

From-Below Approximations in Boolean Matrix Factorization: Geometry and New Algorithm II

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INVESTMENTS IN EDUCATION DEVELOPMENT

List of Algorithms for Boolean Matrix Factorization (BMF)

- Algorithm 1
- GreConD
- ASSO
- Tiling (LTM algorithm)
- PANDA
- HYPER (HYPER+)

ASSO Algorithm in Short

- Author: Pauli Miettinen from Max-Planck-Institut für Informatik, originally from Finland.
- First introduced in: The Discrete Basis Problem (2006), MSC thesis as Association algorithm.
- Probably currently the most discussed BMF algorithm in the data mining literature.

ASSO Algorithm Pseudocode

Input: A matrix $C \in \{0, 1\}^{n \times m}$ for data, a positive integer k , a threshold value $\tau \in [0, 1]$, and real-valued weights w^+ and w^- .

for $i = 1, \dots, m$ **do**

 Construct matrix \mathcal{A} row by row

$$a_i \leftarrow (1(c(i \Rightarrow j, C) \geq \tau))_{j=1}^m$$

end

\mathcal{B} and \mathcal{S} are empty matrices

$$\mathcal{B} \leftarrow [], \mathcal{S} \leftarrow []$$

Select the k basis vectors from \mathcal{A}

for $l = 1, \dots, k$ **do**

$$(a_i, s) \leftarrow \max_{a_i, s \in \{0, 1\}^{n \times 1}} \text{cover}\left(\begin{bmatrix} \mathcal{B} \\ a_i \end{bmatrix}, [\mathcal{S} \ s], C, w^+, w^-\right)$$

$$\mathcal{B} \leftarrow \begin{bmatrix} \mathcal{B} \\ a_i \end{bmatrix}, \mathcal{S} \leftarrow [\mathcal{S} \ s]$$

end

return \mathcal{B} a \mathcal{S}

ASSO Continuation

- $cover(\mathcal{B}, \mathcal{S}, \mathcal{C}, w^+, w^-)$ is equal to:

$$w^+ |\{(i, j) : c_{ij} = 1, (\mathcal{S} \circ \mathcal{B})_{ij} = 1\}| - w^- |\{(i, j) : c_{ij} = 0, (\mathcal{S} \circ \mathcal{B})_{ij} = 1\}|.$$

- Unfortunately it is more complicated.
- Vector s is compute from a_i in greedy manner: $cover$ function is compute separately for each row of input matrix C , $s_j = 1$ if value of $cover > 0$ for row C_j .
- Take to account covered part (not mentioned in pseudocode).
- $cover$ function does not count covered part.

Example (ASSO Algorithm, Miettinen's implementation - First Factor)

$$k = 3, \tau = 0.9, w^+ = 1, w^- = 1$$

$$C = \begin{pmatrix} 0 & 0 & 1 & 0 \\ 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 0 \end{pmatrix}, A = \begin{pmatrix} 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 1 & 1 & 1 \end{pmatrix}, \text{covered} = \begin{pmatrix} 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \end{pmatrix}$$

cover for the first row: $-2 + 4 + 2 + 0 = 6$

cover for the second row: $0 + 2 + 2 + 2 = 6$

cover for the third row: $1 + 1 + 1 + 1 = 4$

cover for the fourth row: $-1 + 3 + 3 + 1 = \mathbf{7}$

$$\begin{pmatrix} 0 \\ 1 \\ 1 \\ 1 \end{pmatrix} \circ (0 \ 1 \ 1 \ 1) = \begin{pmatrix} 0 & 0 & 0 & 0 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \end{pmatrix}$$

Example (ASSO Algorithm, Miettinen's implementation - Second Factor)

$$k = 3, \tau = 0.9, w^+ = 1, w^- = 1$$

$$C = \begin{pmatrix} 0 & 0 & 1 & 0 \\ 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 0 \end{pmatrix}, A = \begin{pmatrix} 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 1 & 1 & 1 \end{pmatrix}, \text{covered} = \begin{pmatrix} 0 & 0 & 0 & 0 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \end{pmatrix}$$

cover for the first row: $-2 + 1 - 1 - 1 = 1$

cover for the second row: $0 + 0 + 0 + 0 = 0$

cover for the third row: $1 + 0 + 0 + 0 = \mathbf{1}$

cover for the fourth row: $-1 + 0 + 0 + 0 = 0$

$$\begin{pmatrix} 0 & 1 \\ 1 & 0 \\ 1 & 0 \\ 1 & 0 \end{pmatrix} \circ \begin{pmatrix} 0 & 1 & 1 & 1 \\ 0 & 0 & 1 & 0 \end{pmatrix} = \begin{pmatrix} 0 & 0 & 1 & 0 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \end{pmatrix}$$

