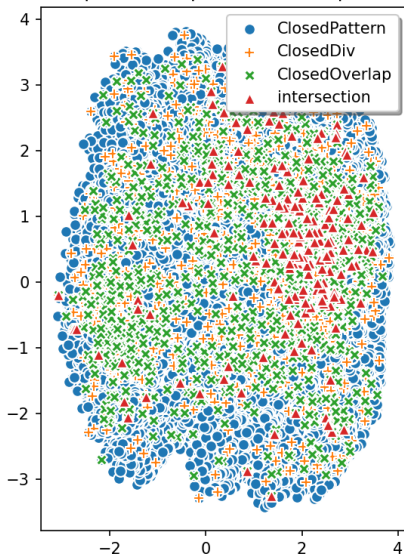
$$\text{ClosedOverlap}_{\mathcal{D},\theta}(X, c, T_{max}, \mathcal{H}) \equiv \begin{cases} \text{OverlapDiv}_{\mathcal{D},\theta}(X, T_{max}, \mathcal{H}) & (1) \\ \text{CoverSize}_{\mathcal{D},\theta}(X, c) \wedge c \geq \theta & (2) \\ \text{CoverClosure}_{\mathcal{D}}(X) & (3) \end{cases}$$

- **OverlapDiv**: Enforces diversity by controlling the overlap between the current itemset and the previously mined itemsets in $\mathcal{H}$.
- **CoverSize (schaus et al., 2017)**: Ensures frequency by requiring that the cover size of the itemset satisfies: $\text{Freq}_{\mathcal{D}}(X) \geq \theta$.
- **CoverClosure**: Guarantees closure by ensuring that the itemset is maximal w.r.t. the frequency measure, i.e., $\text{Clos}_{\text{freq}}(X) = X$.



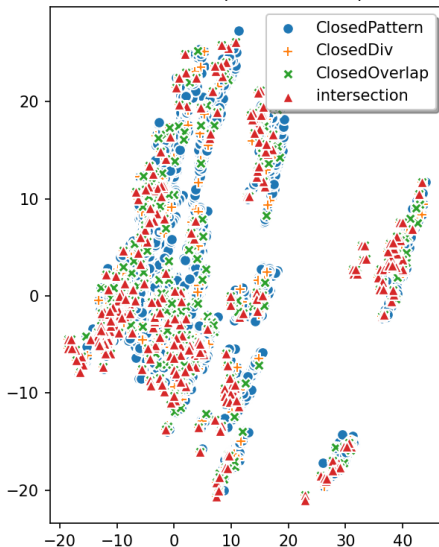
PCA of Extracted Pattern Landscapes

hepatitis, Freq: 10.0%, MinSup: 14



(g)

mushroom, Freq: 5.0%, MinSup: 407



(h)



Optimization model or mining k -diverse frequent itemsets

Both `ClosedOverlap` and `CLOSED DIVERSITY` are sensitive to order in which patterns are explored $\Rightarrow$ can significantly influence the output

Formulate the initial problem as an optimization one: $\Rightarrow$ select the best pattern set S^* of size k maximizing some quality measure ϕ over all subsets of $\mathcal{P}$

Joint entropy measure as a quality measure:

Definition (Joint entropy of an itemset)

Let $P = \{i_1, \dots, i_n\}$ be an itemset and $c = (c_1, \dots, c_n) \in \{0, 1\}^n$ a tuple of binary values. The joint entropy of P is given as

$$H(P) = - \sum_{c_i \in \{0,1\}^n} p(i_1 = c_1, \dots, i_n = c_n) \log_2 p(i_1 = c_1, \dots, i_n = c_n) \quad (1)$$



Join entropy-based formulation for minimizing k -pattern set redundancy

Each pattern $H_i \in \mathcal{H}$ is represented as a **binary vector**, within a $|\mathcal{D}| \times |\mathcal{H}|$ matrix, indicating the transactions where H_i is present.

$$\mathcal{H} = \{ABCD, AE, BE\}$$

$\{A, B, C, D\}$	$\{A, E\}$	$\{B, E\}$
0	0	1
0	0	0
0	0	0
1	1	1
1	0	0
0	0	0
0	1	0
0	1	0



Join entropy-based formulation for minimizing k -pattern set redundancy

The Entropy Computation exploits the binary vectors $c \in \{0, 1\}^k$ derived from the binary matrix:

$$H(\mathcal{H}) = - \sum_{c \in \{0,1\}^k} \frac{\text{freq}(H_1 = c_1, \dots, H_k = c_k)}{|\mathcal{D}|} \log_2 \left(\frac{\text{freq}(H_1 = c_1, \dots, H_k = c_k)}{|\mathcal{D}|} \right)$$

