Mining the potential relationship between cancer cases and industrial pollution based on high influence ordered pair patterns

**Abstract** This supplementary document presents the proofs and the detail of the function "searchRI".

1. The influence index does not meet the downward closure property.

For example, in the mining results of Table 3, we can see that

$$pc_1 = < \{waste \ management \ plant\}, \{prostate \ cancer\} >$$
  
 $pc_2 = < \{ , a \}, \{ e \ cancer\} > ,$ 

the influence index of the pattern  $pc_1$  is 0.611, and the influence index of the pattern  $pc_2$  is 0.668;  $pc_1 \subseteq pc_2$ , but  $PII(pc_1) < PII(pc_2)$ . So the influence index does not meet the downward closure property.

**Lemma 1 (Conditional Monotonicity)** If influence ordered pair patterns have the same influence features, the influence index is anti-monotone as the size of patterns increase.

**Proof.** Given two influence ordered pair patterns  $pc = \langle \mathit{IFS}_{pc}, \mathit{RFS}_{pc} \rangle$ ,  $pc' = \langle \mathit{IFS}_{pc}, \mathit{RFS}_{pc'} \rangle$  where  $\mathit{RFS}_{pc'} \subseteq \mathit{RFS}_{pc}$ .

For a reference feature  $c_j \in (RFS_{pc} \cap RFS_{pc})$ , any instance of  $c_j$  participating in a row instance of the pattern pc also certainly participates in a row instance of the pattern pc', so  $FIR(c_j,pc) \leq FIR(c_j,pc')$ , that is, the influence ratio of the feature is antimonotone. The influence index of the pattern is also antimonotonic

because:

$$PII(pc) = min_{c_j \in RFS_{pc}}(FIR(c_j, pc)) \leq min_{c_j \in RFS_{pc}}(FIR(c_j, pc')) \leq min_{c_j \in RFS_{pc'}}(FIR(c_j, pc')) = PII(pc').$$

**Lemma 2** The limit influence index of a pattern is an upper bound of the influence index of the pattern.

**Proof.** The maximum of the superimposed influence of  $c_j t$  is **max superimposed** influence of  $c_j t$ , so  $SII(c_j t) \le MSII(c_j t)$ .

$$\begin{split} FIR(c_{j},pc) &= \sum_{c_{j},t \in \pi_{c_{j}}(TI(pc))} SII(c_{j},t) / FIS(c_{j}) \leq \sum_{c_{j},t \in \pi_{c_{j}}(TI(pc))} SII(c_{j},t) / FIS(c_{j}) = LIR(c_{j},pc) \\ PII(pc) &= \min_{c_{i} \in R} (FIR(c_{j},pc)) \leq \min_{c_{i} \in R} (LIR(c_{j},pc)) = LII(pc) \,. \end{split}$$

**Lemma 3** The limit influence index is anti-monotone as the size of patterns increase.

**Proof.** Given two influence ordered pair patterns  $C = \langle I, R \rangle$ ,  $C' = \langle I', R' \rangle$  and a feature  $f_k$  where  $C' \subseteq C$ ,  $I' \cup R' \cup \{f_k\} = I \cup R$ .

(1) For an influence feature  $c_i \in (I \cap I')$ , any instance of  $c_i$  that participates in a row

instance of the pattern C also certainly participates in a row instance of the pattern C', so  $LIR(c_i, C) \le LIR(c_i, C')$ , that is, the limit influence ratio is antimonotone.

(2) 1) if  $f_k$  is a reference feature,  $\langle I', R' \cup \{f_k\} \rangle = \langle I, R \rangle = C$ 

