# Should we Teach Parallelism Throughout our Data Structures and Algorithms Courses?

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# Why bother?

Parallelisms works even on modest machines

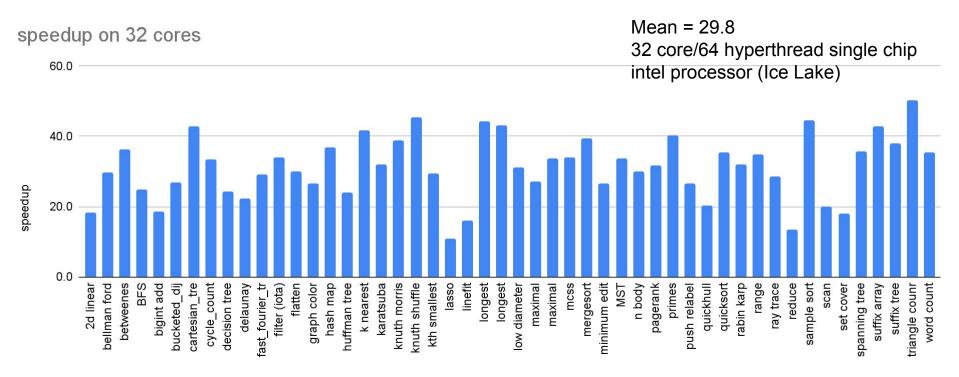
Hard to talk about scalability without parallelism

In many cases more natural and simple to think parallel

Selfish reason: To help Guy, Kunal, Yihan ... get well educated graduate students

#### Lead to good efficiency?

Natural question: do ideas taught in corse help in practice



### How hard is it?

We (at CMU) have been teaching parallelism in our intro (sophomore level) algorithms + data structure course (500 students/year) for 12 years.

• Not just a 2 lecture add on

**Outline**: Will give an overview of what we do and lessons learned.

**Feedback**: Would love to know any blocking factors for others, or what others are doing, and other ideas.

# Philosophy

At the right level of abstraction there is no or little fundamental difference between parallel and sequential data structures and algorithms

- Need to recognize when parallelism is already present
- Need to understand how to make solutions more parallel
- Difference is only in costs and some added techniques

Emphasis on "**Parallel Thinking**": i.e., ideas that are "natural" and transcend the particular model.

#### **Pros and Cons**

Pros:

- Do not have to diverge much from standard algorithms course structure
- Emphasizes that parallelism is not esoteric
- Learn to think about parallelism abstractly and naturally

Cons:

• Requires revising an algorithms course throughout (significant work for faculty), although by how much is up to the instructor.

# **Course Outline**

- Cost models and analysis
- Techniques: D&C, brute force, ...
- Algorithms on sequences: merging, mergesort, scan, ...
- Randomization: quicksort, ...
- BSTs and Priority Queues
- Graph algorithms (search, shortest paths, MSTs, ...)
- Dynamic programming

Traditional at this level of detail

#### **Parallel Model**

(e1 || e2) : evaluate e1 and e2 in parallel, returning a pair when both are done

Sequence operations: e.g.

- Map : [f(x) : x in S]
- Tabulate : [f(i) : 0 <= i < n]
- Filter : [x in S | f(x)]

All calls to f are in parallel, returning a sequence

Avoid race conditions (schedule/interleaving) does not matter

#### Sidebar: easy to e.g. express parallelism in python

Even if implementation does not support it, e.g.,

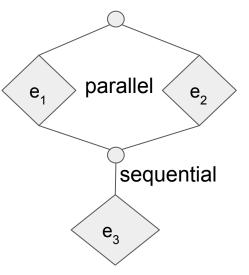
for i in range(n)  $\Rightarrow$  [f(x) for x in S] r[i] = f(S[i])

r = 0for x in S: r = r + xreduce(add, S)

#### Cost Model : Work and Span

Work: Total number of operations (like sequential time) Span: Critical path (assuming unbounded number of processors)

	Sequential composition	Parallel composition
Work	add	add
Span	add	max



No PRAM

#### Cost model

Priorities:

- Most important is to reduce the work
- Then to reduce the span ignore logarithmic factors.

