Solutions for Homework Assignment #3

Answer to Question 1.

a. The scheduler S can be implemented by an augmented AVL tree D. Each node u of D contains information about a thread t of S. A node u contains the following fields:

- id(u): the integer id of the thread.
- status(u): the status of the thread, namely A, R or S.
- ready(u): a boolean variable that is True if and only if the subtree rooted at u (including u) has a node v such that status(v) = R. Note that the ready field of each node u of D satisfies the identity:

$$ready(u) = ready(lchild(u)) \lor (status(u) = R) \lor ready(rchild(u))$$
 (*)

where $ready(NIL) = FALSE.^{1}$

- lchild(u), rchild(u) and parent(u): pointers to the left child, right child, and the parent of u.
- BF(u): the balance factor of u.

The id field is the key of the node in the AVL tree D — thus, an in-order traversal of D visits all the threads in non-decreasing id order.

- **b.** The scheduler's operations are implemented as follows:
 - NewThread(t): Given a thread t = (i, status), where $status \in \{A, R, S\}$, inserts a node representing t into D; assumes that i is different from the id of every other thread in D.
 - (1) Prepare a new node u with id(u) = i and status(u) = A to represent thread t, and use the ordinary BST insertion algorithm to insert node u into D with i as the insertion key. After this insertion, the node u that represents t is a leaf of D.
 - (2) If the status of t is R then traverse the path from the new leaf u to the root of the tree D to update the ready fields: for each node v along this path set:

$$ready(v) := ready(lchild(v)) \lor (status(v) = R) \lor ready(rchild(v))$$

(3) Traverse the path from the new leaf u to the root of the tree D again, performing rotations and updating the balance factors as required by the ordinary AVL insertion algorithm. In addition, update the ready field of every node v involved in a rotation, according to the identity (*). Stop the rotations and updates to the balance factors as in the ordinary AVL insertion algorithm. Remark: The two traversals described above can be done in single pass where we first update the ready field of the node under consideration and then we do the balancing/rotations part (which may also require a few additional ready field updates).

Suppose that D contains n threads. Since D is an AVL tree, the height of any subtree of D is $O(\log n)$. The time required by NewThread(t) is that required by the ordinary AVL insertion algorithm, i.e., $O(\log n)$, plus the time required to update the ready fields of the new node's ancestors (if the new node's status is R), and the ready fields of the nodes involved in a rotation (a few nodes for each rotation). Since there are $O(\log n)$ such nodes and each ready update takes O(1) time, the additional time required to update all the ready fields is also $O(\log n)$. Thus, the worst-case time complexity of NewThread(t) is $O(\log n)$.

¹NIL denotes an empty pointer.

• FIND(i, x): Given a thread id i and a pointer to a node x of D, returns a pointer to the thread t = (i, -) in the subtree of D rooted at x; if no such thread exists, then it returns -1.

This is a straightforward BST search algorithm:

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\begin{aligned} &\operatorname{FIND}(i,x) \\ & \text{if } x = \operatorname{NIL} \text{ then} \\ & \operatorname{return} \ -1 \\ & \text{else if } id(x) < i \text{ then} \\ & \operatorname{return} \ \operatorname{FIND}(i,lchild(x)) \\ & \text{else if } id(x) = i \text{ then} \\ & \operatorname{return} \ x \\ & \text{else} \\ & \operatorname{return} \ \operatorname{FIND}(i,rchild(x)) \\ & \text{end if} \end{aligned}
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Note that FIND(i) is simply FIND(i,r) where r points to the root of the tree D.²

The worst-case time complexity of FIND(i, x) is $O(\log n)$. To see why, note that: (i) each call at a node u results in at most one recursive call, at the left or the right subtree of u; and (ii) each call involves a constant amount of work. So the time to execute FIND(i, x) is proportional to the height the subtree of D rooted at x, i.e., it is $O(\log n)$.

• Completed(i): If S has a thread t = (i, -) then remove t from S, else return -1.

First call Find(i) to determine whether S contains a thread t = (i, -). If it does not, i.e., Find(i) returns -1, then return -1. If it does, i.e., Find(i) returns a pointer x to the node of D that contains t = (i, -), proceed as follows:

- (1) Use the ordinary BST deletion algorithm to delete the node pointed to by x from D. Since D is an AVL tree, this deletion ultimately results in the removal of a leaf from the tree. Let z be the parent of that leaf. The rest of the algorithm is very similar to parts (2) and (3) of NewThread(t), except that the traversals described below start from node z rather than the newly added leaf in NewThread(t).
- (2) Traverse the path from z to the root of the tree D to update the ready fields: for each node v along this path, set $ready(v) := ready(lchild(v)) \lor (status(v) = R) \lor ready(rchild(v))$.
- (3) Traverse the path from z to the root of the tree D again, performing rotations and updating the balance factors as required by the ordinary AVL deletion algorithm. In addition, update the ready field of every node v involved in a rotation, according to the identity (*). Stop the rotations and updates to the balance factors as in the ordinary AVL deletion algorithm. Remark: The two traversals described above can be done in single pass where we first update the ready field of the node under consideration and then we do the balancing/rotations part (which may also require a few additional ready field updates).

