Searching with Dice A survey on randomized data structures

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Papers We Love
$$f(x) = x$$

- Introduction
- Skip lists
- Randomized binary search trees







R. Karp

N. C. Metropolis

M. O. Rabin

The usefulnees of randomization in the design of algorithms has been known for a long time:

- Metropolis' algorithms
- Rabin's primality test
- Rabin-Karp's string search



M.N. Wegman

- Hashing is another early success of randomization for the design of data structures.
- Selecting the hash function from a universal class (Carter and Wegman, 1977) guarantees expected performance

Randomization yields algorithms:

- Simple and elegant
- Practical
- With guaranteed expected performance
- Without assumptions on the probabilistic distribution of the input

- The usual worst-case analysis is not useful for randomized algorithms
- The probabilistic model to use in the analysis is under control; it is not a working hypothesis, but built-in

In this talk:

- Skip lists
- Randomized binary search trees

- Introduction
- Skip lists
- Randomized binary search trees



W. Pugh

- Skip lists were invented by William Pugh (C. ACM, 1990) as a simple alternative to balanced trees
- The algorithms to search, insert, delete, etc. are very simple to understand and to implement, and they have very good expected performance—independent of any assumption on the input

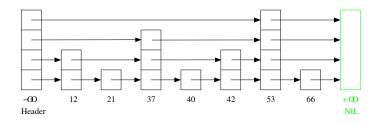


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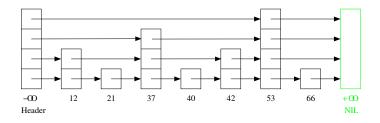
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A skip list S for a set X consists of:

- **1** A sorted linked list L_1 , called level **1**, contains all elements of X
- ② A collection of non-empty sorted lists L_2, L_3, \ldots , called level 2, level 3, . . . such that for all $i \geq 1$, if an element x belongs to L_i then x belongs to L_{i+1} with probability q, for some 0 < q < 1, p := 1 q



To implement this, we store the items of X in a collection of nodes each holding an item and a variable-size array of pointers to the item's successor at each level; an additional dummy node gives access to the first item of each level



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```
template <typename Key, typename Value>
class Dictionary {
public:
private:
  struct node_skip_list {
     Key k;
    Value v;
     vector < node_skip_list *> next;
     node_skip_list(const Key& the_key, const Value& the_value, int h) :
                    k(the_key), v(the_value), next(h, nullptr) {
 };
  node_skip_list* header;
 int height;
 double p; // e.g., p = 0.5
```

- The level or height of a node x, height(x), is the number of lists it belongs to.
- It is given by a geometric r.v. of parameter p:

$$\Pr\{\text{height}(x) = k\} = pq^{k-1}, \qquad q = 1 - p$$

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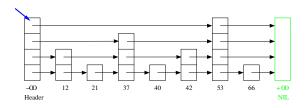
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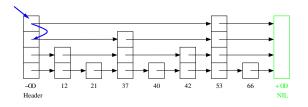
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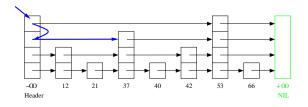
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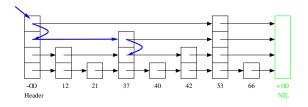
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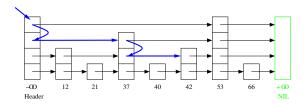
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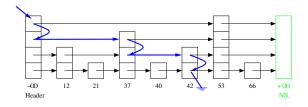


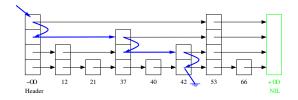


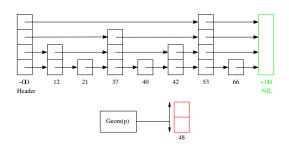


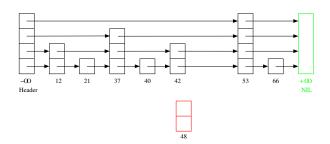


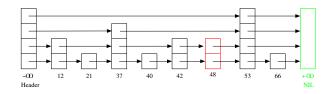












To insert a new item we go through four phases:

- Search the given key. The search loop is slightly different from before, since we need to keep track of the last node seen at each level before descending from that level to the one immediately below.
- 2) If the given key is already present we only update the associated value and finish.

