Introduction to Data Science (IDS) course

## **Sequence Mining - Instruction**Lecture 13

[DS-L13-]





### **Sequence Mining**

In the Sequence Mining lecture you have seen AprioriAll, an algorithm for mining sequences.

It can be considered a generalized version of Apriori.



# Can you tell the differences between Apriori and AprioriAll?

- What are the goals of the two algorithms?
- What is the definition of pattern of each algorithm?
- Where do the two algorithms look for patterns?



#### **Answers!**

Apriori looks at items grouped together.

 AprioriAll looks at items with a specific partial ordering.



#### **Answers!**

- Apriori looks at items grouped together.
- Patterns are sets of items.

- AprioriAll looks at items with a specific partial ordering.
- Patterns are sequences of sets of items.



#### **Answers!**

- Apriori looks at items grouped together.
- Patterns are sets of items.
- Apriori finds patterns within sets.

- AprioriAll looks at items with a specific partial ordering.
- Patterns are sequences of sets of items.
- AprioriAll finds patterns between sets.



### Support

A central concept in pattern mining is *support* (and *support count*): the frequency of appearance (relative or absolute) of a certain pattern within the database.



### Support

```
In this database, find the
support count of:
(bc)(de)
b(de)
(bc)de
(ac)(bc)
(bc)(ac)
```

```
D = [
<a(bc)d(eb)>,
<(ac)(bc)de>,
<(ac)b(cd)>,
<ab(bc)(cde)>,
<(bc)(bd)(bde)>,
<(abc)(ac)(bc)de>,
<a(bd)c(de)>
```

```
In this database, find the
support of:
(bc)(de): 2
b(de)
(bc)de
(ac)(bc)
(bc)(ac)
```

```
D = [
<a(bc)d(eb)>,
<(ac)(bc)de>,
<(ac)b(cd)>,
<ab(bc)(cde)>,
<(bc)(bd)(bde)>,
<(abc)(ac)(bc)de>,
<a(bd)c(de)>
```

```
In this database, find the support of:
```

(bc)(de): 2

b(de): 3

(bc)de

(ac)(bc)

(bc)(ac)

```
D = [
<a(bc)d(eb)>,
<(ac)(bc)de>,
<(ac)b(cd)>,
<ab(bc)(cde)>,
<(bc)(bd)(bde)>,
<(abc)(ac)(bc)de>,
<a(bd)c(de)>
```

```
In this database, find the support of:
```

(bc)(de): 2

b(de): 3

(bc)de: 4

(ac)(bc)

(bc)(ac)

```
D = [
<a(bc)d(eb)>,
<(ac)(bc)de>,
<(ac)b(cd)>,
<ab(bc)(cde)>,
<(bc)(bd)(bde)>,
<(abc)(ac)(bc)de>,
<a(bd)c(de)>
```

```
In this database, find the support of:
```

(bc)(de): 2

b(de): 3

(bc)de: 4

(ac)(bc): 2

(bc)(ac)

```
D = [
<a(bc)d(eb)>,
<(ac)(bc)de>,
<(ac)b(cd)>,
<ab(bc)(cde)>,
<(bc)(bd)(bde)>,
<(abc)(ac)(bc)de>,
<a(bd)c(de)>
```

```
In this database, find the support of:
```

(bc)(de): 2

b(de): 3

(bc)de: 4

(ac)(bc): 2

(bc)(ac): 1

```
D = [
<a(bc)d(eb)>,
<(ac)(bc)de>,
<(ac)b(cd)>,
<ab(bc)(cde)>,
<(bc)(bd)(bde)>,
<(abc)(ac)(bc)de>,
<a(bd)c(de)>
```