

5/14/2020 5:05 AM Questions about RaTG13 PCR amplicon sequences Incoming Received

From 544728460@qq.com
To sra@ncbi.nlm.nih.gov;sra@ncbi.nlm.nih.gov;
Attachment # 1

Dear editor,

I am writing to ask how to add PCR amplicon sequences to the submitted Sequence Read Archive (SRA) data?

I want to add PCR amplicon sequences to my submission : SUB6959238 (the SRA accession is SRR11085797). Could you help me with this or indicate me the way to updating ? Actually, I went throught the SRA Data Updates (<https://www.ncbi.nlm.nih.gov/sra/docs/submitupdate/#how-do-i-add-more-samples-to-my->) but didn't find a solution.

The PCR amplicon sequence files are attached to this E-mail.

Thanks for your kindly help! And look forward to your timely reply!

Best regards,

Si Haorui

Wuhan Institute of Virology

5/14/2020 9:59 AM Re: case #CAS-536334-L2Y4B3: Questions about RaTG13 PCR amplicon sequences TRACKING:000414000006481 Outgoing Sent

From nlm-support@nlm.nih.gov
To 544728460@qq.com;
Attachment #

Dear Si Haorui,

As it says in Update guide, you do not add sequences to your submission - you add new data to your bioproject.

Your bioproject accession is [PRJNA606165](#)

In order to add new data to your bioproject and/or biosample you need to make new SRA submission and on General Info page indicate that you already have bioproject for the dataset that you want to submit and provide bioproject's accession [PRJNA606165](#)

If your biosample is the same as in your previous submission - [SAMN14082201](#)- you also indicate that you already have biosample and provide its accession.

If the biosample is not the same, you indicate that you do not have biosample for this dataset.

After you click "Continue" you will be directed to proceed the submission accordingly.

Please follow step-by-step instructions on this page <https://www.ncbi.nlm.nih.gov/sra/docs/submitportal/>

Best regards,

Svetlana

If you have any questions or concerns regarding your **SRA** submission please don't hesitate to contact sra@ncbi.nlm.nih.gov (applies to new questions). We *normally* respond *within 2 business days*.

Svetlana Iazvovskaia
The NCBI SRA database submission staff

* PLEASE DO NOT MODIFY THE SUBJECT LINE OF THIS EMAIL WHEN RESPONDING TO ENSURE CORRECT TRACKING *

Case Information:

Case #: CAS-536334-L2Y4B3

Customer Name: 独钓寒江雪

Customer Email: 544728460@qq.com

Case Created: 2020-05-14T09:06:01Z

Summary: Questions about RaTG13 PCR amplicon sequences

Details:

Dear editor,

I am writing to ask how to add PCR amplicon sequences to the submitted Sequence Read Archive (SRA) data?

I want to add PCR amplicon sequences to my submission : SUB6959238 (the SRA accession is SRR11085797). Could you help me with this or indicate me the way to updating ? Actually, I went throught the SRA Data Updates (<https://www.ncbi.nlm.nih.gov/sra/docs/submitupdate/#how-do-i-add-more-samples-to-my->) but didn't find a solution.

The PCR amplicon sequence files are attached to this E-mail.

Thanks for your kindly help! And look forward to your timely reply!

Best regards,

Si Haorui

Wuhan Institute of Virology

5/17/2020 5:47 AM Undeliverable: Case CAS-536334-L2Y4B3 - National Library of Medicine Customer Service confirmation TRACKING:000435000441374 Incoming Received

From MAILER-DAEMON@nihcesxwayst05.hub.nih.gov

To 544728460@qq.com;

Attachment # 1

The following message to <544728460@qq.com> was undeliverable.

The reason for the problem:

5.4.7 - Delivery expired (message too old) 'timeout'

5/17/2020 10:26 AM Undeliverable: case #CAS-536334-L2Y4B3: Questions about RaTG13 PCR amplicon sequences TRACKING:000414000006481 Incoming Received

From MAILER-DAEMON@nihcesxwayst03.hub.nih.gov

To 544728460@qq.com;

Attachment # 1

The following message to <544728460@qq.com> was undeliverable.

