CSE 220: Handout 29 Disjoint Sets

#### Motivation

- Suppose we have a database of people
- We want to figure out who is related to whom
- Initially, we only have a list of people, and information about relations is gained by updates of the form "alice is related to bob"
- Key property: If alice and bob are related, and bob and carol are related, then so are alice and carol
- Queries of the form: is alice related to bob?

#### Equivalence Relations

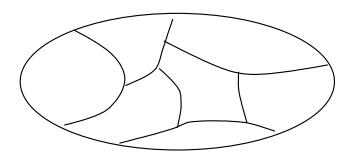
A binary relation R over a set S is called an equivalence relation if it has following properties

- 1. Reflexivity: for all element x, xRx
- 2. Symmetry: for all elements x and y, xRy if and only if yRx
- 3. Transitivity: for all elements x, y and z, if xRy and yRz then zRz

The relation "is related to" is an equivalence relation over the set of people

### The Disjoint Sets view

- ullet An equivalence relation R over a set S can be viewed as a partitioning of S into disjoint sets
- Each set of the partition is called an equivalence class of R (all elements that are related to each other according to R)

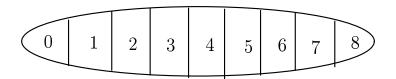


#### Union-Find Data Structure

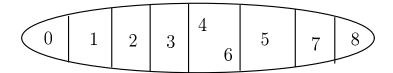
- The disjoint set is an abstract data type that supports two operations: union and find
- union(x,y) means merge the set containing x with the set containing y. In other words, declare x and y to be equivalent
- find(x) should return some representation of the set containing x.
- We can choose what find(x) returns as long as the following holds:
  find(x) == find(y) if and only if x and y are in the same set

## Sample Sequence

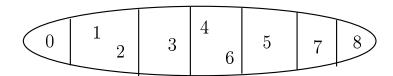
Initial



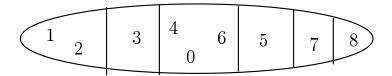
Union(4,6)



Union (1,2)



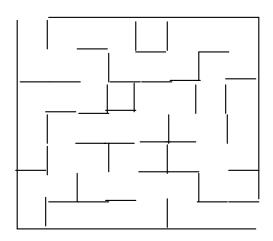
Union (0,4)



### Application: Maze generation

Goal: Knock down "enough" walls randomly so that cells 0 and 55 are connected

	0	1	2	3	4	5	6	7
	8	9	10	11	12	13	14	15
•								
_								
	48	49	50	51	52	53	54	55



### Algorithm

Initialize set S of all 56 cells as union/find ADT

### A Simple Implementation

- We will assume that elements are 0 to n-1
- Maintain an array A: for each element i, A[i] is the name of the set containing i
- find(i) returns A[i]: O(1)
- union(i,j) requires scanning entire array: O(n)

• In better implementations, union is less expensive (find will no longer be O(1), but overall performance will be improved)

#### Forest Data Structure

- Maintain the set S as a collection of trees, one per partition
- ullet Initially, there are n trees, each containing a single element
- find(i) returns the root of the tree containing *i*
- union(i,j) merges the trees containing i and j
- Typical tree traversal not required, so no need for pointers to children, instead we need a pointer to parent
- Parent pointers can be stored in an array: parent[i] (set to -1 if *i* is root)

### Tree-based Implementation

```
Initialization:
    for (i=0; i<n; i++)
        parent[i] = -1;

find(i):
    // traverse to the root
    for (j=i;
        parent[j]>=0;
        j=parent[j]);
    return j

union(i,j):
    root1 = find(i);
    root2 = find(j);
    if (root1 != root2)
        parent[root2] = root1;
```

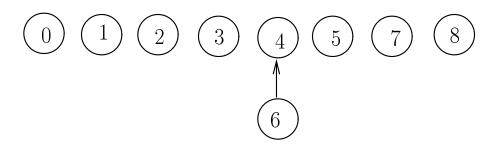
## Initially for n = 9

0	. 1	2	3	4	5	6	7	8
-1	-1	-1	-1	-1	-1	-1	-1	-1



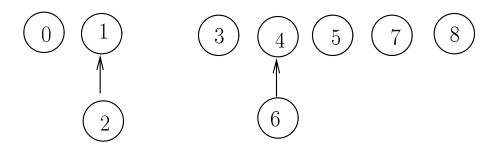
## After union(4,6)