```
void insert_skip_list(const Key& k, const Value& v) {
  // search for insertion point for the new item with key k
  // (or detect it is duplicate)
   node_skip_list* p = header;
   int 1 = height - 1;
   vector < node_skip_list *> pred(height, header);
   while (1 >= 0)
      if (p \rightarrow next[1] == nullptr or k <= p <math>\rightarrow next[1] \rightarrow k) {
         pred[1] = p; // <===== keep track of predecessor at level 1</pre>
         --1;
      } else {
         p = p -> next[1];
   if (p -> next[0] == nullptr or p -> next[0] -> k != k) {
      // k is not present, add new node here
   else // k is present, update associated value
      p -> next[0] -> v = v;
```

- 3) When k is not present, create a new node with k and v, and assign a random level r to the new node, using geometric distribution
- 4) Link the new node in the first r lists, adding empty lists if r is larger than the maximum level of the skip list

```
void insert_skip_list(...) {
  // adding new node
  // generate random height
  // each call to rng() produces a (pseudo)random
  // number uniformly distr. in (0,1)
   int h = 1; while (rng() > p) + h;
   // create new node
   node_skip_list* nn = new node_skip_list(k, v, h);
   if (h > height) {
     // add new levels to the header and to pred, if necessary
     // make pred[i] = _header for all i = _height .. h-1
      (header -> next).resize(h. nullptr):
      pred.resize(h. header);
   // link the new node to h linked lists
   for (int i = h - 1; i >= 0; --i) {
       nn -> next[i] = pred[i] -> next[i];
       pred[i] -> next[i] = nn:
```

- Deletions are also very easy to implement
- Ordered iterators are trivially implemented
- Skip list can also support many other operations, e.g., merging search and deletion by rank, finger search, . . .
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A preliminary rough analysis considers the search path backwards. Imagine we are at some node x and level i:

- ullet The height of x is >i and we come from level i+1 since the sought key k is smaller than the key of the successor of x at level i+1
- The height of x is i and we come from x's predecessor at level i since k is larger or equal to the key at x

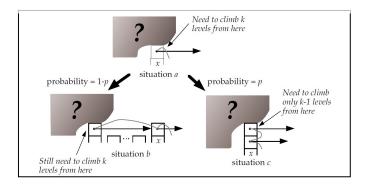


Figure from W. Pugh's *Skip Lists: A Probabilistic Alternative to Balanced Trees* (C. ACM, 1990)—the meaning of p is the opposite of what we have used!

The expected number ${\cal C}(k)$ of steps to "climb" k levels in an infinite list

$$C(k) = p(1 + C(k)) + (1 - p)(1 + C(k - 1))$$

$$= 1 + pC(k) + qC(k - 1) = \frac{1}{q}(1 + qC(k - 1))$$

$$= \frac{1}{q} + C(k - 1) = k/q$$

since C(0) = 0.

The analysis above is pessimistic since the list is not infinite and we might "bump" into the header. Then all remaining backward steps to climb up to a level k are vertical—no more horizontal steps. Thus the expected number of steps to climb up to level L_n is

$$\leq (L_n-1)/q$$

• L_n = the largest level L for which

$$\mathop{\mathbb{E}}[\text{\# of nodes with height} \geq L] \leq 1/q$$

• Probability that a node has height $\geq k$ is

$$\Pr\{\mathsf{height}(x) \ge k\} = \sum_{i \ge k} pq^{i-1} = pq^{k-1} \sum_{i \ge 0} q^i = q^{k-1}$$

• Number of nodes with height $\geq k$ is a binomial r.v. with parameters n and q^{k-1} , hence

$$\mathop{\mathbb{E}}[\#$$
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$$nq^{L_n-1} = 1/q \implies L_n = \log_q(1/n) = \log_{1/q} n$$

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Then the steps remaining to reach H_n (=the height of a random skip list of size n) can analyzed this way:

- we need not more horizontal steps than nodes with height $\geq L_n$, the expected number is $\leq 1/q$, by definition
- the probability that $H_n > k$ is

$$1 - \left(1 - q^k\right)^n \le nq^k$$

It follows that

$$\mathbb{E}[H_n] \le L_n + 1/p$$

and the expected additional vertical steps need to reach H_n from L_n is $\leq 1/p$

Summing up, the expected path length of a search is

$$\leq (L_n - 1)/q + 1/q + 1/p = \frac{1}{q} \log_{1/q} n + 1/p$$

On the other hand, the average number of pointers per node is 1/p so there is a trade-off between space and time:

- $ullet p o 0, q o 1 \implies$ very tall "nodes", short horizontal cost
- $ullet p o 1, q o 0 \implies {\sf flat\ skip\ lists}$
- Pugh suggests p=3/4, optimal choice minimizes factor $(q\ln(1/q))^{-1}$ is $q=e^{-1}=0.36\ldots, p=1-e^{-1}\approx 0.632\ldots$

Analysis of the height





W. Szpankowski

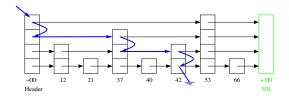
V. Rego

Theorem (Szpankowski and Rego,1990)

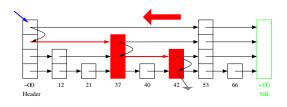
