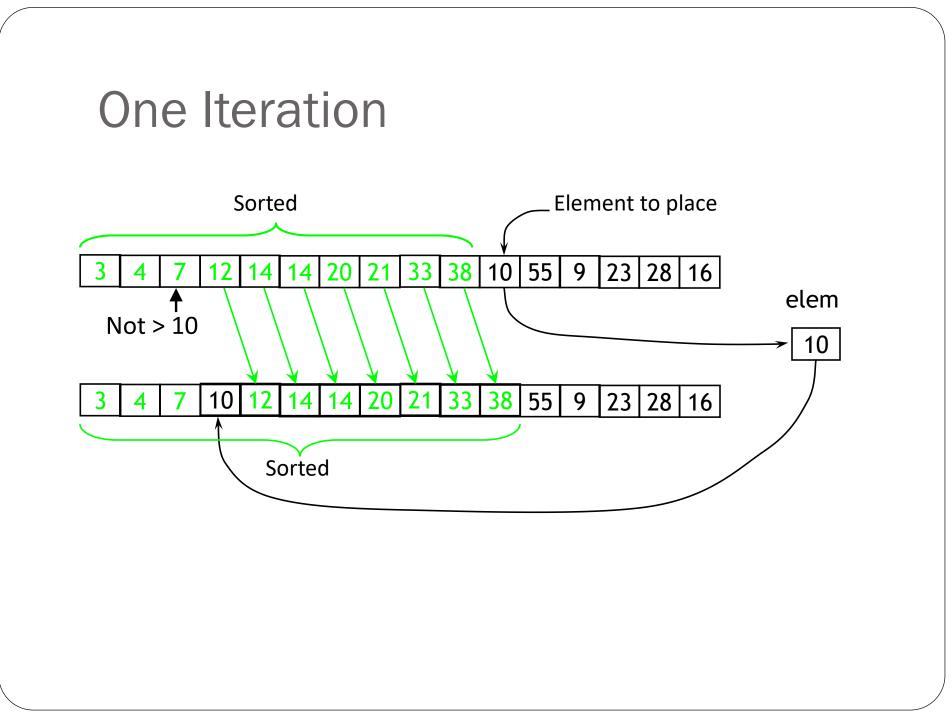
Improving on Insertion Sort

Shell Sort

Insertion Sort

Fast (O(N)) when sequence nearly sorted; otherwise s...l...o...w

```
for (idxToInsert = 1; idxToInsert < v.size ();</pre>
     ++idxToInsert)
{
  k = idxToInsert;
  elem = v[k];
  while (k \ge 1 \&\& e e \le v[k - 1])
  {
    v[k] = v[k - 1];
    k = k - 1;
  }
  v[k] = elem;
}
```



How to Improve?

- Each time we insert an element other elements get nudged *one step* closer to where they ought to be
- What if we move elements a *much longer distance* each time?
- Move each element long distances initially, and decrease that distance to 1 eventually
 - This leads to *Shell sort*

Sorting Subsequences

Vector to be sorted



- Insertion sort red elements
- Insertion sort yellow elements ...
- ... and finally purple elements
- Resultant array is sorted?

Elements Compared

0, 5, 10, 15, 20 1, 6, 11, 16, 21 2, 7, 12, 17, 22 3, 8, 13, 18, 23 4, 9, 14, 19

h-Sorting

- We sorted 5 sequences of elements spaced 5 apart a (single) *h-sort* with h=5
 - Insertion sort is a 1-sort
- What if we follow the 5-sort with a 1-sort?
 - Expect each insertion would involve moving fewer elements
 - Resulting vector would be sorted

Values of 'h'

- For large vectors we don't want to start with a 5-sort
- Start with h = f (v.size ())
- Reduce h to 1
- Values of h form *increment* or *decrement sequence*

Values of 'h'

- Hibbard suggests $< 1, 3, 7, ..., 2^{k} 1 >$
 - To find initial 'h': for (h = 1; h <= N / 4; h = h * 2 + 1) /* empty */;
 - Repeat while h > 0
 - Do h-sort
 - h = h / 2
 - Worst case O(N^{1.5})

Increment Sequences

- Performance sensitive to increment/decrement sequence
- Optimal sequence not known
 - Shell proposed decrement seq. < 1, 2, 4, 8, ... >
 - Good?
 - One from Donald Knuth: < 1, 4, 13, ... >
 - Decrement by dividing by 3
 - <u>Many others...</u>
- So what does the code for h-sorting look like?

Analysis

- You cut the size of the array, N, by some fixed amount (N = N / k)
- Consequently, you have about log N stages
- Each stage takes **O(N)** time
- Hence, the algorithm takes **O(N log N)** time
- Right?

Analysis

• Wrong!

- This analysis assumes that each stage actually moves elements closer to where they ought to be, by a fairly large amount
- What if all the red cells, for instance, contain the largest numbers in the array?

• In fact, if we just cut the size in half each time, sometimes we get $O(N^2)$ behavior!

Analysis

• What is the *real* complexity?

• Depends on sequence

• Sometimes unknown

• Some complexities determined empirically