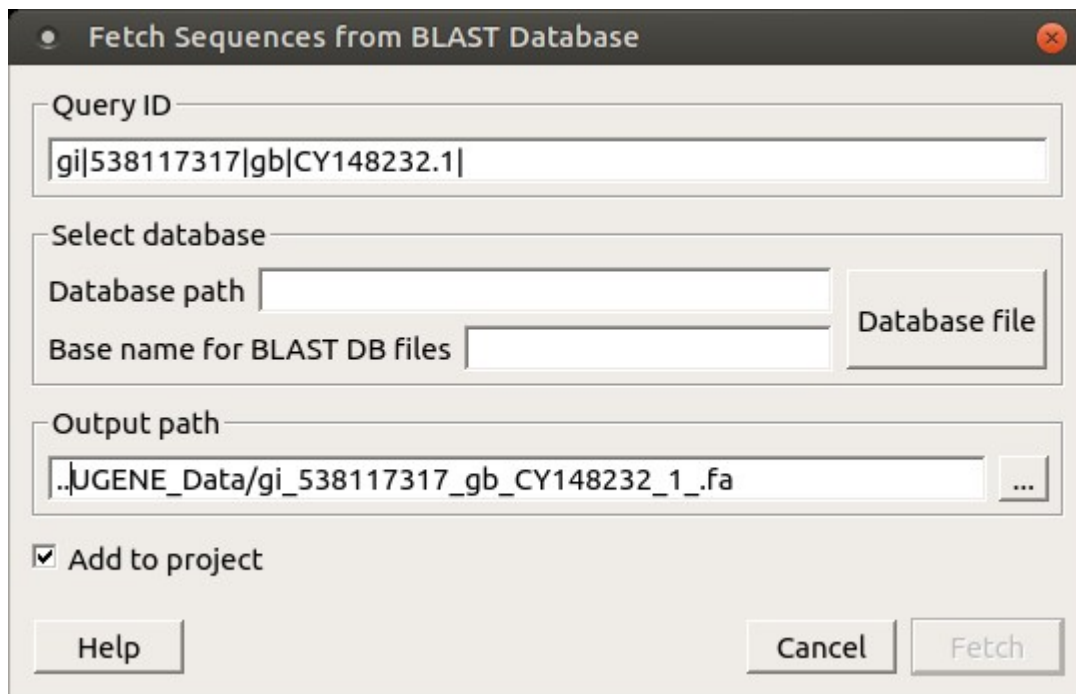


Fetching Sequences from Local BLAST Database

To fetch sequences from the local BLAST+ database use the *Fetch sequences from local BLAST+ database->Fetch sequences by 'id' from 'blast result'* context menu item of the blast result. The following dialog will appear:



The dialog box is titled "Fetch Sequences from BLAST Database". It contains the following fields and controls:

- Query ID:** A text field containing the text "gi|538117317|gb|CY148232.1|".
- Select database:** A section containing two text fields: "Database path" and "Base name for BLAST DB files". To the right of these fields is a button labeled "Database file".
- Output path:** A text field containing the text "..\\UGENE_Data\\gi_538117317_gb_CY148232_1_.fa". To the right of the field is a button with three dots "...".
- Add to project:** A checkbox that is checked, with the label "Add to project".
- Buttons:** At the bottom of the dialog are three buttons: "Help", "Cancel", and "Fetch".

Here you need to select a query ID, database, type of file(s) and output path. After that click on the *Fetch* button. To fetch sequences for several annotations at the same time select the blast results with *Ctrl* key and call the *Fetch sequences from local BLAST database->Fetch sequences by 'id' from 'blast result'* context menu item.