



NCBI Genome Reference Consortium User Survey

OMB Control Number: 0925-0648

Expiration Date: 03/31/2018

Public reporting burden for this collection of information is estimated to average 3 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.

NCBI is conducting this short survey on behalf of the Genome Reference Consortium (GRC) in order to gain insight for future improvements by the GRC to its user resources.

Next



NCBI Genome Reference Consortium User Survey

GRC Promotion

*** 1. How did you initially find out about the Genome Reference Consortium (GRC)?**

- Workshop/Conference
- From Colleagues
- Publication
- I didn't, I'm here by accident
- Other (please specify)

Prev

Next



NCBI Genome Reference Consortium User Survey

Purpose

*** 2. What was your primary reason for visiting the GRC website today?**

- Get general information about an assembly
- Search for information about one or more specific assembly regions
- Download data
- Contact the GRC with a question
- Other (please specify)

*** 3. Did the GRC website provide you with what you needed on this visit?**

- Yes
- No

Additional comments

4. What additional features would you add to this website?

Prev

Next



NCBI Genome Reference Consortium User Survey

Data Usage

*** 5. How would you characterize your primary interest in genomic sequence?**

- Basic research
- Clinical related
- Educational
- Other (please specify)

*** 6. How would you characterize your typical use of genomic sequence?**

- Single organism, Genome-scale
- Single organism, Single gene/gene family
- Multiple organism, Genome-scale
- Multiple organism, Single gene/gene family

*** 7. Which organism was the focus of your visit to the GRC website today?**

- Human
- Mouse
- Zebrafish
- Other (please specify)

Prev

Next



NCBI Genome Reference Consortium User Survey

Human Assembly Version

*** 8. What version(s) of the human reference genome assembly do you use?**

- GRCh38
- GRCh37
- Both GRCh37 and GRCh38
- Don't know
- Other (please specify)

Prev

Next



NCBI Genome Reference Consortium User Survey

GRCh37/GRCh38 Usage

*** 9. Have you ever used an alternate loci or patch scaffold in your analyses?**

Yes

No

Prev

Next



NCBI Genome Reference Consortium User Survey

GRCh37/GRCh38 Usage

*** 9. Do you plan to start using GRCh38?**

Yes

No

Additional comments

*** 10. Have you ever used an alternate loci or patch scaffold in your analyses?**

Yes

No

Prev

Next



NCBI Genome Reference Consortium User Survey

Alts and Patches

*** 11. If no, why not?**

- Not relevant to my work
- Lack of tools (e.g. too hard to use)
- Don't see the point
- Waiting for a graph genome representation
- Other (please specify)

Prev

Next



NCBI Genome Reference Consortium User Survey

Thank you

10. Additional comments:

Thank you for your participation. Your feedback is greatly appreciated.

Prev

Done