Parallelism = Work/Span (roughly # of processors can make use of)

Some discussion of scheduling, leading to:

Time = Work/Processors + O(Span). for "greedy" schedulers

# Recurrences and Big-O

Same as sequential algorithms but for work and span instead of time.

E.g. for mergesort:

- $W(n) = 2W(n/2) + O(n) = O(n \log n)$  using sum over recursive calls
- $S(n) = S(n/2) + O(\log n) = O(\log^2 n)$  using max over recursive calls

Assuming for merge W(n) = O(n) and  $S(n) = O(\log n)$ 

# Techniques

- Brute force
- Divide-and-conquer
- Contraction
- Graph search
- Dynamic programming

## Techniques

- Brute force (note that naturally parallel)
- Divide-and-conquer (naturally parallel, and use more aggressively)
- Contraction (not usually covered in sequential algorithms)
- Graph search (BFS is parallel but not DFS)
- Dynamic programming (view more abstractly as a DAG)

# **Divide and Conquer**

Same as with sequential, but:

- Recurrences for span
- More emphasis (use where iteration is used sequentially), e.g., reduce, parenthesis matching, MCSS
- Often need to strengthen induction

#### Divide and Conquer: reduce

Taking the sum with respect to a binary associative function f

```
reduce f identity S =
  Case splitMid(x) of
  Empty => return identity
  | Single(x) => return x
  | Pair(L, R) => return f(reduce f identity L || reduce f identity R)
```

 $W(n) = 2W(n/2) + W_f = O(n)$  assuming  $W_f = S_f = O(1)$  $S(n) = S(n/2) + S_f = O(\log n)$ 

Would normally do iteratively in sequential setting.

# Divide and Conquer: mergeSort

Taking the sum with respect to a binary associative function f

mergesort S =
 case splitMid(x) of
 Empty => return []
 | Single(x) => S
 | Pair(L, R) => return merge(mergesort L || mergesort R)

 $W(n) = 2W(n/2) + O(n) = O(n \log n)$  $S(n) = S(n/2) + \log n = O(\log^2 n)$ 

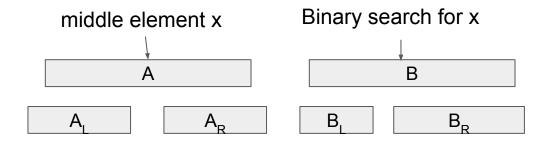
Assuming for merge: W(n) = O(n),  $S(n) = O(\log n)$ 

# Divide and Conquer: mergeSort

Or:

Mergesort S = reduce(merge,[], [[x] : x in S])

# Divide and Conquer: Merging



If use dual binary search:

 $W(n) = 2W(n/2) + O(\log n) = O(n)$  $S(n) = S(n/2) + O(\log n) = O(\log^2 n)$ 

Very different from sequential algorithm. Can get  $S(n) = O(\log n)$  with more advanced approach.

#### **Divide and Conquer: Parenthesis Matching**

Do these parentheses match: (()(())())(()))(()())

**Strengthen the induction** to return number of unmatched parentheses on left and right:

```
parenMatch' S=
Case split_mid(x) of
Empty => return (0,0)
| Single(x) => if x = '(' then return (0,1) else return (1,0)
| Pair(L,R) =>
((la,lb), (ra,rb)) = (pareMatch' L, parenMatch' R)
if (lb > ra) then return (la, lb-ra + rb)
else return (la + ra - lb, rb)
```

```
parenMatch S =
(a,b) = ParenMatch' S
return (a = 0) and (b = 0)
```

## Technique: Contraction

Basic structure:

- Reduce problem to one that is a constant factor smaller
- Recurse on the problem
- Use result to solve the problem

Some applications: Scan, kthSmallest, Graph Contraction

Not usually covered in intro algorithms/data structure course, but important for parallelism.

