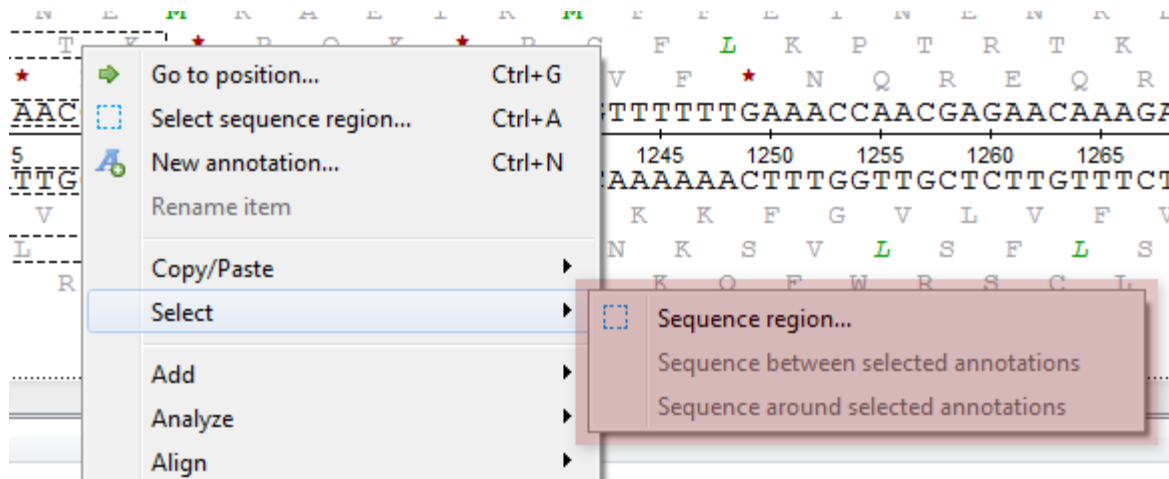
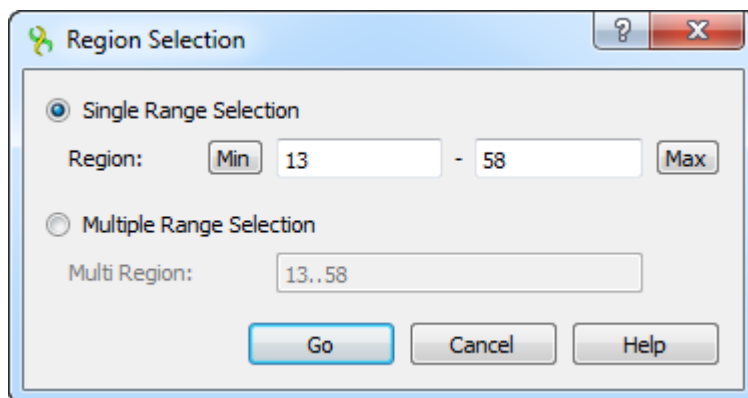


Selecting Sequence

You can use different items from the *Select* submenu of the context menu to select a sequence.



Selecting the *Sequence region* context menu item opens the *Select range* dialog:



Here you can specify the sequence range you would like to select.

You can open the same dialog using the *Select sequence region* button on a sequence toolbar or using the Ctrl-A key sequence.

To use the *Sequence between selected annotations* item, select two annotations in the *Annotations editor* (holding the Ctrl key at the same time):

The screenshot displays a bioinformatics software interface. At the top, a sequence alignment is shown with amino acid residues (N, E, R, P, H, P, V, A, S, L, K, R, H, F, A) and their corresponding DNA codons (M, K, D, P, T, R, R, W, Q, A, S, L, S, N, A, T, L, Q). Below the alignment, a table with 'Name' and 'Value' columns is visible. A context menu is open over the 'misc_feature' entry in the table. The menu includes options like 'Go to position...', 'Select sequence region...', 'New annotation...', 'Copy', 'Select', 'Add', 'Analyze', 'Align', 'Export', 'Edit sequence', 'Remove', 'Rulers...', and 'Annotations highlighting...'. A sub-menu is open for the 'Select' option, showing 'Sequence region...', 'Sequence between selected annotations', and 'Sequence around selected annotations'.

Name	Value
NC_001363 features [murine_copy2.gb]	
CDS (0, 4)	
misc_feature (0, 2)	
misc_feature	2, 500
note	Repeat
misc_feature	33
note	Repeat
source (0, 1)	

And select the *Select Sequence between selected annotations* item in the context menu.

The *Sequence around selected annotations* item selects the selected annotations and the sequences between these annotations.

