Good Morning.

Randomized Algorithms, Lecture 6

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May 13th 2019

Today's Lecture

Data structures

The data-structuring problem

Random Treaps

Hashing fundamentals

Hashing with chaining

Two-level hashing

Maintain disjoint sets S_1, S_2, \ldots of *items* with keys from totally ordered universe \mathcal{U} under

```
MAKESET(S) Create a new empty set S.
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INSERT(i, S) Insert item i with key k(i) into S
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DELETE(
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) Delete item with key k from S .

$$FIND(k, S)$$
 Get item with key k in S

JOIN
$$(S_1, i, S_2)$$
 Given $k(i_1) < k(i) < k(i_2)$ for $i_1 \in S_1, i_2 \in S_2$.
Replace S_1 and S_2 with $S = S_1 \cup \{i\} \cup S_2$.

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 Given $k(i) < k(j)$ for $i \in S_1, j \in S_2$.
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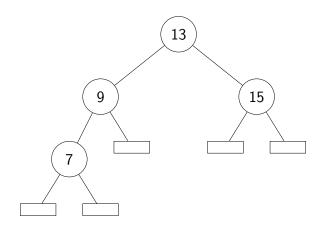
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Binary Search Tree

Suppose we assign a unique $key \ k(i)$ from some totally ordered set to each item of a set S.

A binary tree whose inner nodes store the items of S is in *symmetric order* and is called a *binary search tree* if, for each node with key k, the left subtree contains $\{j \in S \mid k(j) < k\}$ and the right subtree contains $\{j \in S \mid k(j) > k\}$.

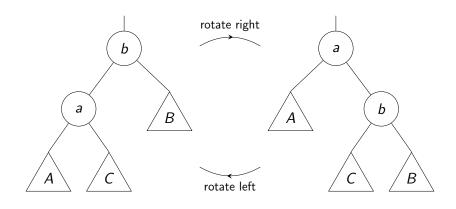
Binary Search Tree, Example



We assume that we only store items in internal nodes.

Leaves are marked on this slide using rectangles, but are left out of the drawing in the following.

Binary Search Tree, Rotations



```
function MAKESET(r)
   Initialize r as a leaf.
function FIND(k, r)
   if k(r.item) < k then
       return FIND(k, r.left)
   else if k(r.item) > k then
                      \triangleright If k \notin S_r this is a leaf and has no item.
function INSERT(i, r)
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function INSERT(i, r)
    v \leftarrow \text{FIND}(k(i), r)
    Replace leaf v with internal node i having 2 leaf children.
```

```
function JOIN(r_1, i, r_2)
   r \leftarrow new internal node having children r_1 and r_2.
    r.item \leftarrow i
    return r
function DELETE(k, r)
   if v.left is a leaf then
   else if v.right is a leaf then
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function JOIN(r_1, i, r_2)
    r \leftarrow new internal node having children r_1 and r_2.
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function DELETE(k, r)
    v \leftarrow \text{FIND}(k, r)
    if v.left is a leaf then
         Let v.right replace v as child of v.parent
    else if v.right is a leaf then
         Let v.left replace v as child of v.parent
    else
         v' \leftarrow \text{FIND}(\infty, v.\text{left}).\text{parent}
         i \leftarrow v'.item
         DELETE(k(i), v.left) \triangleright v' has a leaf as right child
         v.item \leftarrow i
```

```
function CONCAT(r_1, r_2)

i \leftarrow \text{FIND}(\infty, r_1).parent.item

DELETE(k(i), r_1)

return JOIN(r_1, i, r_2)

function SPLIT(k, r)

Use rotations to make k the root key.

return (r.\text{left}, r.\text{right}).
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A number of strategies exist that use rotations to keep the BST *balanced*.

- ► AVL-trees (1962)
- ▶ Red/Black trees (1978)
- ► Splay trees (1985)

AVL-trees explicitly maintain the height of each subtree and ensures the height difference between siblings is at most one. Simple to implement, but uses $\mathcal{O}(\log n)$ rotations per insert/delete in the worst case.

Red/black trees store a single bit of balancing info per node, and use at most 2 (3) rotations per insertion (deletion). They are complicated to implement but good in practice. Time is still worst case $\mathcal{O}(\log n)$ per operation.