The time required by COMPLETED(i) is that required by the ordinary AVL deletion algorithm, i.e., $O(\log n)$, plus the time required to update the ready fields of z's ancestors, and the ready fields of the nodes involved in a rotation (a few nodes for each rotation). Since there are $O(\log n)$ such nodes and each ready update takes O(1) time, the additional time required to update the ready fields is also $O(\log n)$. Thus, the worst-case time complexity of COMPLETED(i) is $O(\log n)$.

²More precisely, if the FIND(i) search is successful, it returns a *pointer* to a node u in D that represents some thread t = (i, status); given this pointer, it is trivial to return the pair t = (i, status) contained in u.

• ChangeStat(x, stat): Given a pointer x to a node containing thread t in D, sets the status of t to stat.

To do so, we first set status(x) to stat. If stat is not R, then we are done and just return. Otherwise, we update the ready field of every node in the path from x to the root of the tree D, according to the identity (*). Note that if the status of any node along this path does not change, this update process can stop (this obvious optimization is not shown in the basic pseudocode below).

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\begin{aligned} & \operatorname{CHANGESTAT}(x, stat) \\ & status(x) := stat \\ & \text{if } stat \neq R \text{ } \mathbf{return} \\ & y := x \\ & \mathbf{repeat} \\ & ready(y) = ready(lchild(y)) \lor (status(y) = R) \lor ready(rchild(y)) \\ & y := parent(y) \\ & \mathbf{until} \ y = \operatorname{NIL} \\ & \mathbf{return} \end{aligned}
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It is clear that the time to execute ChangeStat(x, stat) is is proportional to length of the path from x to the root of D. So the worst-case time complexity of ChangeStat(x, stat) is $O(\log n)$.

• CHANGESTATUS(i, stat): If S has a thread t = (i, -) then set the status of t to stat, else return -1.

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CHANGESTATUS(i, stat)

x := \text{FIND}(i)

if x = -1 then return -1

else return CHANGESTAT(x, stat)
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Since the worst-case time complexity of FIND(i) and CHANGESTAT(x, stat) is $O(\log n)$, the worst-case time complexity of CHANGESTATUS(i, stat) is also $O(\log n)$.

• SCHEDULENEXT(x): Given a pointer to a node x of D, returns a pointer to the thread t that has the smallest id among all the threads with status = R in the subtree of D rooted at x, and sets the status of t to A; if the subtree rooted at x has no thread with status = R, then the procedure returns -1.

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\begin{aligned} & \text{SCHEDULENEXT}(x) \\ & \text{if } x = \text{NIL} \lor ready(x) = \text{False then} \\ & \text{return } -1 \\ & \text{else if } ready(lchild(x)) = \text{True then} \\ & \text{return SCHEDULENEXT}(lchild(x)) \\ & \text{else if } status(x) = R \text{ then} \\ & \text{CHANGESTAT}(x, A) \\ & \text{return } x \\ & \text{else} \\ & \text{return SCHEDULENEXT}(rchild(x)) \\ & \text{end if} \end{aligned}
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Note that SCHEDULENEXT is simply SCHEDULENEXT(r) where r points to the root of the tree D.

The worst-case time complexity of SCHEDULENEXT(x) is $O(\log n)$. To see why, first note that SCHEDULENEXT(x) calls ChangeStat(x, A) at most once, and, as we have argued earlier, the time complexity of ChangeStat(x, A) is $O(\log n)$. Now consider the time required to execute SCHEDULENEXT(x) excluding the time to execute ChangeStat(x, A). Note that: (i) each call at a node x results in at most one recursive call, at the left or the right subtree of x; and (ii) each call involves a constant amount of work other than the recursive call that it makes and the call to ChangeStat(x, x). So the time to execute SCHEDULENEXT(x) excluding the time taken by ChangeStat(x, x) is proportional to the height the subtree of x rooted at x, i.e., it is x0 (log x1). Since the time complexity of ChangeStat(x1, x2) is x3 (log x3), the total time to execute SCHEDULENEXT(x3) is also x4 (log x5) in the worst-case.

Answer to Question 2.

a. If the number of empty slots in T is k before inserting x, what is the probability that x is inserted into an empty slot?