- Diversity reflects the variability or balance in the presence of patterns across transactions:



Join entropy-based formulation for minimizing k -pattern set redundancy

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- Diversity reflects the variability or balance in the presence of patterns across transactions:
 - Pattern $\{A, B, C, D\}$: Appears in 2 transactions (t_4, t_5)
 - Pattern $\{A, E\}$: Appears in 3 transactions (t_4, t_7, t_8)
 - Pattern $\{B, E\}$: Appears in 2 transactions (t_1, t_4)



Join entropy-based formulation for minimizing k -pattern set redundancy

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- Diversity reflects the variability or balance in the presence of patterns across transactions:
 - If all patterns appear in all transactions, diversity is **low** because there is no variability.
 - If each pattern appears in a unique set of transactions, diversity is **high**.



Join entropy-based formulation for minimizing k -pattern set redundancy

The Entropy Computation exploits the binary vectors $c \in \{0, 1\}^k$ derived from the binary matrix:

$$H(\mathcal{H}) = - \sum_{c \in \{0,1\}^k} \frac{\text{freq}(H_1 = c_1, \dots, H_k = c_k)}{|\mathcal{D}|} \log_2 \left(\frac{\text{freq}(H_1 = c_1, \dots, H_k = c_k)}{|\mathcal{D}|} \right)$$

- Unique bit codes in this matrix are: $\{001, 000, 111, 100, 010\}$
- Occurrences: 001: 1, 000: 3, 111: 1, 100: 1, 010: 2
- $P(001) = \frac{1}{8}$, $P(000) = \frac{3}{8}$, $P(111) = \frac{1}{8}$, $P(100) = \frac{1}{8}$, $P(010) = \frac{2}{8} = \frac{1}{4}$
- The joint entropy H is:

$$\begin{aligned} H(\mathcal{H}) &= - \left(\frac{1}{8} \log_2 \frac{1}{8} + \frac{3}{8} \log_2 \frac{3}{8} + \frac{1}{8} \log_2 \frac{1}{8} + \frac{1}{8} \log_2 \frac{1}{8} + \frac{2}{8} \log_2 \frac{2}{8} \right) \\ &= \mathbf{2.1556 \text{ bits}} \end{aligned}$$



Join entropy-based formulation for minimizing k -pattern set redundancy

- Note that the joint entropy is always between 0 and k .
- It is maximal when patterns are fully uncorrelated and uniformly distributed.
- For 3 binary variables, there are $2^3 = 8$ possible combinations.
- If each occurs exactly once, then:

$$\begin{aligned} H(P_1, P_2, P_3) &= - \left(\sum_{i=1}^8 \frac{1}{8} \log_2 \frac{1}{8} \right) \\ &= - \left(8 \times \frac{1}{8} \times (-3) \right) = \mathbf{3 \text{ bits}} \end{aligned}$$



Join entropy-based formulation for minimizing k -pattern set redundancy

A Greedy Optimization Approach: Iteratively integrates patterns with the highest entropy contribution.

{A B C D}	{A E}	{B E}
0	0	1
0	0	0
0	0	0
1	1	1
1	0	0
0	0	0
0	1	0
0	1	0

Joint Entropy = 2.1556

→

{E}	{A E}	{B E}
1	0	1
1	0	0
0	0	0
1	1	1
0	0	0
0	0	0
1	1	0
1	1	0

Joint Entropy = 2.1556

→

{A B C D}	{E}	{B E}
0	1	1
0	1	0
0	0	0
1	1	1
1	0	0
0	0	0
0	1	0
0	1	0

Joint Entropy = 2.1556

→

{A B C D}	{A E}	{E}
0	0	1
0	0	1
0	0	0
1	1	1
1	0	0
0	0	0
0	1	1
0	1	1

Joint Entropy = **2.25**

Figure: Maximizing Joint Entropy: finding the best k -pattern set through candidate itemset replacement



