VBRC Base By Base: Switching between Mouse Modes

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Abstract

Viral Bioinformatic Resource Centre

- Provide databases of viral genomic information.
 - Please check the **Organisms** menu to see which viruses we support: we're now focusing on large DNA viruses
 - The VOCs (Virus Orthologous Clusters) database is at the heart of our system.
 - The database links directly to integrated tools for comparative analyses.
 - VOCs sorts genes into ortholog clusters (e.g. RNA polymerase) to simplify data retrieval.
- Provide easy access to the genes, gene families, and genomes of the different virus families.
 - via a unique series of powerful Java tools that support multiple computer platforms (see VBRC Tools menu).
 - design and build software to tackle specific bioinformatics/virology problems, often in collaboration with virologists.
- Rally the research community to provide expert curation of these viral genomes by:
 - Adding value to GenBank sequences through enhancing and updating genome annotations
 - Linking to research reviews/papers for the research community.
- Collaborate with researchers to help on specific bioinformatics problems, e.g.
 - Custom searches of the databases
 - Building new features into our tools
 - Help with genome annotation

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Protocol

Step 1.

BBB contains three different mouse modes: Edit, Block Glue and Select.

All three controlled by the Toolbar (the fourth, fifth, and sixth buttons from the left.)

It is important to keep these three modes straight as using the wrong one can be deleterious to your alignment!

Step 2.

The **Edit** mode allows you to <u>create gaps of any length</u> (in the selected sequences) by clicking and dragging. **DO NOT MISTAKE THIS FOR SELECT MODE.**

Step 3.

The **Block Glue** mode allows you to <u>shrink or remove gaps</u> (in the selected sequences) by clicking and dragging.

Step 4.

The **Select** mode allows you to <u>select regions within one or more sequences</u>.