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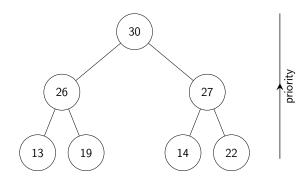
Max-Heap

Suppose we assign a unique *priority* p(i) from some totally ordered set to each item of a set S.

A tree whose inner nodes store the items of S is in max-heap order if the priority of each node is less than the priority of its parent.

Max-Heap, Example

 $\{13, 26, 19, 30, 14, 27, 22\}$



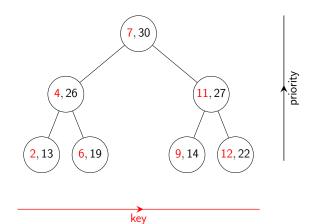
Treap

Suppose we assign both a unique key k(i) and a unique priority p(i) to each item $i \in S$.

A treap for S is a binary tree where each node stores an item from S, such that the nodes are in BST order wrt $k(\cdot)$ and in max-heap order wrt $p(\cdot)$.

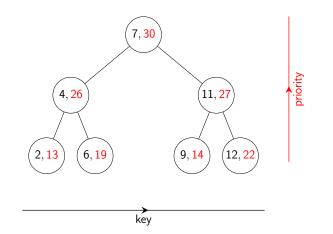
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 $\{(2,13),(4,26),(6,19),(7,30),(9,14),(11,27),(12,22)\}$



Treap, Example

 $\{(2, 13), (4, 26), (6, 19), (7, 30), (9, 14), (11, 27), (12, 22)\}$



Treap, Uniqueness

Theorem

Let $S = \{(k_1, p_1), \dots, (k_n, p_n)\}$ be any set of key-priority pairs such that the keys are distinct and the priorities are distinct. Then there is a unique treap T(S) for it.

Treap, Uniqueness Proof

By induction on n. Trivial for n < 1, so let n > 1 and suppose it holds for all S' with |S'| < n. Let p_i be (unique) highest priority in S. If a treap exists for S, then (k_i, p_i) is the root, $S_1 = \{(k_i, p_i) \mid k_i < k_i\}$ is in the left subtree, and $S_2 = \{(k_i, p_i) \mid k_i > k_i\}$ is in the right subtree. By induction, $T(S_1)$ and $T(S_2)$ exist and are unique, so we can recursively construct the unique T(S) from

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Random Treap

Idea: We can assign distinct uniformly random priorities on insert and use rotations to maintain the treap invariant.

Theorem

Let $x_{(k)}$ denote the node containing the item with the k'th smallest key. The expected depth of $x_{(k)}$ is $H_k + H_{n-k+1} - 1 \in \mathcal{O}(\log n)$.

We will assume in our analysis that the random priorities are distinct. This can be justified by e.g. using the keys (which are required to be distinct) to disambiguate.

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This means that the time for a successful find is $\mathcal{O}(\log n)$.

Finding the time for an *unsuccessful* find is part of the next assignment.

Let $p_{(i)}$ denote the priority of $x_{(i)}$, and let X_{ik} indicate that $x_{(i)}$ is ancestor to $x_{(k)}$.

Then for i < k

$$X_{ik} = 1 \iff p_{(i)} = \max\{p_{(i)}, \dots, p_{(k)}\}$$

$$\mathbb{E}[X_{ik}] = Pr[X_{ik} = 1] = \frac{1}{k+1-i}$$

$$\mathbb{E}\left[\sum_{i \neq k} X_{ik}\right] = \sum_{k=0}^{k} \frac{1}{d} = H_k - 1$$

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Let $x_{(j)}$ be the unique deepest node such that the subtree rooted at $x_{(j)}$ contains both $x_{(i)}$ and $x_{(k)}$. This is also known as the *nearest common ancestor* $x_{(j)} = \operatorname{nca}(x_{(i)}, x_{(k)})$.

By definition, either j = i, or j = k, or $x_{(i)}$ is in the left subtree and $x_{(k)}$ is in the right subtree.

Since $x_{(j)}$ is the root of this subtree, $p_{(j)} \ge \max\{p_{(i)}, \dots, p_{(k)}\}.$

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Expectation of indicator variable.

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Linearity of expectation and change of variable, setting d = k + 1 - i.