Example (ASSO Algorithm, Miettinen's implementation - Third Factor)

$$k = 3, \tau = 0.9, w^+ = 1, w^- = 1$$

$$C = \begin{pmatrix} 0 & 0 & 1 & 0 \\ 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 0 \end{pmatrix}, A = \begin{pmatrix} 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 1 & 1 & 1 \end{pmatrix}, \text{covered} = \begin{pmatrix} 0 & 0 & 1 & 0 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \end{pmatrix}$$

cover for the first row: $-3 + 1 + 0 - 1 = \mathbf{1}$

cover for the second row: $-1 + 0 + 0 + 0 = 0$

cover for the third row: $0 + 0 + 0 + 0 = 0$

cover for the fourth row: $-2 + 0 + 0 + 0 = 0$

$$\begin{pmatrix} 0 & 1 & 0 \\ 1 & 0 & 1 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \end{pmatrix} \circ \begin{pmatrix} 0 & 1 & 1 & 1 \\ 0 & 0 & 1 & 0 \\ 1 & 1 & 1 & 1 \end{pmatrix} = \begin{pmatrix} 0 & 0 & 1 & 0 \\ 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 0 & 1 & 1 & 1 \end{pmatrix}$$

ASSO Algorithm Remark

There exist several variants of ASSO:

- ASSO + transpose
- ASSO + clustering
- ASSO + exhaustive search
- ASSO + iterative search

- One specific feature: **There is no way how to remove E_o !**
- Empirical time complexity: 4-5x slower than GreConD.

Tiling Algorithm in Short

- First introduced in: Tiling Databases (2004).
- Originally called LTM algorithm.
- Base on idea from CHARM, LPMiner and BAMBOO algorithm.
- Introduced notion of tile $\tau(I, D) = \{(tid, i) \in cover(I, D), i \in I\}$.
- Tiles can be seen as rectangles in binary data.

Tiling Algorithm Pseudocode

Input: Binary database D , minimal size of tile δ , itemset I initially called with $I = \emptyset$.

$T \leftarrow \emptyset$

$D \leftarrow \text{PRUNE}(D, \delta, I)$

forall the i occurring in D do

if $|\text{cover}(\{i\})| \cdot (|I| + 1) \geq \delta$ **then**

$T \leftarrow T \cup \tau(I \cup \{i\})$

end

$D^i \leftarrow \emptyset$

forall the j occurring in D such that $j > i$ do

$C \leftarrow \text{cover}(\{i\}) \cap \text{cover}(\{j\})$

 Add (j, C) to D^i

 Recursively compute for $I \cup \{i\}$

end

end

PRUNE Procedure Pseudocode

Input: D, δ, I .

repeat

forall the i occurring in D **do**

if $UB_{I \cup \{i\}} < \delta$ **then**

 | Remove i from D

end

forall the $tid \in cover(\{i\})$ **do**

if $size(tid) < ML_{I \cup \{i\}}$ **then**

 | Remove tid from $cover(\{i\})$

end

end

end

until *nothing change*

Explain Prune Procedure

$$UB_{I \cup \{i\}} = \max\{(|I| + l) \cdot \text{supp}_{\geq l}(i, D^I) \mid l \in \{1, 2, \dots\}\}$$

- If $UB_{I \cup \{i\}} < \delta$ then remove item (column) i from D .
- Upper bound of the largest possible tile containing $I \cup \{i\}$.

$$ML_{I \cup \{i\}} = \min\{l \mid l \cdot \text{supp}_{\geq l}(i, D^I) \geq \delta\}$$

- If $\text{size}(tid) < ML_{I \cup \{i\}}$ then remove transaction (row) tid from D .
- Minimal size of transaction containing i , that can still generate a tile with area at least δ .

