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Started on	Tuesday, 4 June 2024, 10:22 PM
State	Finished
Completed on	Wednesday, 5 June 2024, 12:52 PM
Time taken	14 hours 29 mins
Marks	5.00/5.00
Grade	100.00 out of 100.00

Question **1**

Correct

Mark 1.00 out of 1.00

Write a program to eliminate the common elements in the given 2 arrays and print only the non-repeating elements and the total number of such non-repeating elements.

Input Format:

The first line contains space-separated values, denoting the size of the two arrays in integer format respectively.

The next two lines contain the space-separated integer arrays to be compared.

[Sample](#) Input:

```
5 4
1 2 8 6 5
2 6 8 10
```

[Sample](#) Output:

```
1 5 10
3
```

[Sample](#) Input:

```
5 5
1 2 3 4 5
1 2 3 4 5
```

[Sample](#) Output:

```
NO SUCH ELEMENTS
```

**For example:**

Input	Result
5 4 1 2 8 6 5 2 6 8 10	1 5 10 3
5 5 1 2 3 4 5 1 2 3 4 5	NO SUCH ELEMENTS

**Answer:** (penalty regime: 0 %)

```
1 def find_non_repeating_elements():
2     n,m=map(int, input().split())
3     arr1=list(map(int, input().split()))
4     arr2=list(map(int, input().split()))
5     set1=set(arr1)
6     set2=set(arr2)
```

```

/      non_repeating_elements = set1.symmetric_difference(set2)
8 ▾    if len(non_repeating_elements) == 0:
9      print("NO SUCH ELEMENTS")
10 ▾   else:
11     print(' '.join(map(str, non_repeating_elements)))
12     print(len(non_repeating_elements))
13 find_non_repeating_elements()
```

	Input	Expected	Got	
✓	5 4 1 2 8 6 5 2 6 8 10	1 5 10 3	1 5 10 3	✓
✓	3 3 10 10 10 10 11 12	11 12 2	11 12 2	✓
✓	5 5 1 2 3 4 5 1 2 3 4 5	NO SUCH ELEMENTS	NO SUCH ELEMENTS	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question **2**

Correct

Mark 1.00 out of 1.00

The **DNA sequence** is composed of a series of nucleotides abbreviated as 'A', 'C', 'G', and 'T'.

- For example, "ACGAATTCG" is a **DNA sequence**.

When studying **DNA**, it is useful to identify repeated sequences within the DNA.

Given a string `s` that represents a **DNA sequence**, return all the **10-letter-long** sequences (substrings) that occur more than once in a DNA molecule. You may return the answer in **any order**.

**Example 1:**

Input: `s = "AAAAACCCCCAAAAACCCCCAAAAGGGTTT"`

Output: `["AAAAACCCCC", "CCCCAAAAA"]`

**Example 2:**

Input: `s = "AAAAAAAAAAAA"`

Output: `["AAAAAAAAA"]`

**For example:**

Input	Result
AAAAACCCCCAAAAACCCCCAAAAGGGTTT	AAAAACCCCC CCCCAAAAA

**Answer:** (penalty regime: 0 %)

```

1 s=input()
2 substring_counts={}
3 for i in range(len(s)-9):
4     substring=s[i:i+10]
5     substring_counts[substring]=substring_counts.get(substring,0)+1
6 repeated_substrings=[substring for substring, count in substring_counts.items() if count>1]
7 for substring in repeated_substrings:
8     print(substring)

```

	Input	Expected	Got	
✓	AAAAACCCCAAAAACCCCAAAAAGGGTTT	AAAAACCCCAAAAACCCCAAAA	AAAAACCCCAAAAACCCCAAAA	✓
✓	AAAAAAAAAAAA	AAAAAAAAAAAA	AAAAAAAAAAAA	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question **3**

Correct

Mark 1.00 out of 1.00

Given an array of integers `nums` containing  $n + 1$  integers where each integer is in the range  $[1, n]$  inclusive. There is only **one repeated number** in `nums`, return *this repeated number*. Solve the problem using [set](#).

