



## Searching for Data

Sequence Read Archive Submissions Staff

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This section of the Sequence Read Archive (SRA) Knowledge Base contains:

- Instructions for searching for specific data types
- Suggestions for search terms
- Instructions for using SRA search tools
- Answers to common problems that happen while searching
- Information on how to interpret your search results.

**To begin searching this section of the Sequence Read Archive Knowledge Base, you can either:**

- **Enter your search word(s)** text in the text box **at the top of the page** and click on the “Go” button,

**OR**

- **Clickon** any of the “SRA Knowledge Base” **sub-categories listed** in the “Contents” section **below** to go to the sub-category of your choice.

Searching for Species-Specific Data

Searching for Specific Study Types

Searching for Specific Molecule Types

Searching for Data Generated by a Specific Platform or Instrument

Searching for Recently Submitted Data Accession Numbers

Searching for New Studies added to SRA

Searching for Data Generated by a Specific Next-Generation Sequencing Application

Searching for Data by Submitter or Author

Searching for PubMed Articles Associated with SRA Data

Searching for Data Referenced in a Publication

Searching using BLAST

Problems Searching SRA

Understanding SRA Search Results