## Hashing

To store n keys, we need a table slightly bigger than n of size m

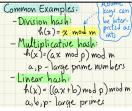
Load factor  $\lambda = n/m$ . Running time increases as  $\lambda \to 1$ 

 $\text{Hash Function}:\,h:\texttt{Keys}\to\mathbb{R}$ 

Scatters keys randomly and need to handle collision

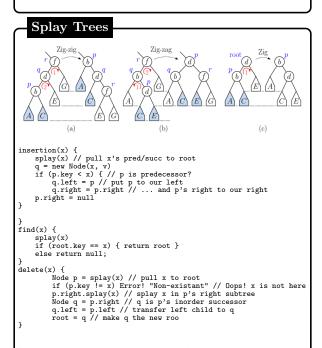
Good Hash Function: Efficient to compute, low collisions

Use every bit of the key and break up clusters



Modding by prime scatters keys, m may not be primed

If quadratic probe, table size m prime, |m/2| probe sequences distinct



Start with an empty dict, any sequence of m accesses take time

O(mlogn+nlogn) DFingerThm: Cost of a search starting from element y to x is the log of the of elements between

Working-Set Thm: if we accessed an element t steps ago, then the time to access it now is rougly log t.

## Treaps

Insertion times increase as we walk down the tree

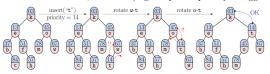
If keys are inserted in random order, expected height of BST O(logn).

Treap behave as if keys are inserted in random order

Timestamp increases from root to leaf. Assign random priority to keys

Treap has O(logn) expected height over all n! orderings

Insertion: BST insert, rotate left/right if parent priority is bigger



Deletion: Set node priority to  $\infty$ , rotate to leaf, delete

With standard binary search trees, the expectation was over all n! insertion orders. With treaps, the expectation was over all n! orders of the priority values. The latter is preferred, because the data structure's expected performance is not dependent on the insertion order

If just two keys have the same priority, their parent/child relationship might be affected, but the structure will be fine.

1) If 
$$a > b^k$$
, then  $T(n) = \Theta(n^{\log b})$ 

2) If 
$$a = b^k$$

a. If 
$$p > -1$$
, then  $T(n) = \Theta(n^{\log_b^a} \log^{p+1} n)$ 

b. If 
$$p = -1$$
, then  $T(n) = \Theta(n^{\log_b^a} \log \log n)$ 

c. If 
$$p < -1$$
, then  $T(n) = \Theta(n^{\log a})$ 

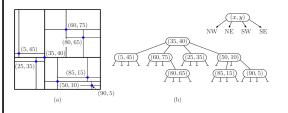
3) If  $a < b^k$ 

a. If  $p \ge 0$ , then  $T(n) = \Theta(n^k \log^p n)$ 

b. If p < 0, then  $T(n) = O(n^k)$ 

## QuadTrees

Each child is a division of a subspace/quadrant. In d-dimensions, we have  $2^d$  child. Above dim 3, these trees are too large



## Skip List

```
Total space = \sum_{i=0}^{\infty} n/2^i = 2n
Value forwardSearch(SkipNode p, Key y) {
   int i = 0 // start at the lowest level
   while (i >= 0) { // (level will rise then fall)
        if (p.next[i].key <= y) { // can we move forward?</pre>
            if (i+1 < p.next.length) i++ // move up, if possible
            else p = p.next[i] // move horizontal, if not
        else i-- // drop down a level
   return (p.key == y ? p.value : null) // return value if found
Value getMinK(int k) {
   int i = topmostLevel // start at topmost nonempty level
   SkipNode p = head // start at head node
   int count = k // number of items remaining
   while (count > 0) { // repeat until exhausting the span
        if (p.span[i] <= count) {</pre>
            count -= p.span[i] // decrement count by the number span
            p = p.next[i] // advance along same level
        }
        else i-- // drop down a level
   return p.key // return final element
```

Insertion: Save references on nodes where next is greater than the new value

In expectation the number of nodes visited at any level is 2

We start with n nodes at level 0, on average pn nodes survive to level 1,  $p^2n$  survive to level 2, and in general we expect  $p^in$  to survive to level i.

Let h denote the number of levels in the skip list. (We actually don't care what this value is.) Summing the number of nodes that contribute to each level of the skip list, and employing the fact that, for 0 < c < 1,  $\sum_{i=0}^{\infty} c^i = 1/(1-c)$ , the total expected number of links is

$$\sum_{i=0}^{h} p^{i} n \leq n \sum_{i=0}^{\infty} p^{i} = \frac{n}{1-p};$$

Observe that for any constant p, this is O(n).

## Kd-Tree

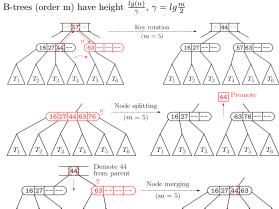
Deletion: Find minimum node in the right subtree with the minimum x/y according to cutdim. If right is empty, find min node in left and make left

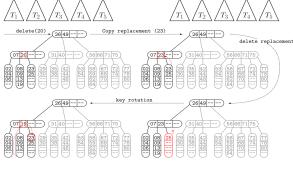
the new right. This can lead to  $O(\sqrt{n})$  height over large insert/deletes Splay: Any individual operation can be  $\Omega(n)$ 

If each element of a splay tree is accessed in ascending (or descending) order the total time for all these accesses is O(n). (Scanning theorem)

#### B-Trees

Designed so that each node fits in one page of memory B-tree: of order m ( $\geq 3$ ) Root of leaf or has  $\geq 2$  children Non root nodes have  $\lceil m/2 \rceil$  to m children, all leaves same level





