<u>Dashboard</u> / <u>My courses</u> / <u>PSPP/PUP</u> / <u>Experiments based on Tuples, Sets and its operations</u> / <u>Week7 Coding</u>

Started on	Friday, 24 May 2024, 9:28 AM
State	Finished
Completed on	Sunday, 26 May 2024, 9:28 AM
Time taken	2 days
Marks	4.00/5.00
Grade	80.00 out of 100.00

Question 1
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 <u>set</u>.

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 
    def find_duplicate(nums):
 2
        seen = set()
 3 ▼
        for num in nums:
            if num in seen:
 4 ▼
 5
                 return num
            seen.add(num)
 6
 7
   nums = list(map(int, input().split()))
   print(find_duplicate(nums))
 8
 9
10
```

	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 **2**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 <u>set</u>.

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 ▼ def is_binary_string(s):
 2
        binary_set = {'0', '1'}
 3
        return set(s).issubset(binary_set)
 4
 5 ▼ def main():
        s = input().strip()
 6
        if is_binary_string(s):
 7 ▼
 8
            print("Yes")
 9 🔻
        else:
10
            print("No")
11
12 v if __name__ == "__main__":
13
        main()
```

	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.

Question **3**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 %)

```
t = tuple(map(int, input().split(',')))
   K = int(input())
 2
 3
 4
   seen = {}
 5
   distinct_pairs = set()
 6
 7 v for num in t:
        complement = K - num
 8
        if complement in seen and seen[complement] > 0:
 9 🔻
            distinct_pairs.add((min(num, complement), max(num, complement)))
10
11
            seen[complement] -= 1
        else:
12 🔻
13
            seen[num] = seen.get(num, 0) + 1
14
15 print(len(distinct_pairs))
```

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

Passed all tests! <

Correct

Marks for this submission: 1.00/1.00.

Question **4**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

12865

26810

Sample Output:

1 5 10

3

Sample Input:

5 5

12345

12345

Sample Output:

NO SUCH ELEMENTS

For example:

Input				R	es	ult	
5	4				1	5	10
1	2	8	6	5	3		
2	6	8	16	9			

Answer: (penalty regime: 0 %)

	Input	Expected	Got	
~	5 4	1 5 10	1 5 10	~
	1 2 8 6 5	3	3	
	2 6 8 10			
~	3 3	11 12	11 12	~
	10 10 10	2	2	
	10 11 12			

Passed all tests! ✓

Correct

Marks for this submission: 1.00/1.00.

Question **5**Incorrect
Mark 0.00 out of

1.00

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

• For example, "ACGAATTCCG" 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 = "AAAAACCCCCAAAAAACCCCCCAAAAAGGGTTT"
Output: ["AAAAACCCCC","CCCCCAAAAA"]
```

Example 2:

```
Input: s = "AAAAAAAAAAA"
Output: ["AAAAAAAAAA"]
```

For example:

Input	Result
AAAAACCCCCAAAAACCCCCCAAAAAGGGTTT	AAAAACCCCC CCCCAAAAA

Answer: (penalty regime: 0 %)

```
s = input()
   A = set()
 2
 3
   B = set()
 4 v for i in range(len(s) - 9):
        C = s[i:i + 10]
 5
 6 ▼
        if C in A:
 7
            B.add(C)
 8 🔻
        else:
 9
            A.add(C)
10 v for seq in B:
        print(seq)
11
```

	Input	Expected	Got	
×	AAAAACCCCCAAAAACCCCCCAAAAAGGGTTT		CCCCCAAAAA AAAAACCCCC	×
~	АААААААААА	АААААААА	АААААААА	~

Your code must pass all tests to earn any marks. Try again.

Show differences

Incorrect

Marks for this submission: 0.00/1.00.

■ Week7_MCQ

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Dictionary ►