Chapter 7

Algorithm-Design Technique: Contraction

Contraction, an inductive technique for designing parallel algorithms, is probably one of the most important algorithm-design techniques. Like divide-and-conquer algorithms, contraction algorithms involve solving a smaller instance of the same problem, but unlike in divide-and-conquer, there is only one subproblem to solve at a time. A contraction algorithm for problem P has the following structure.

Base Case: If the problem instance is sufficiently small, then compute and return the answer, possibly using another algorithm.

Inductive Step: If the problem instance is sufficiently large, then apply the following three steps (possibly multiple times).

- 1. *Contract:* "contract", i.e., map the instance of the problem *P* to a smaller instance of *P*.
- 2. *Solve:* solve the smaller instance recursively.
- 3. *Expand:* use the solution to solve the original instance.

Contraction algorithms have several nice properties. First, their inductive structure enables establishing correctness and efficiency properties in a relatively straightforward fashion using principles of induction. For example, to prove a contraction algorithm correct, we first prove correctness for the base case, and then prove the general (inductive) case by using strong induction, which allows us to assume that the recursive call is correct. Similarly, the work and span of a contraction algorithm can be expressed as a recursive (inductive) relation, which essentially reflects the structure of the algorithm itself. We then establish the bounds for work and span by using known, well-understood techniques for solving recursive relations as briefly discussed in Chapter 4. Second, contraction algorithms can be efficient, if they can reduce the problem size geometrically (by a constant factor greater than 1) at each contraction step and if the contraction and the expansions steps are efficient. Similarly, if size of the problem instance decreases geometrically, and if contraction and expansion steps low spans, then the algorithm will have a low span, and thus high parallelism.

In this chapter, we will consider several applications of the contraction technique and analyze the resulting algorithms.

7.1 Example 1: Implementing Reduce with Contraction

As our first example, we describe how to perform reduction on a sequence by using contraction. Recall that the type signature for reduce is as follows.

```
reduce (f: \alpha \times \alpha \rightarrow \alpha) (id: \alpha) (A: \mathbb{S}_{\alpha}): \alpha
```

where f is an associative function, A is the sequence, and id is the identity element of f. Even though we have defined reduce broadly for both associative and non-associative functions, throughout this chapter, we restrict ourselves the case with associative functions, i.e., we assume that the function f is associative.

The crux in using the contraction technique is to design an algorithm for reducing an instance of the problem to a geometrically smaller instance by performing a parallel contraction step. In the case of reduce, there are many ways to perform such a contraction. For example, we can reduce the problem to a smaller problem by removing the first element of the sequence. This approach, however, would not result in a geometric reduction in the size of the problem instance. Consider instead applying the function f to consecutive pairs of the input. For example if we wish to compute the sum of the input sequence

```
\langle 2, 1, 3, 2, 2, 5, 4, 1 \rangle
```

by using the addition function, we can contract the sequence to

```
\langle 3, 5, 7, 5 \rangle.
```

By using this contraction step, we have reduced the input size by a factor of two. Note that the contraction step can be performed in parallel, because each pair can be considered independently in parallel.

Having reduced the size of the problem with the contraction step, we next solve the resulting problem by invoking the same algorithm and apply expansion to construct the final result. It is not difficult to see that by solving the smaller problem, we have actually solved the original problem, because the sum of the sequence remains the same as that of the original. Thus, the expansion step requires no additional computation. We can thus express our algorithm as follows; for simplicity, we assume that the input size is a power of two.

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Algorithm 7.1. [Reduce with contract] (* Assumption: |A| is a power of 2 *) reduce_contract f id A =if isSingleton A then A[0]else let $B = \langle f(A[2i], A[2i+1]) : 0 \le i < \lfloor |A|/2 \rfloor \rangle$ in reduce_contract f id B end

We can usually express the work and span of a contraction algorithm using a recursive relation. Assuming that the function being reduced over performs constant work, parallel tabulate in the contraction step requires linear work, we can thus write the work of this algorithm as follows.

```
W(n) = W(n/2) + n.
```

It is not difficult to solve this recursive formula to prove that the algorithm therefore performs O(n) work. Assuming that the function being reduced over performs constant span, parallel tabulate in the contraction step requires constant span; we can thus write the work of this algorithm as follows.

$$S(n) = S(n/2) + 1.$$

The algorithm therefore performs $O(\log n)$ span.