Tiling Algorithm

- Produces all tiles with support $> \delta$.
- Easily can be change to find top- k largest tiles.
- Can be modified to solve (approximately) Minimum tiling problem (completely cover input database with the smallest number of tiles).
- We must modify algorithm cost function - it must take into account already covered area.
- In each step is extracted tile with the highest still uncovered part.
- Step = run whole Tiling algorithm!

Tiling Algorithm Remarks

- Modification of Tiling algorithm for BMF, is very slow!
- Produce the same results as Algorithm 1 (in special settings, fix order of tiles).
- In normal case there is more tiles with the same uncovered part - leads to the clash of decision function for selecting tiles.

PANDA Algorithm in Short

- First introduced in: Mining Top-K Patterns from Binary Datasets in Presence of Noise.
- Basic idea: Select noise-free pattern and extend this pattern with some noise.
- Notion of pattern $P = \langle P_T, P_I \rangle$ in binary database, where $P_T \in \{0, 1\}^n$, $P_I \in \{0, 1\}^m$.
- Pattern can be seen as rectangle in binary database.
- Compute the set of patterns Π with the smallest cost.
- Cost function of PANDA algorithm for set of patterns Π and database \mathcal{D} :

$$\gamma(\Pi, \mathcal{D}) = \sum_{P \in \Pi} \gamma_P(P_T, P_I) + \gamma_N(N)$$

where $\gamma_P(P_T, P_I) = |P_T| + |P_I|$ and $\gamma_N(N) = |N|$.

PANDA Algorithm Pseudocode

Input: \mathcal{D} binary database, k number of extracted patterns

$\Pi \leftarrow \emptyset$

$\mathcal{D}_{\mathcal{R}} \leftarrow \mathcal{D}$

for $i = 1 \dots k$ **do**

$C, E \leftarrow \text{FIND-CORE}(\mathcal{D}_{\mathcal{R}}, \Pi, \mathcal{D})$

$C^+ \leftarrow \text{EXTEND-CORE}(C, E, \Pi, \mathcal{D})$

if $\gamma(\Pi, \mathcal{D}) \leq \gamma(\Pi \cup C^+, \mathcal{D})$ **then**

 | break

end

$\Pi \leftarrow \Pi \cup C^+$

$\mathcal{D}_{\mathcal{R}}(i, j) \leftarrow 0 \forall i, j \text{ s.t. } C_T^+(i) = 1 \wedge C_I^+(i) = 1$

end

return Π

Core Pattern Discovery, FIND-CORE Procedure

Input: $\mathcal{D}_{\mathcal{R}}, \Pi, \mathcal{D}$

$E \leftarrow \emptyset$

$S = \{s_1, \dots, s_m\} \leftarrow \text{SORT-ITEMS-IN-DB}(\mathcal{D}_{\mathcal{R}})$

$C \leftarrow \langle C_T = 0^n, C_I = 0^m \rangle$

$C_I(s_1) \leftarrow 1$

$C_T(i) \leftarrow 1 \forall i \text{ s.t. } \mathcal{D}_{\mathcal{R}}(i, s_1) = 1$

for $h = 2, \dots, m$ **do**

$C^* \leftarrow C$

$C_I^*(s_h) \leftarrow 1$

$C_T^*(i) \leftarrow 0 \forall i \text{ s.t. } \mathcal{D}_{\mathcal{R}}(i, s_h) = 0$

if $\gamma(\Pi \cup C^*, \mathcal{D}) \leq \gamma(\Pi \cup C, \mathcal{D})$ **then**

$C \leftarrow C^*$

else

$E.\text{append}(s_h)$

end

end

return C, E

Extension of Core Pattern, EXTEND-CORE Procedure

Input: C, E, Π, \mathcal{D}

```
while  $E \neq \emptyset$  do  
   $e \leftarrow E.\text{pop}()$   
   $C^* \leftarrow C$   
   $C_I^*(e) \leftarrow 1$   
  if  $\gamma(\Pi \cup C^*, \mathcal{D}) \leq \gamma(\Pi \cup C, \mathcal{D})$  then  
     $C \leftarrow C^*$   
  end  
  for  $i = 1, \dots, n$  s.t.  $C_T^*(i) = 0$  do  
     $C^* \leftarrow C$   
     $C_T^*(i) \leftarrow 1$   
    if  $\gamma(\Pi \cup C^*, \mathcal{D}) \leq \gamma(\Pi \cup C, \mathcal{D})$  then  
       $C \leftarrow C^*$   
    end  
  end  
end  
return  $C$ 
```

PANDA Remarks

- Sorting items in database:
 - Frequency (best results)
 - Correlation
 - Prefix-tree
 - Couples frequency
- Empirical time complexity: 5x slower than GreConD.

