C'mon K-mers!

A useful operation on strings, specially when analyzing certain types of data, like genomic data, is to find all the substrings of length k in a string. These substrings are called the string's k-mers and we are typically interested in finding out how many times each k-mer appears within the string. For example, consider the following string:

aaabaab

This string has three unique 2-mers:

aaabaab

```
aa
aa
ab
ba
aa
```

More specifically, the 2-mers are aa (appearing three times), ab (appearing two times), and ba (appearing once). Notice how counting the k-mers considers overlapping occurrences of each substring. For example, the first two occurrences of aa above overlap with each other.

We could do a similar analysis for the string's 3-mers:

aaabaab

```
aaa
aab
aba
baa
aab
```

In this case, there are four unique 3-mers: aaa (appearing once), aab (appearing twice), aba (appearing once), and baa (appearing once).

In this problem, you will write a function kmers that, given a potentially large string, a value for k, and some k-mers, will determine how many times each k-mer appears in the string. The function should return a list of integers where the ith integer is the number of times the ith mer in the list of mers occurs in the sequence.

Here are some sample inputs and outputs:

Sequence	K-Mers	Results
aaabaab	ab ba aa	2 1 3

Sequence	K-Mers	Results
aaabaab	aaa aab aaa baa xyz	1 2 1 1 0

Here is the skeleton code for this task:

```
import java.util.List;
import java.util.HashMap;
public class Problem4 {
    /**
     * kmers: Takes in a sequence of letters, a kmer length, and a list of k-mers and
     * returns a dictionary of k-mers and their counts in the sequence.
     * Arguments:
        seq: A string of letters
         k: The length of the k-mers represented as an integer
         k-mers: An array of the k-mers of interest
     * Returns: an array where the ith entry is the number
         of times that the ith k-mer in kmers occurs on the sequence
     **/
   public static int[] kmers(String seq, int k, String[] mers) {
        // Return included to allow the skeleton code to compile
        return null;
    }
}
```

When you take the placement exam, you will be expected to copy the skeleton code into a file and then complete the function. For the practice problems, we have provided a file named Problem4.java that includes the above header for your convenience.