The reason for the problem:

5.4.7 - Delivery expired (message too old) 'timeout'



Original Cc line: taxonomy@ncbi.nlm.nih.gov, turner@ncbi.nlm.nih.gov,
andrei.vereshchagin@nih.gov, genomes@ncbi.nlm.nih.gov
Original To line: mmail_submit@ncbi.nlm.nih.gov

54962 silage metagenome

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606030](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606030)

54966 biofilm metagenome

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606093](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606093)

54967 soil metagenome

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606094](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606094)

54968 Psychroflexus sp. YR1-1

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606115](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606115)

54969 Aliifodinibius halophilus

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606119](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606119)

54970 Balneolaceae bacterium YR4-1

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606120](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606120)

54971 soil metagenome

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606122](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606122)

54976 Acinetobacter sp. Acinetobacter sp

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submitter must provide a strain/isolate identifier

=====

http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606152

54977 Pseudoalteromonas sp. Pseudoalteromonas sp.

=====
submitter must provide a strain/isolate identifier
=====

http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606154

54978 Ochrobactrum sp. Ochrobactrum sp.

=====
submitter must provide a strain/isolate identifier
=====

http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606156

54979 Jeotgalicoccus sp. Jeotgalicoccus sp.

=====
submitter must provide a strain/isolate identifier
=====

http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606157

54980 Delftia sp. Delftia sp.

=====
submitter must provide a strain/isolate identifier
=====

http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606158

54981 Rhodococcus sp. Rhodococcus sp.

=====
submitter must provide a strain/isolate identifier
=====

http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606160

-- Se n Turner

-----Original Message-----

From: Taxonomy <taxonomy-bounces@ncbi.nlm.nih.gov> on behalf of
NLM/NCBI mmml_submit <mmml_submit@ncbi.nlm.nih.gov>
Reply-To: NLM/NCBI mmml_submit <mmml_submit@ncbi.nlm.nih.gov>
Date: Wednesday, February 12, 2020 at 10:00 AM
To: NLM/NCBI List taxonomy <taxonomy@ncbi.nlm.nih.gov>
Subject: TaxConsult: submission_portal 14674 [bioproject_ids
PRJNA606024, PRJNA606030, PRJNA606039, PRJNA606073, PRJNA606081,
PRJNA606093, PRJNA606094, PRJNA606115, PRJ] [reply_thread_id=6457]

54961 *Vitis* hybrid
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606024](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606024)

54962 Culture Media Silage Microbiome
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606030](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606030)

54963 *Paenibacillus* sp. ALJ109b
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606039](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606039)

54964 *Wolbachia* endosymbiont of *Tyrophagus putrescentiae*
=====

[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606073](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606073)

54965 *Starkeya* sp. ORNL1
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606081](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606081)

54966 Carbon felt biofilm
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606093](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606093)

54967 Soil sample
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606094](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606094)

54968 *Psychroflexus* sp. YR1-1 *Psychroflexus* bacterium
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606115](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606115)

54969 *Aliifodinibius halophilus* 2W32
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[http://intrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606119](http://intrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606119)

54970 Balneolaceae gen. nov. sp. YR4-1 microbial
=====

[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606120](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606120)

54971 soil sample
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606122](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606122)

54972 Cattle gut metagenome
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606123](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606123)

54973 Rhodobacter sp. ETT8
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606125](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606125)

54974 Caulobacter sp. 17J65-9
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606126](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606126)

54975 Noviherbaspirillum sp. 17J57-3
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606127](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606127)

54976 Acinetobacter sp. Acinetobacter sp
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606152](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606152)

54977 Pseudoalteromonas sp. Pseudoalteromonas sp.
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606154](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606154)

54978 Ochrobactrum sp. Ochrobactrum sp.
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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606156](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606156)

54979 Jeotgalicoccus sp. Jeotgalicoccus sp.

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606157](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606157)

54980 Delftia sp. Delftia sp.

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606158](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606158)

54981 Rhodococcus sp. Rhodococcus sp.

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606160](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606160)

54982 Bat coronavirus isolate RaTG13

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[http://inrawebdev2.be-
md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606165](http://inrawebdev2.be-md.ncbi.nlm.nih.gov/projects/bp/bpedit.cgi?pid=PRJNA606165)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov
Mon Feb 17 06:10:39 EST 2020

Dear Zhengli Shi,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6959189
BioProject ID: PRJNA606165
Title:
Locus tag prefix:
G3591

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6959189/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA606165 with your correspondence and your data submissions. Use the registered locus tag prefix when you include annotation in your submission.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff
Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov
Wed Feb 12 11:10:18 EST 2020

Dear Zhengli Shi,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6959189
BioProject ID: PRJNA606165
Title:
Locus tag prefix:
G3591

has been successfully registered with the BioProject database. After review by the database staff, your project information will be accessible with the following link, usually within a few days of the release date that you set (or the release of linked data, whichever is first):

<http://www.ncbi.nlm.nih.gov/bioproject/606165>

Please use the BioProject ID PRJNA606165 with your correspondence and your data submissions.
Use the registered locus tag prefix when you include annotation in your submission.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff
Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov
Wed May 20 06:20:38 EDT 2020

Dear Zhengli Shi,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6959189
BioProject ID: PRJNA606165
Title:
Locus tag prefix:
G3591 (SAMN14082201)

