# Jutge.org

The Virtual Learning Environment for Computer Programming

# **Protein Domain Overlap**

Z99185\_en

Proteins contain different *domains* (structural or functional units responsible for a particular function).

We need a program that, given a list of domains in a protein, and their positions in it, determines domain overlapping regions in the sequence.

#### Input

The input is a list of protein domains found in a DNA sequence. For each domain, the id of the protein, the name of the domain, and its position span in the protein are given. The input consists of an integer N (the number of protein domain records), followed by N lines, each consisting of two strings, and two integers:

protein\_id domain\_name start\_position end\_position
where:

- protein\_id (string): The protein's identifier.
- domain\_name (string): The name of the domain.
- start\_position, end\_position (integers): The span of the domain within the protein sequence.

#### Output

List proteins in alphabetical order. For each protein, list its domains in order of starting position. If any domain overlaps with the previous one, mark it with "OVERLAP". Print a summary list of the overlapping domains at the end of each protein. If no overlaps exist, print "No overlaps".

Follow the format of the examples.

#### Sample input 1

```
6
P12X43 Kinase 5 50
T5678A Phosphatase 10 40
T5678A Transmembrane 50 90
P12X43 SH3 30 70
T5678A Immunoglobulin 35 60
P12X43 Pleckstrin 80 100
```

#### Sample input 2

```
8
P12X31 RRM 5 20
R3012Y Collagen 35 45
H2F127 FN3 35 50
R3012Y EGF 5 15
```

#### Sample output 1

```
P12X43:

Kinase (5-50)

SH3 (30-70) OVERLAP

Pleckstrin (80-100)

Overlapping domains in protein P12X43: Kinase-SH3
T5678A:

Phosphatase (10-40)

Immunoglobulin (35-60) OVERLAP

Transmembrane (50-90) OVERLAP

Overlapping domains in protein T5678A: Phosphatase-In
```

```
R3012Y Cadherin 20 30
P12X31 Catalase 25 40
P12X31 Kinase 45 60
H2F127 RRM 10 30
```

#### Sample output 2

```
H2F127:

RRM (10-30)

FN3 (35-50)

No overlaps
P12X31:

RRM (5-20)
```

## Sample input 3

```
12
P448X1 SH3 85 110
P59S87 Catalase 5 25
P59S87 HTH 10 35
P59S87 Immunoglobulin 75 90
P448X1 Transmembrane 10 40
P448X1 Kinase 30 60
P59S87 7TM 40 50
P59S87 PDZ 45 70
M32101 Pleckstrin 5 15
P448X1 SH2 55 90
P59S87 WD40 5 25
M32101 Porin 20 30
```

#### **Problem information**

Author: Lluís Padró

Generation: 2025-03-24 17:51:18

© *Jutge.org*, 2006–2025. https://jutge.org

```
Catalase (25-40)
Kinase (45-60)
No overlaps
R3012Y:
EGF (5-15)
Cadherin (20-30)
Collagen (35-45)
No overlaps
```

### Sample output 3

```
M32101:
 Pleckstrin (5-15)
 Porin (20-30)
 No overlaps
P448X1:
 Transmembrane (10-40)
 Kinase (30-60) OVERLAP
 SH2 (55-90) OVERLAP
 SH3 (85-110) OVERLAP
 Overlapping domains in protein P448X1: Transmembrane
P59S87:
 Catalase (5-25)
 WD40 (5-25) OVERLAP
 HTH (10-35) OVERLAP
 7TM (40-50)
 PDZ (45-70) OVERLAP
 Immunoglobulin (75-90)
  Overlapping domains in protein P59S87: Catalase-WD40
```