

Copying Qualifier Text

Use the Copy/Paste->*Copy qualifier '[the qualifier name]'* value context menu item to copy the qualifier value.

The screenshot displays the NCBI SRA Explorer interface. On the left, a tree view shows the features of the sequence NC_004718, including 3'UTR, 5'UTR, and CDS. The CDS feature is selected, and a context menu is open over it. The menu contains various actions such as 'Find qualifier...', 'Invert annotation selection', 'Add \'db_xref\' column', 'Go to position...', 'Select sequence region...', 'New annotation...', 'Copy/Paste', 'Select', 'Add', 'Analyze', 'Align', 'Cloning', 'Export', 'Edit', and 'Remove'. The 'Copy/Paste' option is highlighted, and a sub-menu is visible showing options like 'Copy selected sequence', 'Copy selected complementary 5'-3\' sequence', 'Copy amino acids', 'Copy amino acids of complementary 5'-3\' strand', 'Copy annotation sequence', 'Copy annotation amino acids', 'Copy qualifier \'db_xref\' value', and 'Paste sequence'. The background shows a list of features for NC_004718, including 3'UTR, 5'UTR, and CDS. The CDS feature is selected, and a context menu is open over it.