

## 1. Exercise 15.2.4 (c) and (d)

- (a)
- Read  $R$  into some in-memory data structure that supports efficient *delete*, e.g. a hash table or a balanced binary search tree.
  - Read  $S$  into memory one page at a time.
  - For each tuple  $s$  in  $S$ , remove from  $R$  all tuples that agree with  $s$  in the common attributes.
  - Output the remaining tuples of  $R$ .
- (b)
- Read  $S$  into some in-memory data structure that supports efficient *search*, e.g. a hash table or a balanced binary search tree.
  - Read  $R$  into memory one page at a time.
  - For each tuple  $r$  in  $R$ , output  $r$  if  $r$  does *not* agree with any tuple in  $S$  in the common attributes.

## 2. Exercise 15.4.10

Consider TPMMS. Let  $R$  be the relation to be sorted,  $B$  be the number of disk pages for  $R$ , and  $M$  be the number of memory pages. TPMMS consists of the following steps:

- (a) Divide  $R$  into  $k$  sublists, where

$$k = \lceil \frac{B}{M} \rceil \leq M - 1$$

or in other words, each sublist contains the max number of tuples that can fit into memory, and the total number of sublists is less than  $M$  because we need one memory page as the output buffer.

- (b) Read in each sublist and perform in-memory sort, then write out the sorted sublist.  
 (c) Read in one page from each sublist, merge the sublists, and output the results.

Note that the I/O complexity of TPMMS is  $3B$  since all data pages have to be read in, write out, then read in again.

Now suppose the size of the last sublist is  $X$ . If we keep the last sublist in memory, we save  $2X$  I/O, so the problem becomes how we can maximize  $X$ . Note that the best we can do is:

$$X + k - 1 + 1 = M$$

or in other words,  $X$  can be as large as  $M - k$ , because we need  $k - 1$  pages to read in one page from each of the other sublists, and one page for output buffer. Also note that we have

$$k = \lceil \frac{B}{X} \rceil$$

Combine the two equations, we have a quadratic equation:

$$X^2 - MX + B = 0$$

From the quadratic formula,

$$X = \frac{M \pm \sqrt{M^2 - 4B}}{2}$$

So the I/O saving is  $2X = M + \sqrt{M^2 - 4B}$ .

### 3. Exercise 16.2.8

Intuitively, we cannot swap  $MIN$  and  $SUM$  because  $MIN$  has the effect of eliminating duplicates, and duplicates do contribute to  $SUM$ . On the other hand, swapping  $MIN$  and  $MAX$  seems to be OK. However, when you try to prove equation (b), you will notice that it is actually false, too.

(a) Let  $R(a,b) = \{ (1,1), (1,1), (2,2) \}$ .

$$\begin{aligned} LHS &= \gamma_{MIN(a) \rightarrow y, x}(\gamma_{a, SUM(b) \rightarrow x}(R)) \\ &= \gamma_{MIN(a) \rightarrow y, x}(\{(1,2), (2,2)\}) \\ &= \{(1,2)\} \\ RHS &= \gamma_{y, SUM(b) \rightarrow x}(\gamma_{MIN(a) \rightarrow y, b}(R)) \\ &= \gamma_{y, SUM(b) \rightarrow x}(\{(1,1), (2,2)\}) \\ &= \{(1,1), (2,2)\} \end{aligned}$$

Since  $LHS \neq RHS$ , equation (a) is false.

(b) Let  $R(a,b) = \{ (1,4), (1,3), (2,3) \}$ .

$$\begin{aligned} LHS &= \gamma_{MIN(a) \rightarrow y, x}(\gamma_{a, MAX(b) \rightarrow x}(R)) \\ &= \gamma_{MIN(a) \rightarrow y, x}(\{(1,4), (2,3)\}) \\ &= \{(1,4), (2,3)\} \\ RHS &= \gamma_{y, MAX(b) \rightarrow x}(\gamma_{MIN(a) \rightarrow y, b}(R)) \\ &= \gamma_{y, MAX(b) \rightarrow x}(\{(1,4), (1,3)\}) \\ &= \{(1,4)\} \end{aligned}$$

Since  $LHS \neq RHS$ , equation (b) is false.

4. Exercise 16.5.1

Simple estimation gives us

$$Est_s = \frac{T(R)T(S)}{MAX(V(R, Y), V(S, Y))} = \frac{52 \times 78}{20} = 203$$

With the histograms, based on the estimation method in Example 16.27, we have

$$Est_h = 5 \times 10 + 6 \times 8 + 4 \times 5 + 5 \times 3 + 7 \times 2 + 15 \times 2 \times 3 = 237$$

The two estimates are actually quite close, but note that the confidence of  $Est_h$  is higher because we know for sure that the join size is at least 118.

5. Exercise 18.2.4 (e)

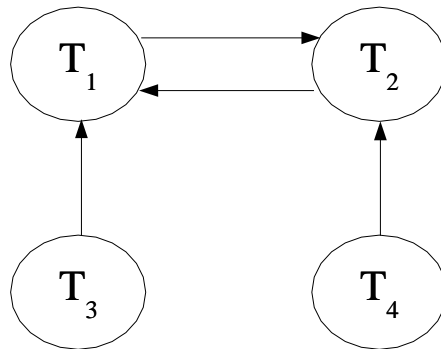


Figure 1: Precedence Graph

This schedule is not conflict-serializable, or serializable for that matter.

6. Exercise 18.4.2 (b)

Three. There are still two interleavings that are equivalent to  $(T_1, T_2)$  (see the online solution for the (a) part of the exercise), but there is only one interleaving that is equivalent to  $(T_2, T_1)$ .

7. Exercise 18.7.3

Based on the locking order, we have the following diagram:

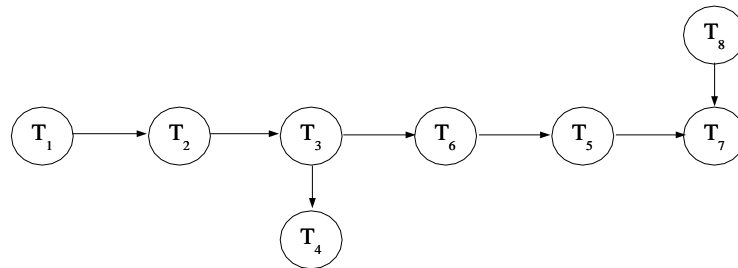


Figure 2: Locking Order

If we remove  $T_8$ , we have four serial orders:

- (a)  $T_1, T_2, T_3, T_4, T_6, T_5, T_7$
- (b)  $T_1, T_2, T_3, T_6, T_4, T_5, T_7$
- (c)  $T_1, T_2, T_3, T_6, T_5, T_4, T_7$
- (d)  $T_1, T_2, T_3, T_6, T_5, T_7, T_4$

Since  $T_8$  must come before  $T_7$ , we have  $7 + 7 + 7 + 6$  ways to add back  $T_8$ , therefore there are total of 27 serial orders.