

# Navigation in Sanger Reads Alignment

You may jump to the previous/next ambiguous characters in the read alignment and edit the characters.

After you finish the editing, use the navigation by previous/next variations to find differences between the reference sequence and the consensus.

Use the main *Actions*→ *Navigation* menu, context menu, or the following toolbar buttons:



To go to selected read use:

- *Actions NavigationGo to selected read* the main menu item
- *NavigationGo to selected read* context menu item
- *Enter*
- *Space*
- *DoubleClick*