

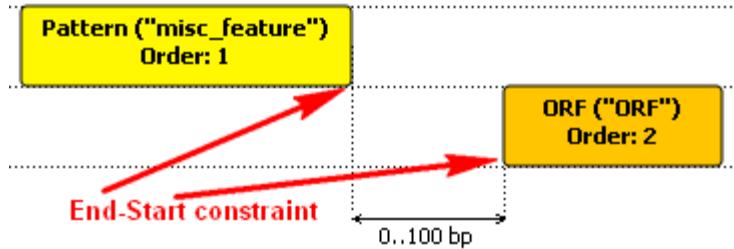
# Constraint Elements

To add a constraint element add the required algorithm elements to the scene and drag&drop the required constraint element.

When a *constraint element* is added, two *algorithm elements* are selected.

There are four types of constraints that you can impose on the positional relationship of the results obtained from the algorithms calculations: *End-Start*, *Start-End*, *End-End* and *Start-Start*.

On the image below you can see a schema with *Pattern* and *ORF* algorithm elements and an *End-Start* constraint:



This means the following:

1. The algorithm elements specify to analyze the sequence with Pattern and ORF algorithms. The results of these analyses are the sets of annotations.
2. The condition says that the distance between a “Pattern annotation end” and an “ORF annotation start” should be within the specified bounds, i.e.:

Let:

```
pattern_annot_end := the last nucleotide of an annotation obtained from the Pattern algorithm calculations  
orf_annot_start := the first nucleotide of an annotation obtained from the ORF algorithm calculations
```

The constraint is:

```
0bp <= Distance(pattern_annot_end, orf_annot_start) <= 100bp
```

Find the details on each constraint element below.

- End-Start Constraint Element
- Start-End Constraint Element
- End-End Constraint Element
- Start-Start Constraint Element