CPU times: CLOSED DIVERSITY vs. ClosedOverlap

Dataset $\mathcal{T} \times \mathcal{I}$	$\theta\%$	CLOSED DIVERSITY		ClosedOverlap	
		CPU-Time (sec)	# Patterns	CPU-Time (sec)	# Patterns
CHESS 75 x 3196 49.33%	20	1.57	64	0.35	56
	15	12.36	237	1.00	224
	10	408.19	1621	15.62	1597
KR-VS-KP 73 x 3196 49.32 %	30	0.24	13	0.11	10
	20	1.41	63	0.35	57
	10	368.50	1608	15.30	1570
CONNECT 129 x 67557 33.33%	30	6.92	18	0.59	11
	18	134.91	140	5.76	104
	15	492.53	296	17.35	251
HEART-CLEVELAND 95 x 296 47.37%	10	40.05	1469	2.26	1394
	8	313.23	4760	12.98	4814
	6	4630.88	20489	281.19	22050
SPICE1 287 x 3190 20.91%	10	11.30	412	2.04	374
	5	2316.70	7919	112.46	5763
	2	TO	NA	112.46	5763
MUSHROOM 112 x 8124 18.75%	5	23.65	547	1.96	551
	1	5172.93	9934	202.43	9323
	0.5	27656.11	23930	1073.44	21974

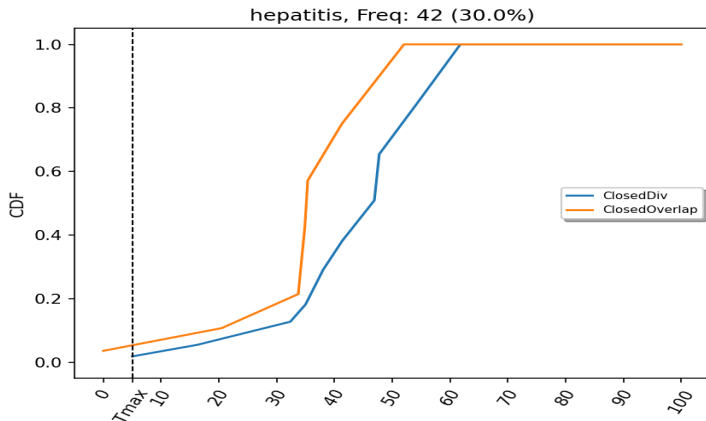


CPU times: CLOSED DIVERSITY vs. ClosedOverlap

Dataset $\mathcal{T} \times \mathcal{I}$	$\theta\%$	CLOSED DIVERSITY		ClosedOverlap	
		CPU-Time (sec)	# Patterns	CPU-Time (sec)	# Patterns
T40I10D100K 942 × 100000 4.20%	8	196.77	124	44.14	114
	5	880.13	283	200.19	271
	1	31213.08	7216	3523.54	5924
PUMSB 2113 × 49046 3.50%	40	31.33	3	2.67	3
	30	126.25	13	10.41	11
	20	431.91	38	35.04	39
T10I4D100K 870 × 100000 1.16%	5	2.26	10	0.62	10
	1	1455.39	359	305.70	351
	0.5	2910.10	606	645.35	595
RETAIL 16470 × 88162 0.06%	5	15.94	11	4.49	11
	1	864.30	104	206.57	103
	0.4	12489.13	514	3500.13	503



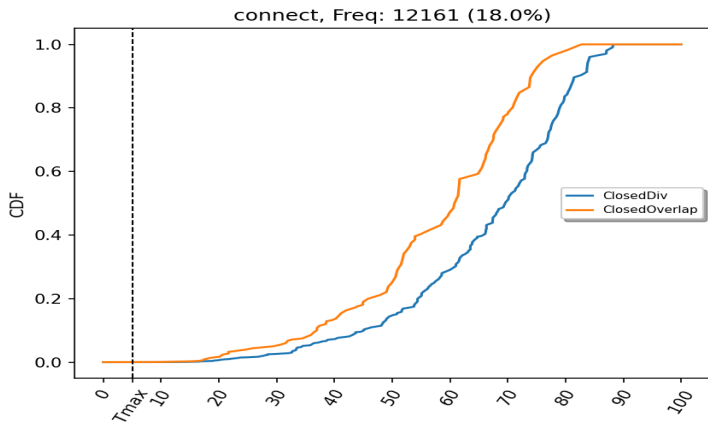
Diversity : CLOSED DIVERSITY vs. ClosedOverlap