HYPER Algorithm

- First introduced in: Summarizing Transactional Databases with Overlapped Hyperrectangles (2011).
- Notion of hyperrectangle $H = T \times I = \{(i, j) : i \in T, j \in I\}$.
- T is subset of all transactions and I is subset of all items.
- Hyperrectangle can be seen as rectangle in binary database.
- Price of hyperrectangle H_i :

$$\gamma(H_i) = \frac{|T_i| + |I_i|}{|T_i \times I_i \setminus R|}$$

where R is already covered part of input database.

HYPER Algorithm Pseudocode

Input: DB binary database, C_α set of frequent itemsets with minimal support α union with singleton sets of itemsets (sets with only one item)

$R \leftarrow \emptyset$

$CBD \leftarrow \emptyset$

while $R \neq DB$ **do**

forall the $H_i \in C_\alpha$ **do**

$X_i \leftarrow \text{FIND-HYPER}(H_i, R)$

end

$H' \leftarrow \min_{X_i} \gamma(X_i)$

$R \leftarrow R \cup H'$

end

return CBD

Procedure for Finding Sub-hyperrectangle with Cheapest Price

Input: Hyperrectangle H , already covered part of input database R

forall the $S = \{t_j\} \times I_i \subseteq H$ **do**

 | calculate the number of uncovered DB elements in S , $|S \setminus R|$

end

sort S according to $|S \setminus R|$ and put in U

$H' \leftarrow$ first hyperrectangle S popped from U

while $U \neq \emptyset$ **do**

 | pop single transaction hyperrectangle S from U

if $\gamma(H' \cup S) > \gamma(H')$ **then**

 | break

else

 | $H' \leftarrow H' \cup S$

end

end

return H'

Example (HYPER Finding Sub-hyperrectangle with Cheapest Price)

	i_2	i_4	i_5	i_7		i_2	i_4	i_5	i_7
t_1	1	1	1	1	t_1	0	1	0	1
t_3	1	1	1	1	t_3	1	0	1	1
$H = t_4$	1	1	1	1	$R = t_4$	0	0	0	0
t_6	1	1	1	1	t_6	0	1	0	1
t_8	1	1	1	1	t_8	0	0	0	0
t_9	1	1	1	1	t_9	1	0	1	1

Sort transaction according to size of their uncovered part: $t_4, t_8, t_1, t_6, t_3, t_9$.

$$H' = t_4 \times I, \gamma(H') = \frac{4+1}{4} = 1.25$$

$$\text{add } t_8 \times I \text{ to } H', \gamma(H') = \frac{4+2}{4+4} = 0.75 \text{ decrease}$$

$$\text{add } t_1 \times I \text{ to } H', \gamma(H') = \frac{4+3}{4+4+4-2} = 0.70 \text{ decrease}$$

$$\text{add } t_6 \times I \text{ to } H', \gamma(H') = \frac{4+4}{4+4+4+4-2-2} = 0.67 \text{ decrease}$$

$$\text{add } t_3 \times I \text{ to } H', \gamma(H') = \frac{4+5}{4+4+4+4+4-2-2-3} = 0.69 \text{ increase - stop}$$

Final hyperrectangle $H' = t_4, t_8, t_1, t_6 \times I$.

HYPER Algorithm Remark

- In basic setting C_α is from Apriori algorithm.
- Variants of HYPER algorithm:
 - Maximal frequent itemset.
 - Frequent closed itemset (best results).
- Prune technique can accelerate HYPER algorithm.
- Empirical time complexity: it depends on size C_α for “reasonable” size is 4-5x slower than GreConD.
- Exact factorization (there is no E_o error BMF).
- HYPER+ approximate factorization (merges hyperrectangles, the merged ones contain 0s).
- HYPER+ is post-processing algorithm.

