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Class- MSc CS – I

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Subject – Bioinformatics

**Topic – Pairwise Sequence
Alignment**

Practical No: 1

Aim: Write a Python/Java code to perform pairwise alignment. Take 2 sequences from user and calculate the score.

Code:

```
se1=input("Enter the first sequence::")
se2=input("Enter the second sequence::")
seq1=list(se1)
seq2=list(se2)
score=[]

def Pairwise_alignment(a,b):
    gap(a,b)
    print(a)
    print(b)
    value=0
    length=len(a)
    for i in range(0,length):
        if(a[i]==b[i]):
            score.append('1')
            value=value+1
        else:
            score.append('0')
    print(score)
    print(value)

def gap(a,b):
    if(len(a)==len(b)):
        print()
    else:
        k=int(input("enter the position to insert::"))
        if (len(a)<len(b)):
            a.insert(k,'-')
        else:
            b.insert(k,'-')
    return(a,b)
Pairwise_alignment(seq1,seq2)
```

Output:

PROBLEMS OUTPUT DEBUG CONSOLE TERMINAL

Try the new cross-platform PowerShell <https://aka.ms/pscore6>

PS E:\Python codes> python -u "e:\Python codes\pairwise.py"

Enter the first sequence::abcvfc

Enter the second sequence::abbcvf

['a', 'b', 'c', 'v', 'f', 'c']

['a', 'b', 'b', 'c', 'v', 'f']

['1', '1', '0', '0', '0', '0']

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