(a) HEPATITIS ($\theta = 30\%$)



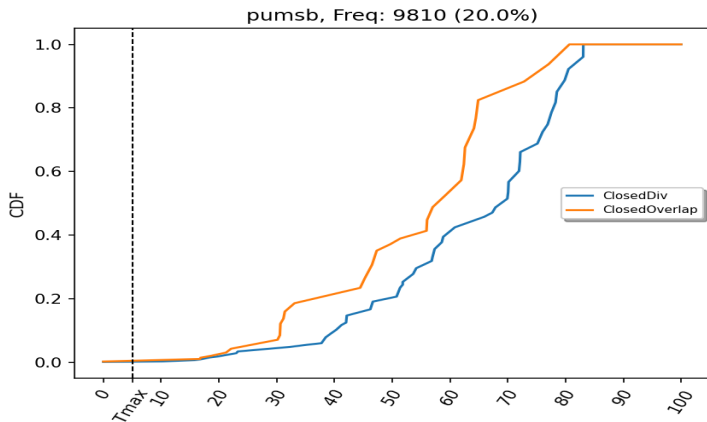
Diversity : CLOSED DIVERSITY vs. ClosedOverlap



(b) CONNECT ($\theta = 18\%$)



Diversity : CLOSED DIVERSITY vs. ClosedOverlap



(c) PUMSB ($\theta = 20\%$)



Search strategies for MOSTDISTANT problem (M. Vavrille et al., 2023)

CLOSEDIVERSITY and ClosedOverlap global constraints work at the **propagation level** of the solver

Idea: Define dedicated new **search strategies** to orient the search towards diverse patterns $\Rightarrow$ no modification of the model nor the solver's inner structure

Task : Given a history $\mathcal{H}$ of solutions (initially empty), the task is to mine new patterns P ensuring a minimum diversity according to a some **History score**



ORIENTED DETERMINISTIC strategy

A vector X of Boolean variables ($X_1, \dots, X_{|I|}$) for representing a pattern P

Definition (History score)

Given a pattern P being constructed, an item i whose associated variables X_i is uninstantiated, and a history $\mathcal{H}$ of solutions, we define the history score as

$$\max_{H \in \mathcal{H}} J_V(P \cup \{i\}, H)$$

The history score ensures a **minimum diversity** between $P \cup \{i\}$ and solutions of $\mathcal{H}$

Branching heuristic : select the next uninstantiated variable $X_{i_{min}}$ that minimizes the score (a low Jaccard means a good diversity):

$$i_{min} \leftarrow \arg \min_{\substack{1 \leq i \leq |I| \\ |\mathcal{D}(X_i)|=2}} \max_{H \in \mathcal{H}} J_V(P \cup \{i\}, H)$$

➡ The search may be stuck in a local optimum, i.e. solutions will be searched repeatedly in the same space.



Select the next uninstantiated variable according to some [weight distribution](#)

Definition (variable weight)

Let W be an array of size $|\mathcal{I}|$. Given a pattern P being constructed, an item i whose associated variables X_i is uninstantiated, and a history $\mathcal{H}$ of solutions, we define the weight of X_i as

$$W[i] = \frac{1}{\max_{H \in \mathcal{H}} J_{\mathcal{V}}(P \cup \{i\}, H) + \epsilon}$$

Instantiated variables are given a weight of 0

To bias the random distribution towards small values, we have to invert the computed history score

Branching heuristic : the next uninstantiated variable is randomly chosen with respect to the weights in W



Choco-Mining: A Java Library For Itemset Mining (Vernerey et al. JOSS 2023)

- Une nouvelle bibliothèque Choco-Mining¹ qui regroupe un ensemble de contraintes pour la fouille de motifs.
- Elle permet à un utilisateur intéressé de facilement réutiliser ces contraintes dans ses propres projets.

¹<https://gitlab.com/chaver/choco-mining>



Choco-Mining: A Java Library For Itemset Mining

(Vernerey et al. JOSS 2023)





A new generic solution for mining diverse patterns

- Exploiting relaxations and search strategies
- Other diversity measures (entropy) ?

Exploiting Jaccard index within search strategies show improved diversity compared to CLOSED DIVERSITY