## Contraction Example: Scan (prefix sums)

For binary associative operator f, return "running sum" of previous elements

```
e.g.: Scan + 0 [2, 1, 3, 1, 2] -> [0, 2, 3, 6, 7], 9
```

Seems inherently sequential

Parallel version has many applications: E.g.: filter, flatten, carry propagation, skyline, ...

# Scan: using contraction

```
scan(f, identity, S) =

If |A| = 0 then return ([], identity)

else if |A| = 1 then return ([identity], S[0])

else

pairSums = [ f(A[2i],A[2i+1]) : 0 <= i < |S|/2 ]

(r,s) = scan(f, identity, pairSums)

return [if (i%2 =0) then r[2i] else r[2i] + a[i]]
```

W(n) = W(n/2) + O(n) = O(n) $S(n) = S(n/2) + O(1) = O(\log n)$ 

Assuming f is constant work:

// pairwise sum the elements// recurse// put together results

### Randomized algorithms

Analyze:

- expectation for work (e.g. E[W<sub>quicksort</sub>(n)] = O(n log n))
- high probability bounds for span (e.g.  $S[W_{quicksort}(n)] = O(\log^2 n) w.h.p.$ )

Problem is there is no equivalent of linearity of expectation for maximum.

Example of students taking an exam.

#### Randomized algorithms: kthSmallest

kthSmallest(A, k) =
 p = pick an element of A uniformly at random
 L = [x in A | x < p]
 R = [x in A | x > p]
 if (k < |L|) then return kthSmallest(L, k)
 else if (k < |A| - |R|) then return p
 else return kthSmallest(R, k - (|A| - |R|))</pre>

Show that number of recursive calls is  $O(\log n)$  with high probability Also work is O(n) in expectation.

#### Sidebar: Deteministic kthSmallest

W(n) = W(7n/10) + W(n/5) + O(n) = O(n)

 $S(n) = S(7n/10) + S(n/5) + O(1) = O(n^{.82..})$ 

Parallelism =  $O(n^{.18})$ 

#### Randomized algorithms: Quicksort

```
quicksort(A) =
    p = pick an element of A uniformly at random
    L = [x in A | x < p]
    R = [x in A | x > p]
    (L', R') = (quicksort(L) || quicksort(R))
    return L ++ [p] ++ R
```

Work =  $O(n \log n)$  in expectation Span =  $O(\log^2 n)$  with high probability

### Binary search trees:

Traditionally teach insertion, deletion, find on sets or dictionaries

None of these have parallelism.

Extend to support:

- Filter, reduce, multi-insert, ...
- Union, intersection, ...

Allows for bulk operations on sets and dictionaries (Parallel Thinking)

Can all be built up from "join", and are highly parallel

## Join for treaps

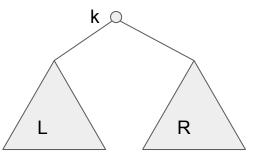
```
join(A, key, B) =
if (p(k(A)) < p(key) and p(k(B)) < p(key)) then
return Node(A ,key ,B)
else if (p(k(A)) > p(k(B))) then
return Node(L(A), key, join(R(A) ,k(A), B))
else
return Node(join(A, k(B), L(B)), key, R(B))
```

```
joinPair(A,B) and
split(A, k) => (AL, ?, AR)
can easily be built on this
```

```
W(n) = S(n) = O(\log n) whp
```

#### Binary search trees: Filter example

```
filter f A =
   case A of
    Leaf => A
    | Node(L,k,R) =>
      (L', R') = (filter f L, filter f R)
      if (f(k)) return join(L',k,R')
      else return joinPair(L', R')
```



Work = O(n)Span =  $O(\log^2 n)$  whp

Join handles the rebalancing. We use treaps for balancing, but could use AVL trees or Red Black trees.

