

Recitation 2

Parenthesis Matching

2.1 Announcements

- *ParenLab* has been released, and is due **Friday afternoon**. It's worth 100 points. This lab is conceptually difficult – if you haven't started yet, do so tonight!
- *SkylineLab* will be released on Friday.

2.2 Parentheses and Matched Sequences

Suppose you are given a sequence of parentheses. You want to determine if it is *matched*, meaning “properly nested”. Let’s begin by defining this more carefully.

Definition 2.1. A *matched sequence of parentheses* p is defined inductively as

$$p ::= \langle \rangle \mid p p \mid (p)$$

In other words, a matched sequence is one of (a) the empty sequence, (b) the concatenation of two matched sequences, or (c) a pair of parentheses surrounding a matched sequence.

To be consistent with ParenLab, we’ll implement parentheses as a custom datatype given in a structure `Paren`.

```
structure Paren =
struct
  datatype t = L | R
  ...
end
```

Our goal is to implement a function

```
val parenMatch : Paren.t Seq.t → bool
```

where `(parenMatch S)` determines whether or not S is a matched sequence.

Note that you will need to familiarize yourself with the 210 library. Documentation can be found on the course website at <http://www.cs.cmu.edu/~15210/docs/>. In particular, you should look closely at the `SEQUENCE` interface and the `ArraySequence` implementation.

2.3 From Left to Right

Task 2.2. Implement `parenMatch` using the sequence function `iterate`.

2.4 Divide and Conquer

Task 2.3. Implement `parenMatch` with a divide-and-conquer approach. Your implementation should satisfy the following work and span recurrences where n is the length of the input.

$$W(n) = 2 W\left(\frac{n}{2}\right) + O(1)$$
$$S(n) = S\left(\frac{n}{2}\right) + O(1)$$

Also briefly justify that your implementation meets the cost bounds shown. You should assume `Seq = ArraySequence` for cost bounds.

Hint: to solve this problem, you'll only need the sequence function `splitMid` and some basic arithmetic. Check out the documentation of `splitMid` on the website if you are not already familiar. You should also use `Primitives.par` for parallelism – the code `Primitives.par (fn () => e1, fn () => e2)` implements the parallel pair $(e_1 \parallel e_2)$. It is logically equivalent to just writing (e_1, e_2) , except that the two expressions are evaluated in parallel.

2.5 Additional Exercises

Exercise 2.4. As implied by the name, the `ArraySequence` implementation of sequences lays out its elements in an array. Describe how to implement `splitMid` (and in general, `subseq`) in $O(1)$ work and span.