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Similarly for i > k

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$$\mathbb{E}[X_{ik}] = Pr[X_{ik} = 1] = \frac{1}{i+1-k}$$

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$$\mathbb{E}[\operatorname{depth}(x_{(k)})] = \mathbb{E}\left[\sum_{i=1}^{n} X_{ik}\right]$$

$$= \mathbb{E}\left[\sum_{i < k} X_{ik}\right] + \mathbb{E}\left[\sum_{i > k} X_{ik}\right] + \mathbb{E}[X_{kk}]$$

$$= (H_k - 1) + (H_{n+1-k} - 1) + 1$$

$$= H_k + H_{n+1-k} - 1$$

The depth of a node is defined as the number of ancestors, including the node itself.

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Linearity of expectation.

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$$egin{aligned} \mathbb{E}[\mathsf{depth}(x_{(k)})] &= \mathbb{E}\Big[\sum_{i=1}^n X_{ik}\Big] \\ &= \mathbb{E}\Big[\sum_{i < k} X_{ik}\Big] + \mathbb{E}\Big[\sum_{i > k} X_{ik}\Big] + \mathbb{E}[X_{kk}] \\ &= (H_k - 1) + (H_{n+1-k} - 1) + 1 \\ &= H_k + H_{n+1-k} - 1 \end{aligned}$$

Previous slides.

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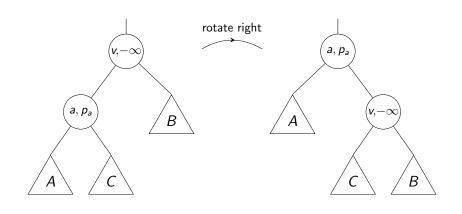
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Random Treap, Deletion

```
We can implement DELETE(k, r) on random treaps
as
 1: function RT-DELETE(k, r)
        v \leftarrow \text{FIND}(k, r)
        v.priority \leftarrow -\infty
 3:
        while v has a non-leaf child do
 4:
            if v.left.priority > v.right.priority then
 5:
                ROTATE-RIGHT(\nu)
 6:
            else
                ROTATE-LEFT(v)
 8:
        Set v.item ← nil and delete v.left and v.right
 9:
```

Random Treap, Deletion



Note that this works, because 1) Setting the priority to $-\infty$ only violates the max-heap invariant for the children of that node, and 2) each rotation moves the violation to a smaller subtree.

Also note that each rotation we do adds exactly one new ancestor to ν .

Random Treap, Deletion

Theorem

The expected number of rotations made by each call to RT-DELETE is less than 2.

Let $v = x_{(k)}$, and let Y_{ik} indicate that $x_{(i)}$ becomes a new ancestor of $x_{(k)}$.

Then for i < k

$$Y_{ik} = 1 \iff p_{(i)} = \max\{p_{(i)}, \dots, p_{(k-1)}, -\infty\}$$

$$\land \neg (p_{(i)} = \max\{p_{(i)}, \dots, p_{(k)}\})$$

$$\iff p_{(k)} > p_{(i)} > \max\{p_{(i+1)}, \dots, p_{(k-1)}\}$$

$$\mathbb{E}[Y_{ik}] = \Pr[Y_{ik} = 1] = \frac{1}{(k-i+1)(k-i)} = \frac{1}{k-i} - \frac{1}{k-i+1}$$

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Finally

$$\mathbb{E}[\#\text{rotations}] = \mathbb{E}[\#\text{new ancestors to } v]$$

$$= \mathbb{E}\left[\sum_{i=1}^{n} Y_{ik}\right]$$

$$= \mathbb{E}\left[\sum_{i=1}^{k-1} Y_{ik}\right] + \mathbb{E}\left[\sum_{i=k+1}^{n} Y_{ik}\right] + \mathbb{E}[Y_{kk}]$$

$$= (1 - \frac{1}{k}) + (1 - \frac{1}{n+1-k}) + 0$$

$$< 2$$

Hash function

Given a (large) universe U of keys, and a positive integer m.

Definition

A hash function $h: U \to [m]$ is a random variable, whose values are functions from $U \to [m]$.

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Definition

A hash function $h: U \to [m]$ is *c-universal* if, for all $x \neq y \in U$: $\Pr[h(x) = h(y)] \leq \frac{c}{m}$.

For many purposes c-universal hash functions for some small constant c are enough. We will see examples of such functions a little later today.

