

Converting Alignment

To convert an alignment from

- DNA to RNA
- RNA to DNA
- Raw to DNA or Amino

use the *Advanced* context menu or use the General tab of Options Panel:

The image shows a sequence alignment viewer interface. On the left, a DNA alignment is displayed with columns numbered 214 to 227. The alignment consists of 15 rows of nucleotides (A, T, C, G) with some gaps. The background of the alignment is color-coded by column: orange for G, green for T, red for C, and blue for A. Above the alignment is a bar chart showing the frequency of each nucleotide across the columns.

On the right, the 'General' tab of the Options Panel is visible. It contains the following sections:

- Reference sequence:** A dropdown menu with the text 'Select and add' and a close button (X).
- Alignment info:** Displays 'Alphabet: Raw', 'Length: 269', and 'Sequences: 52'. Below this are two buttons: 'DNA' (highlighted in red) and 'Amino' (highlighted in red).
- Consensus mode:** Includes a 'Consensus type:' dropdown menu set to 'ClustalW' and a 'Threshold:' slider set to 0%.

A vertical toolbar on the far right contains icons for various functions, including a gear icon, a sequence logo icon, a tree icon, and a bar chart icon.