# CSE Qualifying Exam: High-Performance Computing

### Spring 2020

#### Instructions

- Please answer three of the following four questions. All questions are graded on a scale of 10. If you answer all four, all answers will be graded and the three lowest scores will be used in computing your total.
- Please write clearly and concisely, explain your reasoning, and show all work. Points will be awarded for clarity as well as correctness.
- Unless otherwise specified, assume a distributed memory model of computation where a message of *m* words can be sent in O(τ + μm) time, where τ denotes the latency and μ denotes the per word transfer time. You may assume each processor can send and receive one message within the same parallel communication step.

## Problem 1

Suppose we have p interconnected processors, each with its own memory. Consider n particles where the position of each particle in 3-D space is known. Space is partitioned into p cells. The positions of the particles lying in the kth cell are stored only on processor k. If a pair of particles i and j are separated by a distance d less than a given distance r, we must compute the equal-and-opposite force on particle i and on particle j. This force depends on d and is used to update the position of particles i and j.

(a) Assuming a uniform random distribution of particles in a bounded 3-D domain, derive an asymptotic lower bound on the interprocess communication volume required to update the positions of all *n* particles.

(b) Give an algorithm that attains this lower bound.

# Problem 2

The Hilbert space filling curve is a way to convert coordinates (x, y) into an index h such that two points with consecutive indices are always near each other in the plane.

We can write a Hilbert index as an integer in base 4,  $h = h_1 h_2 \cdots h_n$ , where each  $h_i \in \{0, 1, 2, 3\}$ .

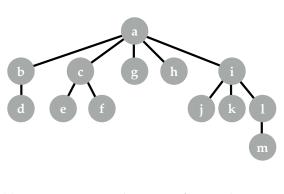
Decoding a Hilbert index can be done recursively: we start with  $x_0 = y_0 = 0$ , and we have four different update rules:

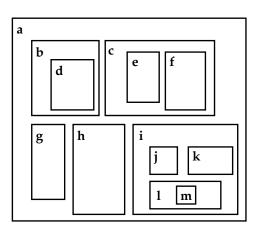
- if  $h_i = 0, x_i \leftarrow y_{i-1}/2, y_i \leftarrow x_{i-1}/2;$
- if  $h_i = 1, x_i \leftarrow (1 + x_{i-1})/2, y_i \leftarrow y_{i-1}/2;$
- if  $h_i = 2$ ,  $x_i \leftarrow (1 + x_{i-1})/2$ ,  $y_i \leftarrow (1 + y_{i-1})/2$ ;
- if  $h_i = 3$ ,  $x_i \leftarrow (1 y_{i-1})/2$ ,  $y_i \leftarrow (2 x_{i-1})/2$ .

Write a parallel algorithm for decoding a Hilbert index. Your algorithm should run in  $O(\log n)$  time.

### Problem 3

**Parallel HTML rendering.** When a web browser downloads a web page, it receives it in HTML format. Let's assume a simplified form of HTML: the downloaded document is represented as a tree of page elements, as shown in Figure 1a. Each vertex is a *page element*; and page elements may be *nested*. If v is an element, let P[v] be its *parent* and let C[v] be the set of its children. For instance, in Figure 1a,  $C[a] = \{b, c, g, h, i\}$  and P[d] = b.





(a) Raw HTML, viewed as a tree of "page elements"

(b) Illustration of a rendered HTML page

Figure 1: Parallel HTML rendering problem

Given such a tree, the browser needs to *render* it, meaning to lay it out physically on the output device, such as a screen or printed page. Figure 1b shows a hypothetical rendering. When fully rendered, every element v has known width and height and a known absolute position on the output device. Let's use  $S_v$  as a shorthand for this "state" of v. The problem is that the input tree (Figure 1a) initially holds only the nesting relationships and the content of each element; it does *not* have any size or position information, other than the nesting structure. In this scenario, we say  $S_v = \emptyset$ , to denote the initially unknown state.

However, suppose you have two special operations that can help resolve  $S_v$ .

- $S_v = \text{place}(S_a, S_b)$ : Let *a* and *b* be any pair of direct siblings, that is P[a] = P[b] = v, with known dimensions in  $S_a$  and  $S_b$ . Then this function will decide how to place these together (e.g., next to each other, on top of each other, or some other arrangement) and returns a new combined state,  $S_v$ , that captures this arrangement and their *relative* positions. You can think of the combined state as a kind of "virtual child" of *v*, so that one may compose placements as place(place(S,T),U) = place(place(S,U),T).
- S'''<sub>v</sub> = merge(S'<sub>v</sub>, S''<sub>v</sub>): Given two partial renderings of v, as might be produced by place(·, ·), this operation reconciles them into a single rendering. That is, S'<sub>v</sub> might represent a partial rendering with some of the children of v while S''<sub>v</sub> does so for a different set of v's children. This operation will combine them. Similar to the above, you may further assume that merge(S', merge(S'', S''')) = merge(merge(S', S''), S''').

Lastly, assume that for any leaf w (i.e.,  $C[w] = \emptyset$ ), the size is easy to compute by a call to  $place(\emptyset, w) = place(w, \emptyset)$ . You may also assume that the above operations cost O(1) time each. If you need more assumptions, state them clearly.

Please answer the following questions.

- a. (70%) Give an efficient parallel algorithm to render the page, that is, to fully resolve  $S_v$  for all v. "Efficient" in this case means with a total work that is as asymptotically close as you can manage to O(n) and a span or depth that is polylogarithmic in n. Analyze your algorithm.
- b. (30%) Critique the assumptions of this problem. That is, which ones pose the "biggest threat" to the efficiency or correctness of your approach, and why?

# Problem 4

#### Scheduling.

- a. Scheduling for a Single Processor: Consider n jobs  $\{j_1, j_2, ..., j_n\}$  that will arrive for execution on a single processor, one after each other. Let their arrival times be  $\{a_1, a_2, ..., a_n\}$ , respectively, with  $a_i \le a_{i+1}$ . Also let  $\{t_1, t_2, ..., t_n\}$  be the respective execution times on that processor. If the processor is free when job  $j_i$  arrives, job starts without any wait and executes for  $t_i$  unit and it exits. Otherwise, job waits in the queue for its turn for execution (in the order it was received). Assume  $a_i$ 's and  $t_i$ 's are distributed evenly across p processors. Design and efficient parallel algorithm to determine the waiting time incurred by each job. (60%)
- b. *Scheduling for Multiple Processors:* Now consider, we have m identical processors that jobs can be executed. Discuss the changes you would do to your algorithm; how they effect the execution time of your algorithm. (40%)