HYPER+ Algorithm Pseudocode

Input: Binary database DB , set of hyperrectangles CBD , final number of hyperrectangles β

$SCDB \leftarrow CBD$

while $\frac{|SCDB^C \setminus DB|}{|DB|} \leq \beta$ **do**

find $H_i, H_j \in SCDB$ whose merge is within the false positive budget

$$\frac{|(SCDB \setminus \{H_i, H_j\} \cup \{H_i \oplus H_j\})^C \setminus DB|}{|DB|} \leq \beta$$

and produce the maximum saving-false positive ratio

$$\frac{|T_i| + |T_j| + |I_i| + |I_j| - |T_i \cup T_j| - |I_i \cup I_j|}{|(H_i \oplus H_j) \setminus SCDB^C|}$$

$SCBD \leftarrow SCBD \setminus \{H_i, H_j\}$

$SCBD \leftarrow SCBD \cup \{H_i \oplus H_j\}$

end

return $SCDB$

Noise in Binary Data

- Big topic in Data Mining in general.
- Current consideration regarding noise in binary data seem not mature (ad hoc).
- For example: in work about ASSO algorithm - 40% of noise (means practically new data).
- It is reasonable to take it as some kind of error.

Experiments Scenario

- We use various random and real datasets.
- We compare GreEss (pessimistic estimate), GreConD, ASSO (basic), Tiling, PANDA, and HYPER (frequent closed itemsets) algorithm.
- Error:

$E(I, A \circ B) = E_u(I, A \circ B) + E_o(I, A \circ B)$, where

$$E_u(I, A \circ B) = |\{\langle i, j \rangle; I_{ij} = 1, (A \circ B)_{ij} = 0\}|,$$

$$E_o(I, A \circ B) = |\{\langle i, j \rangle; I_{ij} = 0, (A \circ B)_{ij} = 1\}|.$$

- We are interested in from-below approximation of binary data.
- For each algorithm we compute coverage quality for each obtained factor.
- Coverage quality:

$$c = 1 - \frac{E_u(I, A \circ B)}{\|I\|}$$

Synthetic Data - Randomly Generated Matrices

- It is senseless compute Boolean matrix factorization on fully random data (there are no rational factors).
- We compute random binary matrices $n \times k$ and $k \times m$.
- Boolean product of these matrices represent our synthetic data with size $n \times m$.
- Such matrix is decomposable with (at most) k factors.

Synthetic Data

dataset	size	basis	dens A	dens B
Set 1	300×100	20	0.10	0.10
Set 2	500×250	20	0.05	0.05
Set 3	500×250	20	0.10	0.05
Set 4	500×250	30	0.12	0.12
Set 5	1000×500	50	0.10	0.10
Set 6	10000×1000	50	0.10	0.10

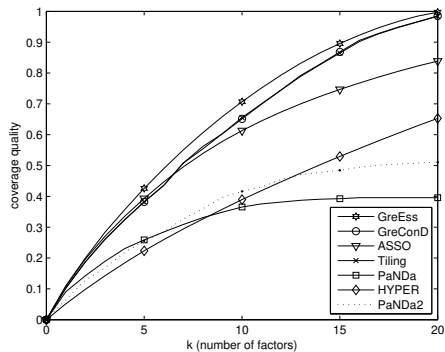
Table: Synthetic datasets $I = A \circ B$

Statistics on Synthetic Data

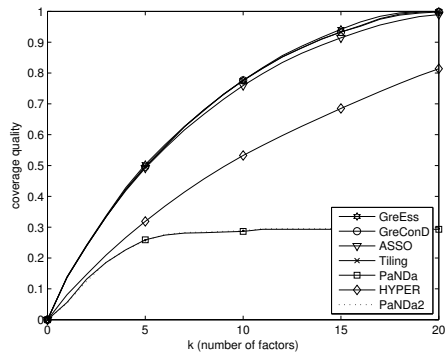
dataset	avg $\ I\ $	avg $\ \mathcal{E}(I)\ $	avg $\ \mathcal{E}(I)\ /\ I\ $
Set 1	5039	2764	0.549
Set 2	11966	6412	0.536
Set 3	22841	5207	0.228
Set 4	44027	8894	0.202
Set 5	195990	34005	0.174
Set 6	3907433	47359	0.012

Table: Essential part of I (synthetic data)

Results for Synthetic Data I.