From lemma 1, it can be known that  $LIR(f_i,\langle I',R' \cup \{f_k\}\rangle) \leq LIR(f_i,\langle I',R'\rangle)$ 

$$\begin{split} &LII(C) = LII(\langle I', R' \cup \{f_k\} \rangle) = \min_{f_i \in R' \cup \{f_k\}} (LIR(f_i, \langle I', R' \cup \{f_k\} \rangle)) \\ &= \min(LIR(f_i, \langle I', R' \cup \{f_k\} \rangle), LIR(f_k, \langle I', R' \cup \{f_k\} \rangle)) \\ &\leq \min_{f_i \in R'} (LIR(f_i, \langle I', R' \cup \{f_k\} \rangle)) \\ &\leq \min_{f_i \in R'} (LIR(f_i, \langle I', R' \cup \{f_k\} \rangle)) \end{split}$$

$$\leq \min_{f_i \in R'} (LIR(f_i, \langle I', R' \rangle)) = LII(C')$$

2) if  $f_k$  is an influence feature,  $\langle I' \cup \{f_k\}, R' \rangle = \langle I, R \rangle = C$  and it can be seen from 1 that  $LIR(c_j, C) \leq LIR(c_j, C')$ .

$$LII(C) = \min_{f_i \in R} (LIR(f_i, C)) = \min_{f_i \in R'} (LIR(f_i, C)) \leq \min_{f_i \in R'} (LIR(f_i, C')) = LII(C')$$

so limit influence index is antimonotone.

**Lemma 4** The participating instances of  $f_i$  in an influence ordered pair pattern pc must be included in  $CPIS(f_i, pc)$ , i.e.,  $PIS(f_i, pc) \subseteq CPIS(f_i, pc)$ .

**Proof.**  $\forall f_i.j \in PIS(f_i,pc)$ , there must be a row instance containing  $f_i.j$ . According to the join method, if  $f_i.j$  participate in the row instance of pc, then  $f_i.j$  must participate in row instance of  $pc_1$  and row instance of  $pc_2$  at the same time, i.e.,

$$f_i.j \in \{PIS(f_i, pc_1) \cap PIS(f_i, pc_2)\}$$
, so  $PIS(f_i, pc) \subseteq CPIS(f_i, pc)$ .

**Lemma 5** The influence ratio of  $f_i$  in pc based on candidate participating instance set is an upper bound of the true influence ratio of  $f_i$  in pc.

**Proof.** The influence ratio of the feature  $f_i$  in pc based on candidate participating instance set is denoted as  $CFIR(f_i, pc)$ .

$$\therefore PIS(f_i, pc) \subseteq CPIS(f_i, pc)$$

$$\therefore CFIR(f_i, pc) = \sum_{f_i.j \in CFIR(f_i, pc)} SII(f_i.j) / FIS(f_i) \geq \sum_{f_i.j \in PIS(f_i, pc)} SII(f_i.j) / FIS(f_i) = FIR(f_i, pc)$$

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Algorithm 3: RI = \text{searchRI}(f_i, j, pc)
1) k = pc. length()
2) RI .resize(k) //The capacity of RI is set to k
3) OssArr = \emptyset
4) for f_p \in \{pc - \{f_i\}\} do:
      get Oss(f_i \cdot j, f_p, pc) and OssArr[f_p] = Oss(f_i \cdot j, f_p, pc)
6) end for
7) featurePos = 0
8) f_p = pc[featurePos]
7) for instancePos = 0; instancePos < Oss. size(); instancePos + +
         RI[0] = particiInstanceArr[instancePos] //Take this element
9)
         gen_RI_recursion(RI, OssArr, k, featurePos + 1, 0, k - 1)
         if(verifyRowInstance((RI)) return RI
10)
11)
          gen_RI_recursion( RI , OssArr , k , featurePos , instancePos + 1, k ) //Do not take this
element, and take the next element of the feature
12)
         if(verifyRowInstance((RI)) return RI
```

## Algorithm 4: gen\_Rl\_recursion(RI, OssArr, k, featurePos, instancePos, remainder)

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1) if remainder == 0: return;
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- 2) **if** featurePos + remainder > k: **return**;
- 3)  $f_p = pc[featurePos]$
- 4)  $Oss = OssArr[f_n]$

13) return  $\varnothing$ 

- 5) **if** *instancePos* +1 > *Oss* .size() **return**;
- 6) RI[featurePos] = Oss[instancePos]
- 7) gen\_RI\_recursion( RI, OssArr, k, featurePos+1, 0, remainder-1);
- 8) gen\_Rl\_recursion(RI, OssArr, k, featurePos, instancePos+1, remainder);