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Started on	Wednesday, 5 June 2024, 1:02 PM
State	Finished
Completed on	Wednesday, 5 June 2024, 2:20 PM
Time taken	1 hour 18 mins
Marks	5.00/5.00
Grade	100.00 out of 100.00

Question 1

Correct

Mark 1.00 out of 1.00

There is a malfunctioning keyboard where some letter keys do not work. All other keys on the keyboard work properly.

Given a string text of words separated by a single space (no leading or trailing spaces) and a string brokenLetters of all distinct letter keys that are broken, return the number of words in text you can fully type using this keyboard.

Example 1:

Input: text = "hello world", brokenLetters = "ad"

Output:

1

Explanation: We cannot type "world" because the 'd' key is broken.

For example:

Input	Result
hello world ad	1
Faculty Upskilling in Python Programming ak	2

Answer: (penalty regime: 0 %)

```

1 a=list(input().split())
2 b=list(input())
3 c=0
4 for i in a:
5     d=0
6     for j in b:
7         if j in i.lower():
8             d+=1
9     if d== 0:
10        c+=1
11 print(c)

```

	Input	Expected	Got	
✓	hello world ad	1	1	✓
✓	Welcome to REC e	1	1	✓
✓	Faculty Upskilling in Python Programming ak	2	2	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question 2

Correct

Mark 1.00 out of 1.00

Given a tuple and a positive integer k, the task is to find the count of distinct pairs in the tuple whose sum is equal to **K**.

Examples:**Input:** t = (5, 6, 5, 7, 7, 8), K = 13**Output:** 2**Explanation:**

Pairs with sum K(= 13) are {(5, 8), (6, 7), (6, 7)}.

Therefore, distinct pairs with sum K(= 13) are { (5, 8), (6, 7) }.

Therefore, the required output is 2.

For example:

Input	Result
1,2,1,2,5 3	1
1,2 0	0

Answer: (penalty regime: 0 %)

```

1 x=input()
2 y=int(input())
3 a=x.split(',')
4 t=tuple(int(num) for num in a)
5 ans=set()
6 for i in range(len(t)):
7     for j in range(i+1, len(t)):
8         if t[i]+t[j]==y:
9             pair=(min(t[i],t[j]), max(t[i],t[j]))
10            if pair not in ans:
11                ans.add((t[i],t[j]))
12 print(len(ans))
13

```

	Input	Expected	Got	
✓	5,6,5,7,7,8 13	2	2	✓
✓	1,2,1,2,5 3	1	1	✓
✓	1,2 0	0	0	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question 3

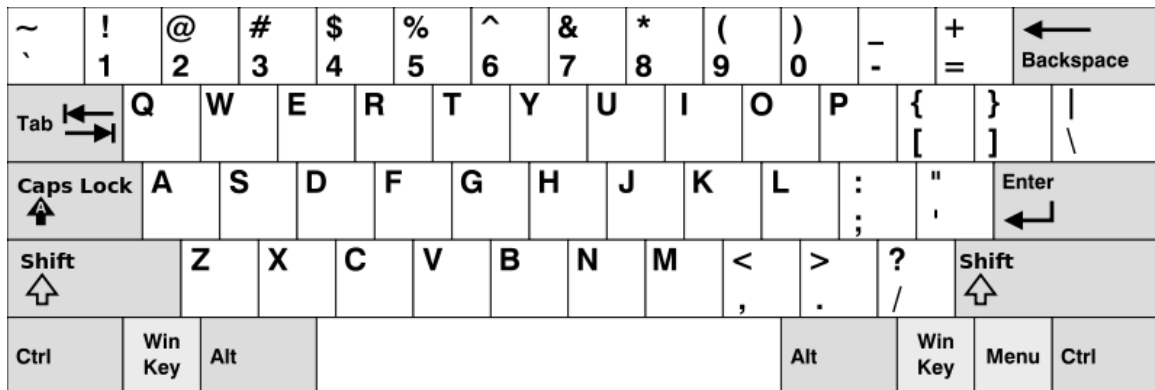
Correct

Mark 1.00 out of 1.00

Given an array of [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 Hello Alaska Dad Peace	Alaska Dad
2 adsdf afd afd	adsdf 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	adsfd afd	adsfd afd	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question 4

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 = "AAAAACCCCCAAAAACCCCCAAAAAGGGTTT"`

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

Example 2:

Input: `s = "AAAAAAAAAAAA"`

Output: `["AAAAAAAAA"]`

For example:

Input	Result
AAAAACCCCCAAAAACCCCCAAAAAGGGTTT	AAAAACCCC 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	
✓	AAAAACCCCCAAAAACCCCCAAAAAGGGTTT	AAAAACCCC CCCCAAAAA	AAAAACCCC CCCCAAAAA	✓
✓	AAAAAAAAAAAAA	AAAAAAAAA	AAAAAAAAA	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question **5**

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	✓
✓	REC123	No	No	✓
✓	010101 10101	No	No	✓

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

◀ Week7_MCQ

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