Let U = [u] and pick prime $p \ge u$. For any $a \in [p]_+ = \{1, \dots, p-1\}, b \in [p], \text{ and }$

$$a \in [p]_+ = \{1,\ldots,p-1\},\ b \in [p],\ ext{and} \ m < u,\ ext{let}\ h^m_{a,b}:U o [m] \ ext{be}$$

$$h_{a,b}^m(x) = ((ax+b) \bmod p) \bmod m$$

$$p_{a,b}^{m}(x) = ((ax + b) \mod p) \mod m$$

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Is this a hash function?

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 $h_{a,b}^m(x) = ((ax + b) \bmod p) \bmod m$

 $m_{a,b}(x) = ((ax + b) \dots a + b) \dots$

Is this a hash function? NO!

Let U = [u] and pick prime $p \ge u$. For any $a \in [p]_+ = \{1, \ldots, p-1\}$, $b \in [p]$, and m < u, let $h_{a,b}^m : U \to [m]$ be

$$h_{a,b}^m(x) = ((ax+b) \bmod p) \bmod m$$

Choose $a \in [p]_+$ and $b \in [p]$ uniformly at random, and let $h(x) := h_{a,b}^m(x)$. Then $h : U \to [m]$ is a universal hash function.

Multiply-shift

Let $U = [2^w]$ and $m = 2^\ell$. For any odd $a \in [2^w]$ define

$$a \in [2^w]$$
 define $h_a(x) := \left | rac{(ax) mod 2^w}{2^{w-\ell}}
ight |$

Choose odd $a \in [2^w]$ uniformly at random, and let $h(x) := h_a(x)$.

Then $h: U \rightarrow [m]$ is a 2-universal hash function.

Multiply-shift, C

```
For U=[2^{64}] the C code looks like this:

#include<stdint.h>

uint64_t hash(uint64_t x, uint64_t 1, uint64_t a) {

return (a*x) >> (64-1);

}
```

Goal is to maintain $S \subseteq U$, |S| = n and for $x \in U$ be able to tell if $x \in S$.

Idea: Let
$$m \ge n$$
 and pick universal $h: U \to [m]$. Store array L , where $L[i] = \text{linked list over } \{y \in S \mid h(y) = i\}.$

Then
$$x \in S \iff x \in L[h(x)]$$
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This can be checked in $\mathcal{O}(|L[h(x)]| + 1)$ time.

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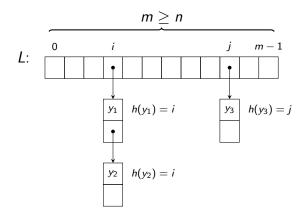
Idea: Let m > n and pick universal

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This can be checked in $\mathcal{O}(|L[h(x)]|+1)$ time.



Theorem

For $x \notin S$, $\mathbb{E}[|L[h(x)]|] \leq 1$

$$\mathbb{E}[|L[h(x)]|] = \mathbb{E}\Big[\big|\{y \in S \mid$$

$$\mathbb{E}[|\mathcal{L}[n(x)]|] - \mathbb{E}[|\{y \in \mathcal{S} \mid n(y) = n(x)\}|]$$

$$=\mathbb{E}\Big[\sum[h(y)=h(x)]\Big]$$

 $\leq |S|^{\frac{1}{m}} = \frac{n}{m} \leq 1$

$$= \sum_{y \in S} \mathbb{E} \Big[[h(y) = h(x)] \Big]$$

$$= n(x)$$

$$=\sum_{x\in X}\Pr[h(y)=h(x)]$$

Theorem

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Proof.

$$\mathbb{E}[|L[h(x)]|] = \mathbb{E}\Big[\big|\{y \in S \mid h(y) = h(x)\}\big|\Big]$$

$$= \mathbb{E}\left[\sum [h(y) = h(x)]\right]$$

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$$= \mathbb{E}\Big[\sum_{x\in C}[h(y) = h(x)]\Big]$$

$$=\sum_{y\in S}\mathbb{E}\Big[[h(y)=h(x)]\Big]$$

$$= \sum_{y \in S} \Pr[h(y) = h(x)]$$

By definition of $L[i] := \{ y \in S \mid h(y) = i \}.$

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$$\mathbb{E}[|L[h(x)]|] = \mathbb{E}\Big[\big|\{y \in S \mid h(y) = h(x)\}\big|\Big]$$
$$= \mathbb{E}\Big[\sum_{x} [h(y) = h(x)]\Big]$$

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 $= \sum \Pr[h(y) = h(x)]$

$$\leq |S|^{\frac{1}{m}} = \frac{n}{m} \leq 1$$

Here we use the *Iverson Bracket* notation

$$[P] := \begin{cases} 1 & \text{if } P \text{ is true} \\ 0 & \text{if } P \text{ is false} \end{cases}$$

This can often be used as a shorthand for an indicator variable.

