

Examples:

Input: str = "01010101010"

Output: Yes

Input: str = "REC101"

Output: No

| Input        | Result |
|--------------|--------|
| 01010101010  | Yes    |
| 010101 10101 | No     |

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## **Binary String**

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.

```
str1=input()
unique_chars1 = set(str1)
if unique_chars1 <= {'0', '1'}:
    print("Yes")
else:
    print("No")</pre>
```

### **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.

| Input     | Result |
|-----------|--------|
| 1,2,1,2,5 | 1      |
| 1,2       | 0      |

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# **Check Pair**

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.

```
t = input()
st = t.split(',')
t1 = tuple(int(item.strip()) for item in st)
K = int(input())
seen = set()
pairs = set()

for number in t1:
    complement = K - number
    if complement in seen:
        pairs.add(tuple(sorted((number, complement))))
    seen.add(number)
print( len(pairs))
```

Input: s = "AAAAACCCCCCAAAAACCCCCCAAAAAGGGTTT"

Output: ["AAAAACCCCC","CCCCCAAAAA"]

Example 2:

Input: s = "AAAAAAAAAAA"

Output: ["AAAAAAAAAA"]

| Input                            | Result                  |
|----------------------------------|-------------------------|
| AAAAACCCCCAAAAACCCCCCAAAAAGGGTTT | AAAAACCCCC<br>CCCCAAAAA |

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## **DNA Sequence**

Name:

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**.

```
def findRepeatedDnaSequences(s):
  if len(s) < 10:
     return []
  sequences = {}
  result = []
  for i in range(len(s) - 9):
     substring = s[i:i+10]
     if substring in sequences:
       sequences[substring] += 1
     else:
       sequences[substring] = 1
  for sequence, count in sequences.items():
     if count > 1:
       result.append(sequence)
  for i in result:
     print(i)
s1=input()
findRepeatedDnaSequences(s1)
```

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

Output: 2

#### Example 2:

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

Output: 3

| Input     | Result |
|-----------|--------|
| 1 3 4 4 2 | 4      |

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## Print repeated no

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  $\underline{set}$ .

```
def find_duplicate(nums):
    seen = set()
    for num in nums:
        if num in seen:
            return num
        seen.add(num)

# Input
I = input()
    nums = list(map(int, I.split()))
# Output
print(find_duplicate(nums))
```

Sample Input:

5 4

12865

26810

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

| Input                        | Result      |
|------------------------------|-------------|
| 5 4<br>1 2 8 6 5<br>2 6 8 10 | 1 5 10<br>3 |

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# Remove repeated

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.

```
a=[int(i) for i in input().split()]
n1=[int(i) for i in input().split()]
n2=[int(i) for i in input().split()]
com=set(n1)&set(n2)
uni=set(n1)|set(n2)
tot=sorted(uni-com)
for i in tot:
    print(i,end=' ')
print('\n',end=")
print(len(tot))
```

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

#### Output:

1

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

| Input             | Result |
|-------------------|--------|
| hello world<br>ad | 1      |

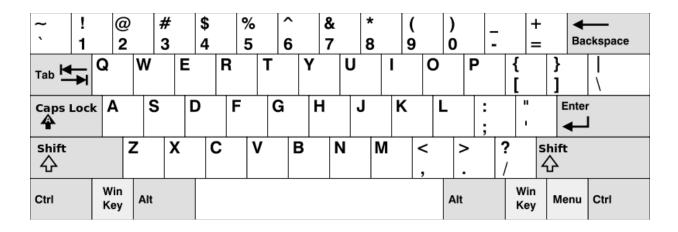
| Ex. No.     | :  | 8.6 | Date: |
|-------------|----|-----|-------|
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## **Malfunctioning Keyboard**

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.

```
a=[i for i in input().split()]
k=list(input())
s=set()
for i in a:
    n=[j for j in i]
    m=[z for z in k if z in n]
    s.update(m)
print(len(s))
```



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

Output: ["Alaska","Dad"]

Example 2:

Input: words = ["omk"]

Output: [] Example 3:

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

| Input                                | Result        |
|--------------------------------------|---------------|
| 4<br>Hello<br>Alaska<br>Dad<br>Peace | Alaska<br>Dad |

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# American keyboard

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

```
n=int(input())
f=0
a=[input() for i in range(n)]
I1=['qwertyuiop','asdfghjkl','zxcvbnm']
I=[[j for j in i] for i in I1]
for i in a:
   n=[j for j in i.lower()]
   if set(n)|set(I[0]) = set(I[0]):
     f=1
      print(i)
     continue
   elif set(n)|set(I[1]) = set(I[1]):
     f=1
     print(i)
     continue
   elif set(n)|set(I[2]) = set(I[2]):
     f=1
      print(i)
     continue
if not f:
   print('No words')
```