**Example 1:**Input: `nums = [1,3,4,2,2]`

Output: 2

**Example 2:**Input: `nums = [3,1,3,4,2]`

Output: 3

**For example:**

Input	Result
1 3 4 4 2	4

**Answer:** (penalty regime: 0 %)

```
1 x=input()
2 y=x.split()
3 z=list(y)
4 a=[]
5 b=[]
6 for element in z:
7     if element in a:
8         b.append(element)
9     else:
10        a.append(element)
11 c=' '.join(map(str,b))
12 print(c)
```

	Input	Expected	Got	
✓	1 3 4 4 2	4	4	✓
✓	1 2 2 3 4 5 6 7	2	2	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question **4**  
Correct  
Mark 1.00 out of 1.00

Coders here is a simple task for you, Given string str. Your task is to check whether it is a binary string or not by using python [set](#).

Examples:

Input: str = "01010101010"

Output: Yes

Input: str = "REC101"

Output: No

For example:

Input	Result
01010101010	Yes
010101 10101	No

Answer: (penalty regime: 0 %)

```
1 a=input()
2 try:
3     int(a)
4     print("Yes")
5 except:
6     print("No")
```

	Input	Expected	Got	
✓	01010101010	Yes	Yes	✓



	Input	Expected	Got	
✓	REC123	No	No	✓
✓	010101 10101	No	No	✓

Passed all tests! ✓

Correct

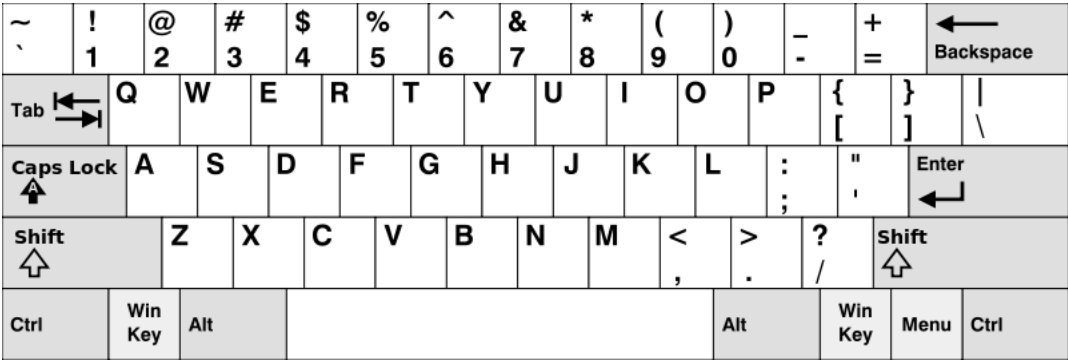
Marks for this submission: 1.00/1.00.

Question 5  
Correct  
Mark 1.00 out of 1.00

Given an array of [strings](#) *words*, return the words that can be typed using letters of the alphabet on only one row of American keyboard like the image below.

In the **American keyboard**:

- the first row consists of the characters "qwertyuiop",
- the second row consists of the characters "asdfghjkl", and
- the third row consists of the characters "zxcvbnm".



Example 1:

Input: words = ["Hello", "Alaska", "Dad", "Peace"]  
Output: ["Alaska", "Dad"]

Example 2:

Input: words = ["omk"]  
Output: []

Example 3:

Input: words = ["adsdf", "sfd"]  
Output: ["adsdf", "sfd"]

For example:

Input	Result
4	Alaska
Hello	Dad
Alaska	
Dad	
Peace	

Input	Result
2	adsfd
adsfd	afd
afd	

**Answer:** (penalty regime: 0 %)

```

1 def findwords(words):
2     row1 = set('qwertyuiop')
3     row2 = set('asdfghjkl')
4     row3 = set('zxcvbnm')
5     result = []
6     for word in words:
7         w = set(word.lower())
8         if w.issubset(row1) or w.issubset(row2) or w.issubset(row3):
9             result.append(word)
10    if len(result) == 0:
11        print("No words")
12    else:
13        for i in result:
14            print(i)
15 a=int(input())
16 arr = [input() for i in range(a)]
17 findwords(arr)

```

	Input	Expected	Got	
✓	4 Hello Alaska Dad Peace	Alaska Dad	Alaska Dad	✓
✓	1 omk	No words	No words	✓
✓	2 adsfd afd afd	adsfd afd	adsfd afd	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

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