Let $E = \{i_1, i_2, \dots, i_k\}$ be the set of *empty* slots of T before inserting x.

The probability that h_1 hashes x into an *empty* slot of T is:

$$Prob[h_1(x) \in E] = Prob[h_1(x) = i_1 \lor h_1(x) = i_2 \lor \dots \lor h_1(x) = i_k]$$

= $\sum_{j=1}^{j=k} Prob[h_1(x) = i_j]$

Since T has m slots and we assume that the hash function h_1 satisfies the Simple Uniform Hashing Assumption (SUHA), we have $Prob[h_1(x) = i_j] = \frac{1}{m}$ for every $j, 1 \le j \le k$. Therefore:

$$Prob[h_1(x) \in E] = \sum_{j=1}^{j=k} \frac{1}{m} = \frac{k}{m}$$

So the probability that h_1 hashes x into a non-empty slot of T is:

$$Prob[h_1(x) \not\in E] = 1 - \frac{k}{m}$$

Since h_2 also satisfies SUHA, the probability that h_2 hashes x into a non-empty slot of T is also:

$$Prob[h_2(x) \not\in E] = 1 - \frac{k}{m}$$

Note that x is inserted in a non-empty slot of T if and only if both h_1 and h_2 hash into a non-empty slot of T. Therefore:

$$Prob[x \text{ is inserted in a } non-empty \text{ slot of } T] = Prob[h_1(x) \notin E \land h_2(x) \notin E]$$

Thus, since the hash functions h_1 and h_2 are independent:

$$Prob[x \text{ is inserted in a } non-empty \text{ slot of } T] = Prob[h_1(x) \notin E] \cdot Prob[h_2(x) \notin E]$$
$$= \left(1 - \frac{k}{m}\right)^2$$

Therefore:

$$Prob[x \text{ is inserted in an } empty \text{ slot of } T] = 1 - Prob[x \text{ is inserted in a } non-empty \text{ slot of } T]$$

$$= 1 - \left(1 - \frac{k}{m}\right)^2$$

$$= \frac{k(2m - k)}{m^2}$$

b. Suppose that m = 4, and T[0], contains 6 elements, T[1] contains 3 elements, and T[2], T[3] contain 9 elements each. What is the *expected* length of the chain x is inserted into, *not counting* x itself?

Since the table has 4 slots, when we insert x using the two independent hash functions h_1 and h_2 there are 16 possible hashing outcomes:

$$S = \{(i, j) \mid \text{ where } 0 \le i = h_1(x) \le 3 \text{ and } 0 \le j = h_2(x) \le 3\}$$
 (S is our sample space)

Note that:

$$Prob[(i,j)] = Prob[h_1(x) = i \land h_2(x) = j]$$
 (by definition)
 $= Prob[h_1(x) = i] \cdot Prob[h_2(x) = j]$ (because h_1 and h_2 are independent)
 $= \frac{1}{4} \cdot \frac{1}{4} = \frac{1}{16}$ (because h_1 and h_2 satisfy SUHA)

Let X be the random variable denoting the length of the chain where x is inserted. That is, for each possible outcome $o = (i, j) \in S$, X(o) is the length of the chain where x is inserted. We are seeking E[X]. Note that:

$$\begin{split} E[X] &= \sum_{o \in S} X(o) \cdot Prob[o] \\ &= \sum_{(i,j) \in S} X[(i,j)] \cdot Prob[(i,j)] \\ &= \frac{1}{16} \sum_{(i,j) \in S} X[(i,j)] \\ &= \frac{1}{16} \Big(X[(0,0)] + X[(0,1)] + X[(0,2)] + X[(0,3)] \\ &+ X[(1,0)] + X[(1,1)] + X[(1,2)] + X[(1,3)] \\ &+ X[(2,0)] + X[(2,1)] + X[(2,2)] + X[(2,3)] \\ &+ X[(3,0)] + X[(3,1)] + X[(3,2)] + X[(3,3)] \Big) \end{split}$$