0	. 1	2	3	4	5	6	7	8
-1	-1	-1	-1	-1	-1	4	-1	-1



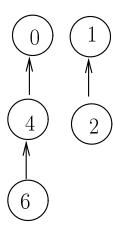
## After union(1,2)

0	. 1	2	3	4	5	6	7	8
-1	-1	1	-1	-1	-1	4	-1	-1



# After union(0,4)

0	. 1	2	3	4	5	6	7	8
-1	-1	1	-1	0	-1	4	-1	-1



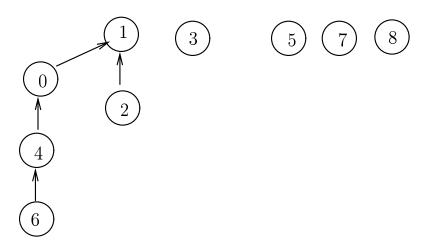






# After union(2,4)

0	. 1	2	3	4	5	6	7	8
1	-1	1	-1	0	-1	4	-1	-1



### Running Time Analysis

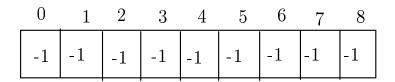
- Running time of find(i) is proportional to the height of the tree containing node *i*
- Running time of an operation is proportional to the maximum height of a tree in the forest
- This can be O(n) in the worst case (but not always)
- Goal: Modify union to ensure that heights stay small

#### Union by Size

- Maintain sizes of all trees, and during union make smaller tree the subtree of the larger one
- Implementation: for each root node i, instead of setting parent [i] to -1, set it to -k if tree rooted at i has k nodes
- Modify the code for union on p. 11

```
union(i,j):
    root1 = find(i);
    root2 = find(j);
    if (root1 != root2)
        if (parent[root1] <= parent[root2]) {
            // first tree has more nodes
            parent[root1] += parent[root2];
            parent[root2] = root1; }
        else { // second tree has more nodes
            parent[root2] += parent[root1];
            parent[root1] = root2; }</pre>
```

### Sample Execution



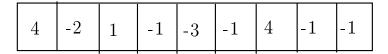
Union (4, 6)

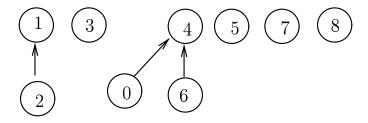


Union (1, 2)



Union (0,4)



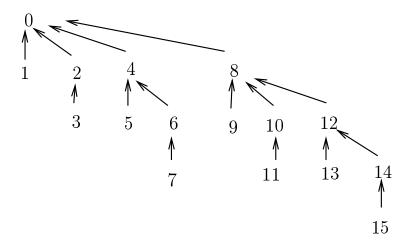


### Running Time Analysis

- Claim: In union-by-size strategy, depth of a node cannot be more than log n
- Each time the depth of a node *i* increases by 1, the size of the tree containing *i* is at least doubled.
- Thus, running time of each operation is log n
- Average over a sequence of operations is almost constant: that is, a sequence of M union/find operations takes a total of O(M) time on average
- Alternative to union-by-size strategy: maintain heights, and during union, make a tree with smaller height a subtree of the other

#### The worst-case scenario

• Worst-case scenario for 16 union operations



- Improving union won't change performance for this sequence of operations (why?: ties must be broken one or the other)
- Can we optimize **find**?

#### Path Compression

- During find(i), as we traverse the path from i to root, update parent entries for all these nodes to the root
- This reduces the heights of all these nodes
- Pay now, and reap benefits later!
  Subsequent find may do less work
- Updated code for find

```
find (i) {
  if (parent[i] < 0)
    return i;
  else return parent[i] = find (parent[i]);
}</pre>
```