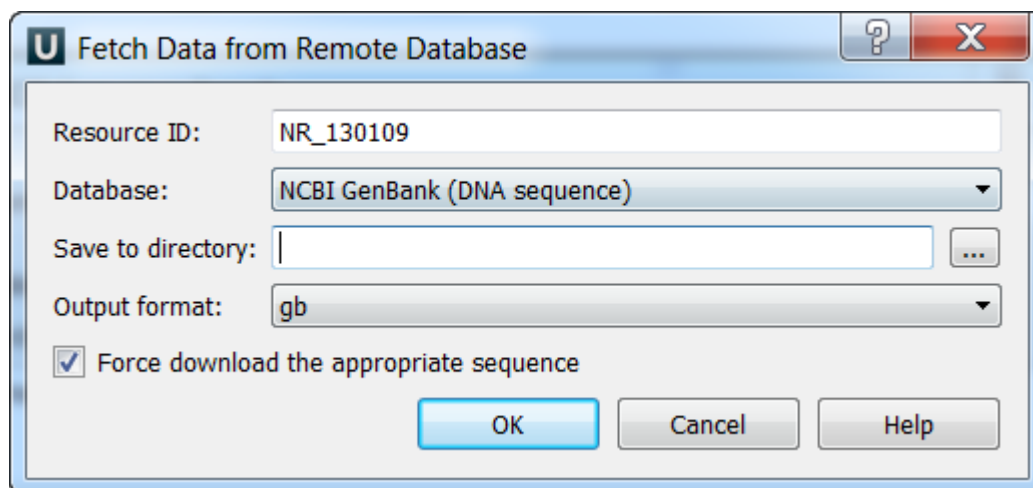


Fetching Data from Remote Database

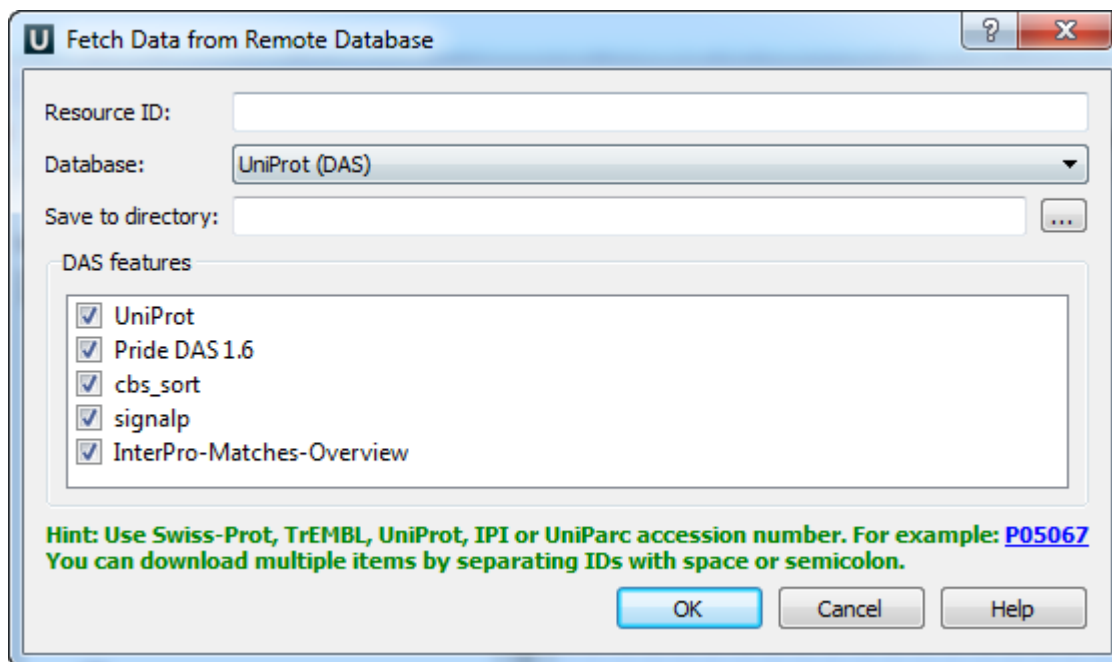
UGENE allows fetching data from remote biological databases such as NCBI GenBank, NCBI protein sequence database and some others.

To fetch data select the *File Access remote database...* item in the main menu.

The dialog will appear:



Here you need to enter unique id of the biological object and choose a database. The following databases are available: NCBI Genbank (DNA sequence), NCBI protein sequence database, ENSEMBL, PDB, SWISS-PROT, UniProtKB/Swiss-Prot, UniProtKB/TrEMBL, UniProt (DAS), Ensembl Human Genes (DAS). Unique identifiers are different for various databases. For example, for NCBI GenBank such unique id could be [Accession Number](#) or [NCBI GI number](#). If you select the UniProt (DAS) or Ensembl Human Genes (DAS) database you can select the DAS features. For example:



Optionally, you can browse for a directory to save the fetched file to.

After you click the *OK* button, UGENE downloads the biological object (DNA sequence, protein sequence, 3d model, etc.) and adds it to the current *project*.

If something goes wrong check the [Log View](#), it will help you to diagnose the problem.