# Misc

Guaranteed to succeed in finding slot: Linear Probing Double hashing, where the table size m and secondary hash function g(x) are relatively prime

```
TreapNode expose(Key x, TreapNode p) {
   if (p == null) // error - key not in tree
       throw Exception("Key not found");
    else if (x < p.key) { // x is smaller - search left
       p.left = expose(x, p.left);
       return rotateRight(p); // rotate the exposed node up
    else if (x > p.key) { // x is larger - search right
       p.right = expose(x, p.right);
       return rotateLeft(p); // rotate the exposed node up
    else { // found it
       p.priority = Integer.MIN_VALUE; // set priority to -infinity
       return p;
```

## Querying Kd-Tree

```
int rangeCount(Rectangle R, KDNode p, Rectangle cell) {
    if (p == null) return 0 // empty subtree
    else if (R.isDisjointFrom(cell)) // no overlap with range?
        return 0
    else if (R.contains(cell)) // the range contains our entire cell?
       return p.size // include all points in the count
    else { // the range stabs this cell
        int count = 0
       if (R.contains(p.point)) // consider this point
        count += 1
        // apply recursively to children
        count += rangeCount(R, p.left, cell.leftPart(p.cutDim, p.point)
        count += rangeCount(R, p.right, cell.rightPart(p.cutDim, p.poin
        return count
   }
Point rayShoot(Point q, KDNode p, Rectangle cell, Point best) {
    if (p == null) // fell out of tree?
        return best
    else if (cell.high.x < q.x || cell.high.y < q.y) // no overlap
       return best
    else {
        if (p.point.x >= q.x && p.point.y >= q.y && p.point.x < best.x)
            best = p.point // p.point is new best
        // get child cells
        Rectangle leftCell = cell.leftPart(p.cutDim, p.point)
        Rectangle rightCell = cell.rightPart(p.cutDim, p.point)
        if (cell.lo.y >= q.y && p.cutDim == 0) {
            if (p.point.x > q.x) // no need to search right
                best = rayShoot(q, p.left, leftCell, best)
            else // no need to search left
                best = rayShoot(q, p.right, rightCell, best)
       } else {
            best = rayShoot(q, p.left, leftCell, best)
            best = rayShoot(q, p.right, rightCell, best)
       }
    return best;
```

 $O(n^{1-1/d})$ , orthogonal range query time

## ScapeGoat Trees

Height will always be O(logn) because of rebuild event

m increment m for inserts, if m > 2n rebuild

 $\overline{\text{Anv}} m \text{ operations take } O(m log m), \text{ amortized} = O(log m)$ 

Must there be a scapegoat? The fact that a child has over 2/3 of the nodes of the entire subtree intuitively means that this subtree has (roughly) more than twice as many nodes as its sibling. We call such a node on the search path a scapegoat candidate. A short way of summarize the above process is "rebuild the scapegoat candidate that is closest to the insertion point.

You might wonder whether we will necessarily encounter an scapegoat candidate when we trace back along the search path. The following lemma shows that this is always the case.

**Lemma:** Given a binary search tree of n nodes, if there exists a node p such that depth(p) > $\log_{3/2} n$ , then p has an ancestor (possibly p itself) that is a scapegoat candidate

**Proof:** The proof is by contradiction. Suppose to the contrary that no node from p to the root is a scapegoat candidate. This means that for every ancestor node u from p to the root, we have  $size(u.child) \le \frac{2}{3} \cdot size(u)$ .

We know that the root has a size of n. Therefore, its child on the search path has size at most (2/3)n, its grandchild has size at most (2/3)((2/3)n) = (4/9)n, and generally the node at depth i along the search path as size at most  $(2/3)^{i}n$ .

Let d denote the depth of p. We know what its subtree rooted at p must have at least one node (namely p itself), and therefore

$$1 \le \text{size}(p) \le \left(\frac{2}{3}\right)^d n.$$

Solving for d, we have

$$\left(\frac{3}{2}\right)^d \le n \implies d \le \log_{3/2} n.$$

```
Node buildSubtree(Key[] A) { // A is a sorted array of keys
   k = A.length
   if (k == 0) return null // empty array
    else {
        j = floor(k/2) // median of the array
        Node p = new Node(A[j]) // ...this is the root
        p.left = buildSubtree(A[0..j-1])
        p.right = buildSubtree(A[j+1..k-1])
        return p // return root of the subtree
```

#### B+Trees

- Internal and leaf nodes are different in structure:
  - Internal nodes store kevs only, no values. The kevs in the internal nodes are used solely for locating the leaf node containing the actual data, so it is not necessary that every key appearing in an internal node need correspond to an actual key-value
- All the kev-value pairs are stored in the leaf nodes. There is no need for child pointers. (This also saves space.)
- Each leaf node has a next-leaf pointer, which points to the next leaf in sorted order.

Storing keys only in the internal nodes saves space, and allows for increased fan-out. This means the tree height is lower, which reduces number of disk accesses. Thus, the internal nodes are merely an index to locating the actual data, which resides at the leaf level, (The policy regarding which keys a subtree contains are changed. Given an internal node with keys  $\langle a_1, \dots, a_{j-1} \rangle$ , subtree  $T_j$  contains keys x such that  $a_{i-1} < x \le a_i$ .)

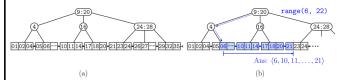


Fig. 9: B+ tree of order m=3, where leaves can hold up to 3 keys.

The next-leaf links enable efficient range reporting queries. In such a query, we are asked to list all the keys in a range  $[x_{\min}, x_{\max}]$ . We simply find the leaf node for  $x_{\min}$  and then follow next-leaf links until exceeding  $x_{max}$ .