



As seen in the image above, users can view genome annotations against a reference “ruler,” with an overhead bar giving a visual indication of chromosome position. The user can navigate by dragging the display left or right or by clicking on navigation buttons; analogous buttons allow the display to be zoomed in or out. Alternatively, users can navigate directly to a region (or feature) of interest by typing the region coordinates (or feature name) into a search box. Additional annotation tracks can be added to the current display by dragging them from a reservoir bar on the left of the screen, and can be removed by dragging them back off the main display. Tracks can, similarly, be reordered by dragging. All track manipulation, as with navigation, is live and requires no page reloads

#### References:

<https://genomebiology.biomedcentral.com/articles/10.1186/s13059-016-0924-1>