

# dbGaP Submission Survey - 2022

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Start of Block: Default Question Block

OMB Statement OMB Control Number: **0925-0648**

Expiration Date: **06/30/2024**

Public reporting burden for this collection of information is estimated to average **7 minutes per response**, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. An agency may not conduct or sponsor, and a person is not required to respond to, a collection of information unless it displays a current valid OMB control number. Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden, to NIH, Project Clearance Branch, 6705 Rockledge Drive, MSC 7974, Bethesda, MD 20892-7974, ATTN: PRA (0925-0648). Do not return the completed form to this address.

*\* Indicates a required question*

Q1 Please select one professional category that best describes you. \*

- Life Sciences Researcher
- Geneticist
- Genetic Counselor
- Laboratory Staff
- Physician
- Other Healthcare Professional
- Bioinformatics Professional
- Computer Scientist / Software Developer
- Educator
- Student
- Librarian / Information Specialist
- Patient and Family
- Other (please specify below) \_\_\_\_\_

Q2 How likely are you to recommend this resource to a friend or colleague? \*

- 0
- 1
- 2
- 3
- 4
- 5
- 6
- 7

- 8
  - 9
  - 10
- 

Q3 Were you able to complete your dbGaP-related work to your satisfaction? \*

- No
  - Unsure
  - Yes
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*Display This Question:*

*If Q3 = No*

Q4a If No to the previous question, was it because: (choose all that apply)

- Submission requirements were not clear
  - SRA data submission, validation, and data processing was not clear
  - Molecular data submission, validation, and data processing was not clear
  - Instructions on how to use Submission Portal were not clear
  - An error was reported that was not clear
  - Additional comments / Other (please specify below)
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Display This Question:

If Q3 = Unsure

Or Q3 = Yes

Q4b We recently released the features listed below. Please rate each based on your level of satisfaction.

	Dissatisfied	Somewhat Dissatisfied	Somewhat Satisfied	Satisfied	Don't Know	Not Applicable
Automated validation from the Submission Portal for faster feedback on errors	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
GaPTools, a stand-alone tool for pre-submission validation	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
New web form for study config data entry	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Updated web form for study data outline with the ability to compare with what was provided at registration	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Ability to upload sequence metadata file via the Submission Portal	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Faster release of studies after all data has been submitted	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

Q5 Is this your first time submitting a study to dbGaP? \*

Yes

No

Display This Question:

If Q5 = No

Q6a Please drag and drop the tasks below in order from easiest (1) to hardest (7).

- \_\_\_\_\_ Registering your study
- \_\_\_\_\_ Preparing data / formatting data for a submission
- \_\_\_\_\_ Finding the status of your submission
- \_\_\_\_\_ Interpreting error / failure reports sent after one or more files are processed by the system
- \_\_\_\_\_ Getting timely answers to your questions
- \_\_\_\_\_ Interpreting messages from curators
- \_\_\_\_\_ Getting your entire submission released as quickly as possible

Display This Question:

If Q5 = Yes

Q6b Please drag and drop the tasks below in order from easiest (1) to hardest (4).

- \_\_\_\_\_ Registering your study
- \_\_\_\_\_ Preparing data / formatting data for a submission
- \_\_\_\_\_ Finding the status of your submission
- \_\_\_\_\_ Interpreting error / failure reports sent after one or more files are processed by the system

Q7 What type of data do you personally submit (choose all that apply)? \*

- Study Metadata (Study Config web form)
- Subject and Sample IDs (Subject Consent and Subject Sample Mapping files)
- Phenotype data (Demographic, Clinical or Exposure on person or Sample attributes)
- Molecular data (Genotype calls, Expression data or counts)
- Sequence data
- Analysis data
- Other (please specify below) \_\_\_\_\_

Q8 We are considering several new features to help with submissions. Please drag / drop the items listed below to indicate your top priorities from the most useful (1) to the least useful (8).

- \_\_\_\_\_ A data model explaining how your data is parsed and stored in dbGaP
  - \_\_\_\_\_ A guide that provides a high-level overview of registration and submission steps
  - \_\_\_\_\_ A process flow diagram explaining your and other parties' roles and responsibilities for various registration and submission steps
  - \_\_\_\_\_ Sample study submission file templates with realistic data for different types of studies
  - \_\_\_\_\_ Tailored submission guide that can be customized to your needs
  - \_\_\_\_\_ Videos that explain registration and submission steps
  - \_\_\_\_\_ Planned and periodic calls between all submitters for your study and dbGaP experts
  - \_\_\_\_\_ Other (please specify)
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Q9 Please share any other suggestions you have for improving dbGaP submissions.

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