

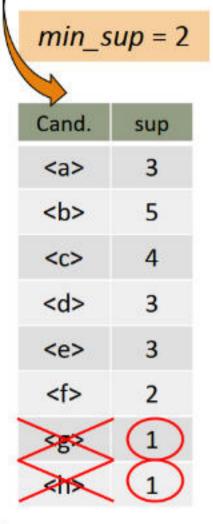
## **GSP: Apriori-Based Sequential Pattern Mining**

Initial candidates: All singleton sequences

<a>, <b>, <c>, <d>, <e>, <f>, <g>, <h>

Scan DB once, count support for each candidate

Generate length-2 candidate sequences



|         | <a></a>   | <b></b>   | <c></c>   | <d></d>   | <e></e>   | <f></f>   |
|---------|-----------|-----------|-----------|-----------|-----------|-----------|
| <a></a> | <aa></aa> | <ab></ab> | <ac></ac> | <ad></ad> | <ae></ae> | <af></af> |
| <b></b> | <ba></ba> | <bb></bb> | <bc></bc> | <bd></bd> | <be></be> | <bf></bf> |
| <c></c> | <ca></ca> | <cb></cb> | <cc></cc> | <cd></cd> | <ce></ce> | <cf></cf> |
| <d></d> | <da></da> | <db></db> | <dc></dc> | <dd></dd> | <de></de> | <df></df> |
| <e></e> | <ea></ea> | <eb></eb> | <ec></ec> | <ed></ed> | <ee></ee> | <ef></ef> |
| <f></f> | <fa></fa> | <fb></fb> | <fc></fc> | <fd></fd> | <fe></fe> | <ff></ff> |

|         | <a></a> | <b></b> | <c></c> | <d></d> | <e></e> | <f></f> |
|---------|---------|---------|---------|---------|---------|---------|
| <a></a> |         | <(ab)>  | <(ac)>  | <(ad)>  | <(ae)>  | <(af)>  |
| <b></b> |         |         | <(bc)>  | <(bd)>  | <(be)>  | <(bf)>  |
| <c></c> |         |         |         | <(cd)>  | <(ce)>  | <(cf)>  |
| <d></d> |         |         |         |         | <(de)>  | <(df)>  |
| <e></e> |         |         |         |         |         | <(ef)>  |
| <f></f> |         | 8       |         | 20      | 0       |         |

| SID | Sequence                        |
|-----|---------------------------------|
| 10  | <(bd)cb(ac)>                    |
| 20  | <(bf)(ce)b(fg)>                 |
| 30  | <(ah)(bf)abf>                   |
| 40  | <(be)(ce)d>                     |
| 50  | <a(bd)bcb(ade)></a(bd)bcb(ade)> |

Length-2 candidates:

36 + 15= 51

■ Without Apriori pruning: 8\*8+8\*7/2=92 candidates

GSP (Generalized Sequential Patterns): Srikant & Agrawal @ EDBT'96)

## **GSP Mining and Pruning**

Sth scan: 1 cand. 1 length-5 seq. pat.

(bd)cba>

Candidates cannot pass min\_sup threshold

4th scan: 8 cand. 7 length-4 seq. pat.

3rd scan: 46 cand. 20 length-3 seq. pat. 20 cand. not in DB at all

2nd scan: 51 cand. 19 length-2 seq. pat.

10 cand. not in DB at all

1st scan: 8 cand. 6 length-1 seq. pat.

(bd)bc> ...

Candidates cannot pass min\_sup threshold

candidates not in DB

candidates cannot pass min\_sup

candidates cannot pass min\_sup

- Repeat (for each level (i.e., length-k))
  - Scan DB to find length-k frequent sequences
  - Generate length-(k+1) candidate sequences from length-k
     frequent sequences using Apriori
  - set k = k+1
- Until no frequent sequence or no candidate can be found

| mm_sup = 2                      |
|---------------------------------|
| Sequence                        |
| <(bd)cb(ac)>                    |
| <(bf)(ce)b(fg)>                 |
| <(ah)(bf)abf>                   |
| <(be)(ce)d>                     |
| <a(bd)bcb(ade)></a(bd)bcb(ade)> |
|                                 |