# **Minimum Skew Problem**

**Input**: A DNA string *Genome*.

**Output**: All integer(s) i minimizing  $Skew_i(Genome)$  among all values of i (from 0 to |Genome|).

## **SAMPLE DATASET:**

Input:

 ${\tt TAAAGACTGCCGAGAGGCCAACACGAGTGCTAGAACGAGGGGCGTAAACGCGGGTCCGAT}$ 

Output:

11 24

The sample dataset is not actually run on your code.

# TEST DATASET 1: Input: ACCG Output:

3

This dataset checks if your code's indexing is off. Specifically, it verifies that your code is not returning an index 1 too high (i.e. 4) or 1 too low (i.e. 2).

# **TEST DATASET 2:**

Input:

Output:

4

This dataset checks to see if your code is missing the last symbol of *Genome*.

## **TEST DATASET 3:**

Input:

CCGGGT

Output:

2

This dataset makes sure you're not accidentally finding the maximum skew instead of the minimum skew.

## **TEST DATASET 4:**

Input:

CCGGCCGG

Output:

2 6

First, this dataset checks if you are only finding 1 index (and not multiple indices). Then, it checks if you are using a delimiter to separate your indices (ideally a space character).