7.2 Example 2: Implementing Scan with Contraction

As an even more interesting example, we describe how to implement the scan sequence primitive efficiently by using contraction. Recall that the scan function has the type signature

```
scan (f: \alpha \times \alpha \rightarrow \alpha) (id: \alpha) (A: \mathbb{S}_{\alpha}) : (\mathbb{S}_{\alpha} \times \alpha)
```

where f is an associative function, A is the sequence, and id is the identity element of f. When evaluated with a function and a sequence, scan can be viewed as applying a reduction to every prefix of the sequence and returning the results of such reductions as a sequence. For example, applying scan ` '+', i.e., "plus scan" on the sequence (2,1,3,2,2,5,4,1) returns

```
(\langle 0, 2, 3, 6, 8, 10, 15, 19 \rangle, 20).
```

We will use this as a running example.

Based on its specification, a direct algorithm for scan is to apply a reduce to all prefixes of the input sequence. Unfortunately, this easily requires quadratic work in the size of the input sequence. We can see that this algorithm is inefficient by noting that it performs lots of redundant computations. In fact, two consecutive prefixes overlap significantly but the algorithm

```
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```

does not take advantage of such overlaps at all, computing the result for each overlap independently. By taking advantage of the fact that any two consecutive prefixes differ by just one element, it is not difficult to give a linear work algorithm (modulo the cost of the application of the argument function) by using iteration. Such an algorithm may be expressed as follows

$$\operatorname{scan} f id A = h \left(\operatorname{iterate} g \left(\langle \rangle, id \right) A \right),$$

```
where g((B,b),a)=((\operatorname{append}\langle\,b\,\rangle\,B),f(b,a)) and
```

the function h(B,b) = ((reverse B),b), where reverse reverses a sequence. This algorithm, while correct, is almost entirely sequential, leaving no room for parallelism.

Considering the fact that it has to compute some value for each prefix, it may seem difficult to give a parallel algorithm for scan. At first glance, we might be inclined to believe that any efficient algorithms will have to keep a cumulative "sum," computing each output value by relying on the "sum" of the all values before it. It is this apparent dependency that makes scan so powerful. Indeed, we often use scan when it seems we need a function that depends on the results of other elements in the sequence.

We can implement scan efficiently using contraction. To this end,, we need to reduce a given problem instance to a geometrically smaller instance by applying a contraction step. As a starting point, let's apply the same idea as we used for reduction in Algorithm 7.1 and see how far that takes use. Applying the contraction step from the reduce algorithm described above, we would reduce the input sequence

```
\langle 2, 1, 3, 2, 2, 5, 4, 1 \rangle
```

to the sequence

$$\langle 3, 5, 7, 5 \rangle$$
,

which if recursively used as input would give us the result

```
(\langle 0, 3, 8, 15 \rangle, 20).
```

Notice that in this sequence, the elements in the first, third, etc., positions are actually consistent with the result expected:

```
(\langle 0, 2, 3, 6, 8, 10, 15, 19 \rangle, 20).
```

The reason for why half of the elements is correct is because the contraction step which pairs up the elements and reduces them, does not affect, by associativity of the function being used, the result at the position that do not fall in between a pair. Thus, we can use an expansion step to compute the missing items. It is actually quite simple: all we have to do is to compute the missing elements by applying the function element-wise to the elements of the input at odd positions in the input sequence and the results of the recursive call to scan.

To illustrate, the diagram below shows how to produce the final output sequence from the original sequence and the result of the recursive call:

```
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```

Input =
$$\langle 2,1,3,2,2,5,4,1 \rangle$$

Partial Output = $(\langle 0,3,8,15\rangle,20)$

Desired Output = $(\langle 0,2,3,6,8,10,15,19\rangle,20)$

This leads to the following code. The algorithm we present works for when n is a power of two.

Let's assume for simplicity that the function being applied has constant work and constant span. We can write out the work and span for the algorithm as a recursive relation as follows.

$$W(n) = W(n/2) + n$$
, and $S(n) = S(n/2) + 1$,

because 1) the contraction step which tabulates the smaller instance of the problem performs linear work in constant span, and 2) the expansion step that constructs the output by tabulating based on the result of the recursive call also performs linear work in constant span.

These recursive relations should look familiar. Indeed, they are the same as those that we ended up with when we analyzed the work and span of our contraction-based implementation of reduce. They yield O(n) work and $O(\log n)$ span.

7.3 Problems

7-1 Ranking sequences

Given a sequence of n natural numbers and and a rank r < n, you want to find the element of the sequence with the given rank. For example if the input is $\langle 1, 3, 2, 4 \rangle$ and you are asked to find the element with rank 0 then the answer is 1 because 1 is the element with rank 0, i.e., the smallest element.

- Using the contraction technique, design an algorithm for finding the element of the sequence with a given rank.
- Analyze the worst-case work and span of your algorithm.
- Analyze the best-case work and span of your algorithm.