#### Binary search trees: Union

union A B = case (A,B) of (Leaf,\_) => B | (\_, Leaf) => A | (Node(AL,k,AR),\_) => (BL,\_,BR) = split(B, k) (L', R') = (union(AL,BL) || union(AR,BR)) return join(L',k,R')

В

BR

Work =  $O(m \log (n/m + 1))$ Span =  $O(\log^2 n)$ 

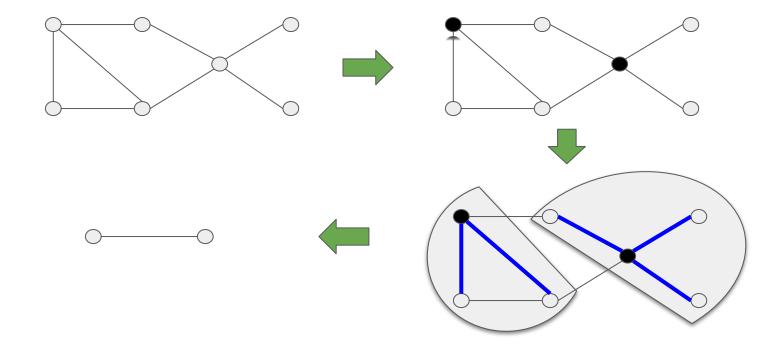
# Graphs:

30% of course is on graphs

Some key points:

- BFS is parallel, span is O(d log n), where d is the diameter of the graph
- DFS and Dijkstra are sequential
- Bellman Ford is parallel: O(mn) work and O(n log n) span.
- Johnson is parallel
- Use graph contraction for connectivity and Min Spanning Tree MST)
   MST uses Boruvka's algorithm, but we also mention Kruskal and Prim

#### Graphs: Example of star contraction



#### Graphs: Star contraction

In one round remove at least  $\frac{1}{4}$  of the vertices in expectation.

Therefore finishes in O(log n) rounds with high probability.

Total work is O(m log n)

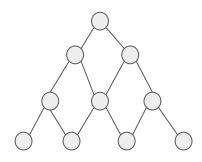
Used for both Connectivity and Boruvkas MST algorithm

# **Dynamic Programming:**

To expose parallelism, we raise level of abstraction.

DP is recursion with sharing. Sharing can be viewed as a DAG.

Recursion Tree -> Dag



Calculate total work across all recursive calls.

Calculate longest path in DAG (same as span for non-shared solution)

# **Dynamic Programming:**

Some examples we cover:

- Subset sum: W(n,k) = O(nk), S(n, k) = O(n)
- Minimum edit distance: W(n,m) = O(nm), S(n,m) = O(n + m)
- Optimal BST:  $W(n) = O(n^3)$ ,  $S(n) = O(n \log n)$
- Matrix chain Product:  $W(n) = O(n^3)$ ,  $S(n) = O(n \log n)$

Discuss bottom up and top down approaches.

#### Lessons Learned

- **Deterministic** parallelism not difficult for the students to understand.
  - They get the hang quickly.
- List of "new" ideas that need to be covered, and which are easier/harder
- resistance by faculty to changing existing courses
  - faculty often not comfortable with topic
- best to do along with considering whole curriculum

Have not yet done a careful study

#### Lessons Learned: effect on whole curriculum

- ideally requires changes in prerequisites
  - parallel thinking earlier
  - programming with comprehensions and functional style earlier
  - tail bounds in probability
- changes to follow up courses
  - how to cover parallelism in Senior level algorithms class?
  - where to cover concurrent and distributed algorithms?
- important to have complementary course on systems issues
  - $\circ$  gpus, distributed computing, locks, ...

# Why are sequential data structures and algorithms so successful?

- abstract away from detail
- broadly applicable
- general techniques
- simple cost model
- easy to program
- elegant and often simple
- help understand scaling, i.e., big-O
- lead to good efficiency in practice
- interesting theory

#### Conclusions

Parallel and sequential algorithms can be integrated such that students

- consider parallelism throughout
- learn "standard" algorithmic topics, e.g. D&C, DP, mergesort, quicksort, BFS, DFS, …
- learn to think parallel, e.g., ask if an algorithm has parallelism, or if and how it can be added

Including in an existing UG curriculum has challenges