

Selecting Annotations

To select one annotation click on it. To select several annotations hold *Ctrl* key while clicking on the annotations.

To invert the selection use the *Invert annotation selection* item in the *Annotations editor* context menu.

| Name | Type | Value |
|--|-------|------------------|
| Auto-annotations [sars.gb NC_004718] | | |
| NC_004718 features [sars.gb] | | |
| 3'UTR (0, 1) | | |
| 3'UTR | 3'UTR | 29389..29751 |
| 5'UTR (0, 1) | | |
| 5'UTR | 5'UTR | 1..264 |
| CDS (0, 14) | | |
| CDS | CDS | 1265..13398,1413 |
| CDS | CDS | 25259 |
| CDS | CDS | 26092 |
| CDS | CDS | 26153 |
| CDS | CDS | 26347 |
| CDS | CDS | 27063 |
| CDS | CDS | 27265 |
| CDS | CDS | 27641 |
| CDS | CDS | 27772 |
| CDS | CDS | 27898 |
| CDS | CDS | 28118 |
| CDS | CDS | 29388 |
| CDS | CDS | 28426 |
| comment (0, 1) | | |

Find qualifier...

Invert annotation selection

Go to position... Ctrl+G

Select sequence region... Ctrl+A

New annotation... Ctrl+N

Copy/Paste

Select

Add

Analyze

Align

Cloning

Fetch sequences from remote database

After inverting, all annotations are highlighted and selected except for the first two CDS.

| Name | Type | Value |
|--|-------|-------------------------------|
| Auto-annotations [sars.gb NC_004718] | | |
| NC_004718 features [sars.gb] | | |
| 3'UTR (0, 1) | | |
| 3'UTR | 3'UTR | 29389..29751 |
| 5'UTR (0, 1) | | |
| 5'UTR | 5'UTR | 1..264 |
| CDS (0, 14) | | |
| CDS | CDS | join(265..13398,13398..21485) |
| CDS | CDS | 265..13413 |
| CDS | CDS | 21492..25259 |
| CDS | CDS | 25268..26092 |
| CDS | CDS | 25689..26153 |
| CDS | CDS | 26117..26347 |
| CDS | CDS | 26398..27063 |
| CDS | CDS | 27074..27265 |
| CDS | CDS | 27273..27641 |
| CDS | CDS | 27638..27772 |
| CDS | CDS | 27779..27898 |
| CDS | CDS | 27864..28118 |
| CDS | CDS | 28120..29388 |
| CDS | CDS | 28130..28426 |
| comment (0, 1) | | |

Double-clicking on an annotation also selects the corresponding sequence region.