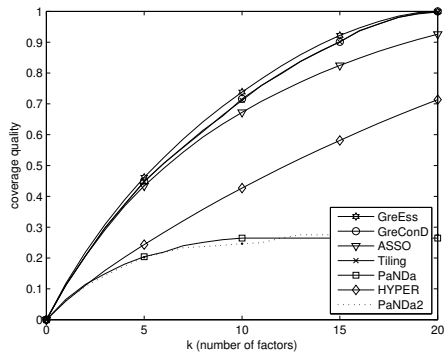
(a) Set 1



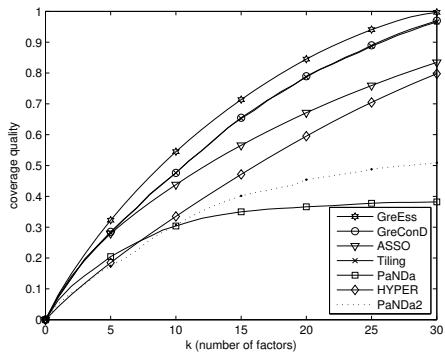
(b) Set 2

Figure: Coverage by the first k factors (synthetic data)

Results for Synthetic Data II.



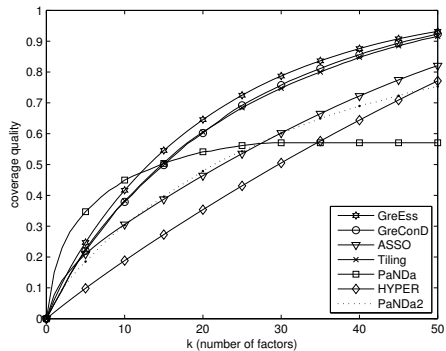
(a) Set 3



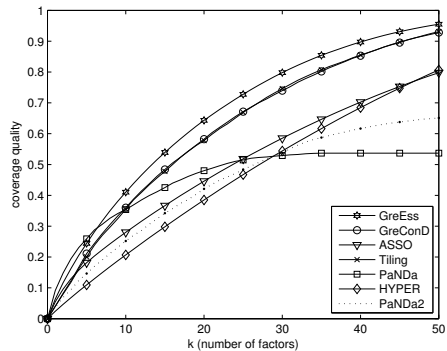
(b) Set 4

Figure: Coverage by the first k factors (synthetic data)

Results for Synthetic Data III.



(a) Set 5



(b) Set 6

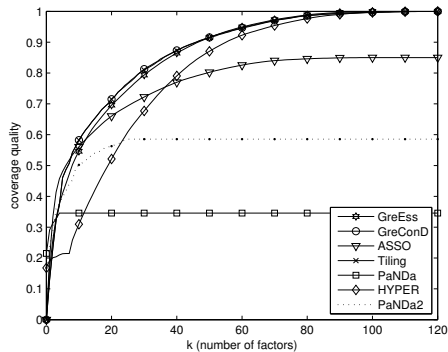
Figure: Coverage by the first k factors (synthetic data)

Real Datasets

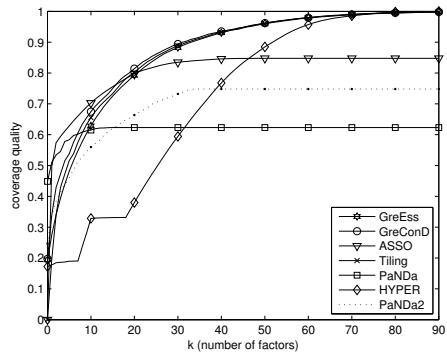
dataset	size	$\ I\ $	$\ \mathcal{E}(I)\ $	$\ \mathcal{E}(I)\ /\ I\ $
Mushroom	8124×119	186852	82965	0.444
DBLP	19×6980	40637	1601	0.039
Paleo	501×139	3537	1906	0.539
Chess	3196×76	118252	71296	0.603
DNA	4590×392	26527	1685	0.064

Table: Essential part of I (real data)

Results for Real Datasets I.



