

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 displays a sequence alignment viewer with a bar chart at the top and a sequence alignment below. The alignment consists of 12 rows of DNA sequences, with columns labeled 214, 216, 218, 220, 222, 224, and 227. The sequences are color-coded: G (orange), T (green), C (red), A (blue), and T (green). The alignment is as follows:

Row	214	216	218	220	222	224	227
1	G	T	C	A	A	T	C
2	A	T	C	A	A	T	G
3	A	T	C	A	A	G	C
4	A	T	C	A	A	T	C
5	A	T	C	A	A	T	C
6	A	T	C	A	A	T	C
7	G	T	C	A	A	T	C
8	G	T	C	A	A	T	C
9	A	T	C	A	A	T	G
10	G	T	C	A	A	T	C
11	A	T	C	A	A	T	C
12	A	T	C	A	A	G	C
13	A	T	C	A	A	T	G
14	A	T	C	A	A	T	G

Overlaid on the right is the 'General' tab of the Options Panel. It contains the following settings:

- Reference sequence:** Select and add
- Alignment info:** Alphabet: Raw; Length: 269; Sequences: 52. Buttons for 'DNA' and 'Amino' are visible.
- Consensus mode:** Consensus type: ClustalW; Threshold: 0%.