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6959189/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA606165 with your correspondence and your data submissions. Use the registered locus tag prefix when you include annotation in your submission.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff
Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov
Thu Apr 30 12:44:58 EDT 2020

Dear Zhengli Shi,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6959189
BioProject ID: PRJNA606165
Title:
Locus tag prefix:
G3591 (SAMN14082201)

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6959189/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA606165 with your correspondence and your data submissions. Use the registered locus tag prefix when you include annotation in your submission.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff
Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov
Mon May 18 14:10:46 EDT 2020

Dear Zhengli Shi,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6959189
BioProject ID: PRJNA606165
Title:
Locus tag prefix:
G3591 (SAMN14082201)

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6959189/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA606165 with your correspondence and your data submissions. Use the registered locus tag prefix when you include annotation in your submission.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff
Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

3/23/2020 3:26 AM change the release time of the Bioprojec : PRJNA606875 Incoming Received

From 907904153@qq.com
To sra@ncbi.nlm.nih.gov;sra@ncbi.nlm.nih.gov;
Attachment #

Dear prof:

I am sorry to trouble you but I would like to change the release time of the Bioproject: PRJNA606875.

Would you please advance the release time of all the seven Biosamples of this project to March 25, 2020.

Thank you very much.

Yours Sincerely,
Ya-Wei Zhang

3/23/2020 5:29 PM Re: case #CAS-508999-C2V8Q9: change the release time of the Bioprojec : PRJNA606... TRACKING:000422000002553 Outgoing Sent

From nlm-support@nlm.nih.gov
To 907904153@qq.com;
Attachment #

Dear Ya-Wei Zhang,
You can review and update release date by yourself anytime:
<https://dataview.ncbi.nlm.nih.gov/object/PRJNA606875>

Feel free to contact us with any questions or concerns.

Thanks,
Inna

==

Inna Belaia
The SRA submissions staff
NCBI/NLM/NIH

* PLEASE DO NOT MODIFY THE SUBJECT LINE OF THIS EMAIL WHEN RESPONDING TO ENSURE CORRECT TRACKING *

Case Information:

Case #: CAS-508999-C2V8Q9

Customer Name: 907904153

Customer Email: 907904153@qq.com

Case Created: 2020-03-23T07:27:00Z

Summary: change the release time of the Bioprojec : PRJNA606875

Details:

Dear prof:

I am sorry to trouble you but I would like to change the release time of the Bioproject: PRJNA606875.

Would you please advance the release time of all the seven Biosamples of this project to March 25, 2020.

Thank you very much.

Yours Sincerely,

Ya-Wei Zhang

3/26/2020 3:44 AM Undeliverable: Case CAS-508999-C2V8Q9 - National Library of Medicine Customer Service confirmation TRACKING:000435000403803 Incoming Received

From MAILER-DAEMON@nihcesxway4.hub.nih.gov
To 907904153@qq.com;

Attachment # 1

The following message to <907904153@qq.com> was undeliverable.

The reason for the problem:

5.4.7 - Delivery expired (message too old) 'timeout'

3/26/2020 6:01 PM Undeliverable: case #CAS-508999-C2V8Q9: change the release time of the Bioprojec : PRJNA606... TRACKING:000422000002553 Incoming Received

From MAILER-DAEMON@nihcesxwayst04.hub.nih.gov
To 907904153@qq.com;

Attachment # 1

The following message to <907904153@qq.com> was undeliverable.
The reason for the problem:
5.4.7 - Delivery expired (message too old) 'timeout'

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov

Sat Feb 15 07:54:27 EST 2020

Dear Yawei Zhang,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6975258 BioProject ID:

PRJNA606875

Title:

has been successfully registered with the BioProject database. After review by the database staff, your project information will be accessible with the following link, usually within a few days of the release date that you set (or the release of linked data, whichever is first):

<http://www.ncbi.nlm.nih.gov/bioproject/606875>

Please use the BioProject ID PRJNA606875 with your correspondence and your data submissions.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff

Bethesda, Maryland USA

(301) 496-2475 (301) 480-2918 (Fax) bioprojecthelp at ncbi.nlm.nih.gov

(for BioProject questions/replies) info at ncbi.nlm.nih.gov (for general

questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov

Thu Mar 26 21:50:46 EDT 2020

Dear Yawei Zhang,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6975258

BioProject ID: PRJNA606875

Title:

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6975258/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA606875 with your correspondence and your data submissions.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff

Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov

Mon May 18 14:10:45 EDT 2020

Dear Yawei Zhang,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6975258

BioProject ID: PRJNA606875

Title:

Locus tag prefixes:

None (SAMN14115940)

None (SAMN14115941)

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6975258/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA606875 with your correspondence and your data submissions.