In this case [h(y) = h(x)] becomes an indicator variable for the event h(y) = h(x).

Linearity of expectation.

Hashing with chaining

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$$\sum [h(y)]$$

$$p(y) =$$

$$= \mathbb{E}\Big[\sum_{x\in C}[h(y)=h(x)]\Big]$$

$$h(y) = h(y)$$

$$=\sum \mathbb{E}\Big[[h(y)=h(x)]\Big]$$

$$[n(y) - n(y)]$$

$$=\sum_{y\in\mathcal{S}}\Pr[h(y)=h(x)]$$

$$\leq |S| \frac{1}{m} = \frac{n}{m} \leq 1$$

Then by definition of a universal hash function
$$h: U \to [m]$$
, $\Pr[h(y) = h(x)] \le \frac{1}{m}$.

Since $x \notin S$ and $y \in S$, we have $x \neq y$.

Two-level hashing

When storing a *static* set, we can get *worst case* constant time lookup with linear space by using two levels of hashing, as follows:

- Step I Pick universal hash function $h: U \to [n]$. Let $S_i = \{x \in S \mid h(x) = i\}, \ s_i = |S_i|,$ and $B = \sum_{i=0}^{n-1} \binom{s_i}{2}$. If $B \ge n$, start over with a new h, else continue to Step II.
- Step II For each $i \in [n]$, pick a universal hash function $h_i: U \to [s_i(s_i-1)]$ until h_i has no collisions on S_i .

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$$h: U \to [n]$$
.
Let $S_i = \{x \in S \mid h(x) = i\}, \ s_i = |S_i|,$
and $B = \sum_{i=0}^{n-1} \binom{s_i}{2}$. If $B \ge n$, start over with a new h , else continue to Step II.

Step II For each
$$i \in [n]$$
, pick a universal hash function $h_i: U \to [s_i(s_i-1)]$ until h_i has no collisions on S_i .

Two-level hashing

When storing a *static* set, we can get *worst case* constant time lookup with linear space by using two levels of hashing, as follows:

Step I Pick universal hash function
$$h: U \to [n]$$
.
Let $S_i = \{x \in S \mid h(x) = i\}, \ s_i = |S_i|,$
and $B = \sum_{i=0}^{n-1} {s_i \choose 2}$. If $B \ge n$, start over with a new h , else continue to Step II.

Step II For each $i \in [n]$, pick a universal hash function $h_i: U \to [s_i(s_i-1)]$ until h_i has no collisions on S_i .

Two-Level hashing

The expected total time for finding h is $\mathcal{O}(n)$.

$$B = \sum_{\substack{x,y \in S \\ x \neq y}} [h(x) = h(y)]$$

$$\mathbb{E}[B] = \sum_{x \neq y} \Pr[h(x) = h(y)] \le \binom{n}{2} \cdot \frac{1}{n} = \frac{n}{2}$$

$$\Pr[B \ge 2\mathbb{E}[B]] \le \frac{1}{2}$$
 (By Markov) is geometrically distributed, so the expected wher of trials before $B < n$ is ≤ 2 , and each trial

B is geometrically distributed, so the expected number of trials before B < n is ≤ 2 , and each trial takes linear time.

Two-Level hashing

 $x,y \in S_i$

takes linear time.

The expected total time for finding h_i is $\mathcal{O}(s_i)$.

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 is $\mathcal{O}(s_i)$.
$$B_i = \sum [h(x) = h(y)]$$

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 $\mathbb{E}[B_i] = \sum_i \Pr[h(x) = h(y)] \le {s_i \choose 2} \cdot \frac{1}{s_i(s_i - 1)} = \frac{1}{2}$

$$\Pr[B \ge 2\mathbb{E}[B]] \le \frac{1}{2}$$
 (By Markov)

 B_i is geometrically distributed, so the expected number of trials before $B_i < 1$ is < 2 , and each trial

- ► First as a simple means of balancing a binary search tree by viewing it as a random treap
- Then we defined the concept of a hash function, and saw some actual practical hash functions.
- Finally we saw a simple application of universal hashing, when we looked at hashing with chaining and two-level hashing.
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