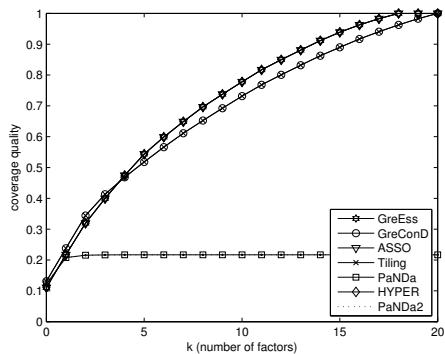
(a) Mushroom



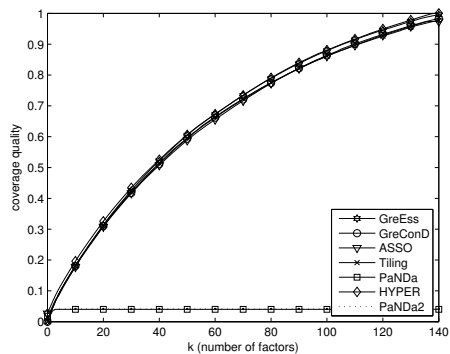
(b) Chess

Figure: Coverage by the first k factors (real data)

Results for Real Datasets II.



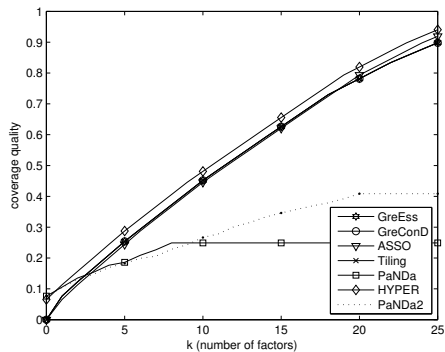
(a) DBLP



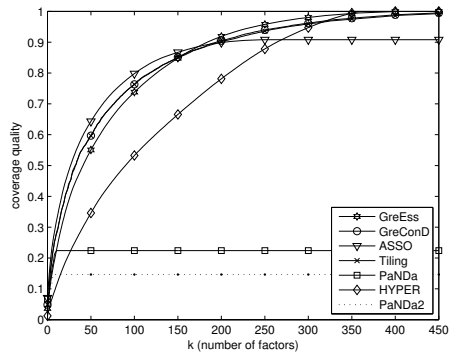
(b) Paleo

Figure: Coverage by the first k factors (real data)

Results for Real Datasets III.



(a) Tic-tac-toe



(b) DNA

Figure: Coverage by the first k factors (real data)

dataset	coverage (100c%)	Tiling	number of factors required for the prescribed coverage					
			ASSO	GreConD	PANDA	PANDA2	HYPER	GreEss
Mushroom	25%	3	2	3	1	2	8	2
	50%	7	6	7	NA	10	19	8
	75%	24	36	24	NA	NA	37	26
	100%	119	NA	120	NA	NA	122	105
DBLP	25%	2	2	2	NA	NA	2	2
	50%	5	5	5	NA	NA	5	5
	75%	10	10	11	NA	NA	10	10
	100%	21	19	20	NA	NA	19	19
Paleo	25%	16	16	16	NA	NA	14	15
	50%	39	40	39	NA	NA	38	38
	75%	75	76	76	NA	NA	73	73
	100%	151	NA	152	NA	NA	139	145
Chess	25%	2	1	1	1	1	9	1
	50%	5	2	4	NA	7	26	6
	75%	16	15	15	NA	NA	39	17
	100%	124	NA	124	NA	NA	90	113
DNA	25%	8	6	8	NA	NA	24	13
	50%	32	27	33	NA	NA	67	41
	75%	94	80	96	NA	NA	155	105
	100%	489	NA	496	NA	NA	392	408
Tic-tac-toe	25%	5	6	5	NA	10	5	5
	50%	12	12	12	NA	NA	11	12
	75%	19	19	19	NA	NA	18	19
	100%	31	29	32	NA	NA	29	32

Conclusion

- GreEss algorithm outperform existing algorithms in both number of factors and coverage quality.
- Empirical time complexity: GreEss is 2x slower than GreConD algorithm.
- Extracted factors seem to be more reasonable than factors from different algorithms (more detailed study is needed for support this claim).