$$\mathbb{E}[H_n] = \log_Q n + \frac{\gamma}{L} - \frac{1}{2} + \chi(\log_Q n) + \mathcal{O}(1/n)$$

with Q:=1/q, $L:=\ln Q$, $\chi(t)$ a fluctuation of period 1, mean 0 and small amplitude.

The number of forward steps $F_{n,k}$ is the number of weak left-to-right maxima in $a_k, a_{k-1}, \ldots, a_1$, with $a_i = \mathsf{height}(x_i)$



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Total unsuccessful search cost

$$C_n = \sum_{0 \le k \le n} C_{n,k} = nH_n + F_n$$

Total forward cost

$$F_n = \sum_{0 \le k \le n} F_{n,k}$$

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P. Kirschenhofer

H. Prodinger

Theorem (Kirschehofer, Prodinger, 1994)

The expected forward cost in a random skip list of size n is

$$\mathbb{E}[F_n] = (Q-1)n\left(\log_Q n + \frac{\gamma - 1}{L} - \frac{1}{2} + \frac{1}{L}\chi(\log_Q n)\right) + \mathcal{O}(\log n),$$

with Q:=1/q, $L=\ln Q$ and χ a periodic fluctuation of period 1, mean 0 and small amplitude.

Skip Lists in Real Life

Usages [edit]

List of applications and frameworks that use skip lists:

- . MemSQL uses skip lists as its prime indexing structure for its database technology.
- Lucene uses skip lists to search delta-encoded posting lists in logarithmic time. [citation needed]
- QMap
 @ (up to Qt 4) template class of Qt that provides a dictionary.
- Redis, an ANSI-C open-source persistent key/value store for Posix systems, uses skip lists in its implementation of ordered sets.
- nessDB @, a very fast key-value embedded Database Storage Engine (Using log-structured-merge (LSM) trees), uses skip lists for its memtable.

- Speed Tables of are a fast key-value datastore for Tcl that use skiplists for indexes and lockless shared memory.
- Con Kolivas' MuQSS^[8] Scheduler for the Linux kernel uses skip lists

Skip lists are used for efficient statistical computations \mathcal{Q} of running medians (also known as moving medians). Skip lists are also used in distributed applications (where the nodes represent physical computers, and pointers represent network connections) and for implementing highly scalable concurrent priority queues with less lock contention, [19] or even without locking [19] as well as lookless concurrent dictionaries. [14] There are also several US patents for using skip lists to implement (lockless) briority queues and concurrent dictionaries. [15]

See also [edit]

• Bloom filter

Source: Wikipedia

To learn more

L. Devroye.

A limit theory for random skip lists.

The Annals of Applied Probability, 2(3):597–609, 1992.

- P. Kirschenhofer and H. Prodinger. The path length of random skip lists. Acta Informatica, 31(8):775–792, 1994.
- P. Kirschenhofer, C. Martnez and H. Prodinger. Analysis of an Optimized Search Algorithm for Skip Lists. Theoretical Computer Science, 144:199–220, 1995.

To learn more (2)

T. Papadakis, J. I. Munro, and P. V. Poblete. Average search and update costs in skip lists. *BIT*, 32:316–332, 1992.

H. Prodinger.

Combinatorics of geometrically distributed random variables: Left-to-right maxima.

Discrete Mathematics, 153:253-270, 1996.

W. Pugh.

Skip lists: a probabilistic alternative to balanced trees. *Comm. ACM*, 33(6):668–676, 1990.

🖥 W. Pugh.

A Skip List Cookbook.

Technical Report UMIACS-TR-89-72.1. U. Maryland, College Park, 1989.

- Introduction
- 2 Skip lists
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C. Aragon

R. Seidel

Two incarnations

- Randomized treaps (tree+heap) invented by Aragon and Seidel (FOCS 1989, Algorithmica 1996) use random priorities and bottom-up balancing
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- In a random binary search tree (built using random insertions) any of its n elements is the root with probability 1/n
- Idea: to insert a new item, insert it at the root with probability 1/(n+1), otherwise proceed recursively
- The random priorities of treaps "simulate" random timestamps (cif. Vuillemin's Cartesian trees 1980); rotations are used to maintain the BST invariant w.r.t. keys and the heap invariant w.r.t. priorities

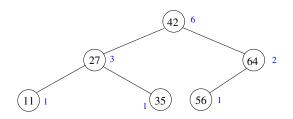
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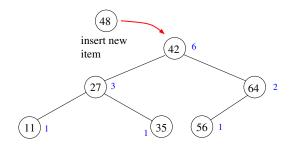
J. Vuillemin

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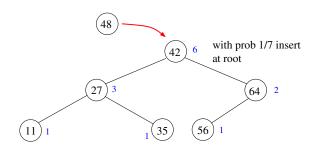
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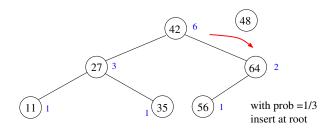
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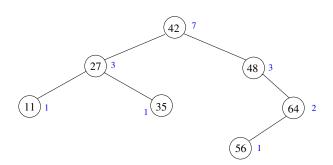
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Inserting an item x=48



Inserting an item x = 48