Recall that T[0] contains 6 elements, T[1] contains 3 elements, and T[2], T[3] contain 9 elements each. From our hashing scheme, when the hashing outcome is (i,j), the length X[(i,j)] of the chain that x enters is as follows: if i=1 or j=1 then X[(i,j)]=3, else if i=0 or j=0 then X[(i,j)]=6 else X[(i,j)]=9. Therefore:

$$E[X] = \frac{1}{16} (6+3+6+6)$$

$$+ 3+3+3+3$$

$$+ 6+3+9+9$$

$$+ 6+3+9+9$$

$$= \frac{87}{16}$$

$$= 5.4375$$

Another way to compute E[X] is as follows. Note that the chain where x is inserted has length 3, 6 or 9. So the only possible value v of the random variable X is 3, 6 and 9. An alternative formula for E[X] is:

$$E[X] = \sum_{v \in \{3,6,9\}} v \cdot Prob[X=v]$$

It now suffices to compute Prob[X=3], Prob[X=6], and Prob[X=9]. Note that X=3 iff the outcome of the hashing is an (i,j) with i=1 or j=1. Therefore:

$$\begin{split} Prob[X=3] &= Prob[\{(0,1),(1,0),(1,1),(1,2),(1,3),(2,1),(3,1)\}] \\ &= Prob[(0,1)] + Prob[(1,0)] + Prob[(1,1)] + Prob[(1,2)] + Prob[(1,3)] + Prob[(2,1)] + Prob[(3,1)]\}] \\ &= 7 \cdot \frac{1}{16} = \frac{7}{16} \end{split}$$

Similarly:

$$\begin{split} Prob[X=6] &= Prob[\{(0,0),(0,2),(0,3),(2,0),(3,0)\}] \\ &= Prob[(0,0)] + Prob[(0,2)] + Prob[(0,3)] + Prob[(2,0)] + Prob[(3,0)]\}] \\ &= 5 \cdot \frac{1}{16} = \frac{5}{16} \\ Prob[X=9] &= Prob[\{(2,2),(2,3),(3,2),(3,3)\}] \\ &= Prob[(2,2)] + Prob[(2,3)] + Prob[(3,2)] + Prob[(3,3)]\}] \\ &= 4 \cdot \frac{1}{16} = \frac{4}{16} \end{split}$$

Therefore:

$$\begin{split} E[X] &= \sum_{v \in \{3,6,9\}} v \cdot Prob[X = v] \\ &= 3 \cdot \frac{7}{16} + 6 \cdot \frac{5}{16} + 9 \cdot \frac{4}{16} \\ &= \frac{87}{16} \end{split}$$

Answer to Question 3.

(1) We will use two data structures for our algorithm: a direct access table A with 26 indices (corresponding to each letter in the English alphabet) and a hash table T.

We assume that the size t of the hash table T is within a constant factor of l; more precisely, we assume that $t = \kappa \cdot l$, for some constant $\kappa > 0$. We will also assume our hashing function satisfies the Simple Uniform Hashing Assumption (SUHA).

(2) The idea for our algorithm is as follows. Initially, we set every value in our direct access table A to null; A will store the current most frequent word starting with each letter. When we receive an input word, we insert (word, 1) into our hash table T using word as our key if it does not already exist in H; otherwise (if it does), we increment its frequency:

$$(word, freq) \rightarrow (word, freq + 1)$$

Then, let the first letter of word be some letter corresponding to value x (e.g. "b" corresponds to 2); if word's frequency is greater than the frequency of the word at A[x] (or if A[x] == null), then set A[x] to word.

To perform a query operation, we simply read our direct access table from indices 1 to 26.

(3) CHAINED-HASH-SEARCH(T, k):

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return pointer to node with key k in list T[h(k)], null if doesn't exist
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CHAINED-HASH-INSERT(T, x):
   insert x at the head of list T[h(x.key)]

INSERT(T, A, word):
   // update hash table storing word frequencies
   u := CHAINED-HASH-SEARCH(T, word)
   if u = null:
      create a new node u s.t. u.key = word and u.value = 1
      CHAINED-HASH-INSERT(T, u)
   else:
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increment u.value by 1
// check if there is a new most frequent word
l = word[0] // first letter of word
x = 1 - 'a' // convert to number
current_max = CHAINED-HASH-SEARCH(T, A[x]).value
if u.value > current_max or (u.value = current_max and word < A[x]):
    A[x] := word

QUERY(A):
for i in 1 ... 26:
    print T[i]</pre>
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Note: it is also possible to store (word, freq) in the direct access table so that you don't have to look up the frequency of the current max. This slightly increases efficiency in exchange for a slight increase in storage.

(4) Under the simple uniform hashing assumption (SUHA), i.e., that each distinct value is equally likely to be hashed into any one of T's slots, the expected length of a linked list in T is α , where $\alpha = l/t$. Since t is $t = \kappa \cdot l$, α is $\Theta(1)$, and so the expected length of a linked list in T is $\Theta(1)$. Therefore the expected time for an INSERT operation is $\Theta(1)$ — a linked list in T is accessed twice; all other operations are clearly constant (in terms of the current length of the sequence). The QUERY operation runs in constant time since accessing every element in A is a constant time operation and |A| = 26; that is, |A| is not dependant on the length of the sequence.