```
int size(node* p) {
    return (T == nullptr) ? 0 : T -> size;
}

void update_size(node* p) {
    if (p != nullptr)
        p -> size = size(p -> left) + size(p -> right) + 1;
}
```

• To insert a new item x at the root of T, we use the algorithm Split that returns two RBSTs T^- and T^+ with element smaller and larger than x, resp.

$$\begin{split} \langle T^-, T^+ \rangle &= \mathsf{Split}(T, x) \\ T^- &= \mathsf{BST} \text{ for } \{y \in T \,|\, y < x\} \\ T^+ &= \mathsf{BST} \text{ for } \{y \in T \,|\, x < y\} \end{split}$$

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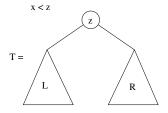
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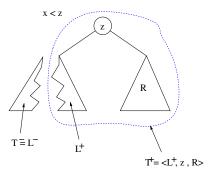
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To split a RBST T around x, we need just to follow the path from the root of T to the leaf where x falls



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Splitting a RBST & Insertion at Root

```
// splits the RBST T (destructively) into two trees, one with
// keys smaller than k, the other with keys larger than k
pair < node * , node * > split (node * T, const Key& k) {
  if (T == nullptr) return make_pair(nullptr, nullptr);
 if (k < T \rightarrow k) {
     pair < node *, node *> result = split(T -> left, k);
     T -> left = result.second;
     update size(T):
    result.second = T:
    return result;
  } else { // idem. change left <-> right
             // first <-> second
node* insert_at_root(node* T, const Key& k, const Value& v) {
  pair < node *. node *> LR = split(T, k): // $LR = \langle T^-, T^+\rangle$
  node* nn = new node(k. v):
  nn -> left = LR.first:
 nn -> right = LR.second:
 update_size(nn);
 return nn;
```

Splitting a RBST

Lemma

Let T^- and T^+ be the BSTs produced by $\operatorname{Split}(T,x)$. If T is a random BST containing the set of keys K, then T^- and T^+ are independent random BSTs containing the sets of keys $K^- = \{y \in T \mid y < x\}$ and $K^+ = \{y \in T \mid y > x\}$, respectively.

Insertion in RBSTs

Theorem

If T is a random BST that contains the set of keys K and x is any key not in K, then $\operatorname{Insert}(T,x)$ produces a random BST containing the set of keys $K \cup \{x\}$.

The Cost of Insertions

- The cost of the insertion at root (measured # of visited nodes) is exactly the same as the cost of the standard insertion
- For a random(ized) BST the cost of insertion is the depth of a random leaf in a random binary searh tree:

$$\mathbb{E}[I_n] = 2\log n + \mathcal{O}(1)$$

• We need to produce $\mathcal{O}(\log n)$ random numbers on average to insert an item

The Cost of Insertions

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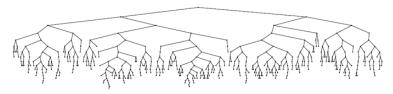
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RBST resulting from the insertion of 500 keys in ascending order Source: R. Sedgewick, Algorithms in C (3rd edition), 1997

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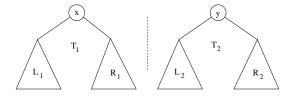
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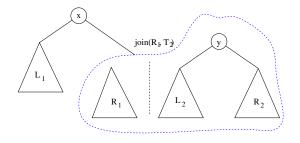
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- These studies showed that deletions degraded performance in the long run

```
node* remove(node* T, const Key& k) {
   if (T == nullptr) return nullptr;
   if (T -> k == k) { // to delete the root of the subtree, join the subtrees
      node* result = join(T -> left, T -> right);
      T -> left = T -> right = nullptr;
      free(T); // release node
      return result;
   }
   if (k < T -> k)
      T -> left = remove(T -> left, k);
   else
      T -> right = remove(T -> right, k);
   update_size(T);
   return T;
}
```

We delete the root using a procedure $\mathsf{Join}(T_1,T_2)$. Given two BSTs such that for all $x \in T_1$ and all $y \in T_2$, $x \le y$, it returns a new BST that contains all the keys in T_1 and T_2 .

$$\begin{aligned} \mathsf{Join}(\square,\square) &= \square \\ \mathsf{Join}(T,\square) &= \mathsf{Join}(\square,T) = T \\ \mathsf{Join}(T_1,T_2) &= ?, \qquad T_1 \neq \square, T_2 \neq \square \end{aligned}$$





- If we systematically choose the root of T_1 as the root of $\mathsf{Join}(T_1,T_2)$, or the other way around, we will introduce an undesirable bias
- Suppose both T_1 and T_2 are random. Let m and n denote their sizes. Then x is the root of T_1 with probability 1/m and y is the root of T_2 with probability 1/n
- Choose x as the common root with probability m/(m+n), choose y with probability n/(m+n)

$$\frac{1}{m} \times \frac{m}{m+n} = \frac{1}{m+n}$$
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Lemma

Let L and R be two independent random BSTs, such that the keys in L are strictly smaller than the keys in R. Let K_L and K_R denote the sets of keys in L and R, respectively. Then $T = \mathsf{Join}(L,R)$ is a random BST that contains the set of keys $K = K_L \cup K_R$.

- The recursion for $Join(T_1, T_2)$ traverses the rightmost branch (right spine) of T_1 and the leftmost branch (left spine) of T_2
- The trees to be joined are the left and right subtrees L and R of the ith item in a RBST of size n; then

length of left spine of L= path length to ith leaf length of right spine of R= path length to (i+1)th leaf

 The cost of the joining phase is the sum of the path lengths to the leaves minus twice the depth of the ith item; the expected cost follows from well-known results

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Theorem

If T is a random BST that contains the set of keys K, then $\mathsf{Delete}(T,x)$ produces a random BST containing the set of keys $K\setminus\{x\}$.

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Corollary

The result of any arbitary sequence of insertions and deletions, starting from an initially empty tree is always a random BST.

- Arbitrary insertions and deletions yield always random BSTs
- A deletion algorithm for BSTs that preserved randomness was a long standing open problem (10-15 yr)
- Properties of random BSTs have been investigated in depth and for a long time
- Treaps only need to generate a single random number per node (with $O(\log n)$ bits)
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- Storing subtree sizes for balancing is more useful: they can be used to implement search and deletion by rank, e.g., find the ith smallest element in the tree
- Other operations, e.g., union and intersection are also efficiently supported by RBSTs
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To learn more



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THANK YOU FOR YOUR ATTENTION!