Use the registered locus tag prefixes when you include annotation in your submission.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff

Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov

Wed May 20 14:11:31 EDT 2020

Dear Yawei Zhang,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6975258

BioProject ID: PRJNA606875

Title:

Locus tag prefixes:

None (SAMN14115940)

None (SAMN14115941)

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6975258/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA606875 with your correspondence and your data submissions.

Use the registered locus tag prefixes when you include annotation in your submission.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff

Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

sra at ncbi.nlm.nih.gov sra at ncbi.nlm.nih.gov

Thu Mar 26 21:47:04 EDT 2020

Dear Yawei Zhang,

This email confirms that NCBI has completed your request to change the release date of the data associated with the BioProject PRJNA606875 so that it will not be publicly-released until 2020-03-28.

Note that the following linked data related to this BioProject will also not be released until your new requested BioProject release date.

BioProject: PRJNA606875

Linked accessions are in the attachment.

You may change the release date in <https://dataview.ncbi.nlm.nih.gov/> if you are ready to release your data sooner than the date above.

Thank you!

----- next part -----

BioSamples:

SAMN14115940

SAMN14115941

SAMN14115942

SAMN14115943

SAMN14115944

SAMN14115945

SAMN14116618

SRA:

SRR11093271

SRR11093270

SRR11093269

SRR11093268

SRR11093267

SRR11093266

SRR11093265

sra at ncbi.nlm.nih.gov sra at ncbi.nlm.nih.gov

Thu Mar 26 21:48:05 EDT 2020

Dear Yawei Zhang,

We've received your request to update the release date for the BioProject below to 2020-03-26.

Note that the following linked data will also be released on the same date:

BioProject: PRJNA606875

Linked accessions are in the attachment.

If you have requested that Sequence Read Archive (SRA) data's release be deferred, you will receive a reminder of your upcoming release 2 weeks prior to the release date.

A final email will be generated to confirm the successful release of your data. Please contact sra at ncbi.nlm.nih.gov with any urgent questions or concerns.

Thank you!

----- next part -----

BioSamples:

SAMN14115940

SAMN14115941

SAMN14115942

SAMN14115943

SAMN14115944

SAMN14115945

SAMN14116618

SRA:

SRR11093271

SRR11093270

SRR11093269

SRR11093268

SRR11093267

SRR11093266

SRR11093265

sra at ncbi.nlm.nih.gov sra at ncbi.nlm.nih.gov

Thu Mar 26 21:49:06 EDT 2020

Dear Yawei Zhang,

This email confirms that NCBI has completed your request to release the data associated with the BioProject on 2020-03-26. Please allow four hours for the data to be indexed and searchable in Entrez.

Note that the following linked data has also been released to the public with the BioProject's release.

BioProject: PRJNA606875

Please contact sra at ncbi.nlm.nih.gov if you need to suppress this data from the public view, make other modifications, or if it has been 2 weeks and you do not see all your data public at <https://www.ncbi.nlm.nih.gov/bioproject/PRJNA606875>.

Thank you!

sra at ncbi.nlm.nih.gov sra at ncbi.nlm.nih.gov

Thu Mar 26 21:46:30 EDT 2020

Dear Yawei Zhang,

We've received your request to update the release date for the BioProject below to 2020-03-28.

Note that the following linked data will also be released on the same date:

BioProject: PRJNA606875

Linked accessions are in the attachment.

If you have requested that Sequence Read Archive (SRA) data's release be deferred, you will receive a reminder of your upcoming release 2 weeks prior to the release date.

A final email will be generated to confirm the successful release of your data. Please contact sra at ncbi.nlm.nih.gov with any urgent questions or concerns.

Thank you!

----- next part -----

BioSamples:

SAMN14115940

SAMN14115941

SAMN14115942

SAMN14115943

SAMN14115944

SAMN14115945

SAMN14116618

SRA:

SRR11093271

SRR11093270

SRR11093269

SRR11093268

SRR11093267

SRR11093266

SRR11093265

3/25/2020 9:45 PM change the release time of the Bioprojec : PRJNA606875 Incoming Received

From 15202206702@163.com
To sra@ncbi.nlm.nih.gov;sra@ncbi.nlm.nih.gov;
Attachment #

Dear prof:

I am sorry to trouble you but I would like to change the release time of the Bioproject : PRJNA606875.

Would you please advance the release time of all the seven Biosamples of this project to March 26, 2020, or assoon as possible.

Thank you very much.

Yours Sincerely,
Ya-Wei Zhang

3/26/2020 11:47 AM case #CAS-510715-D9N5B2: RE: change the release time of the Bioprojec : PRJN... TRACKING:000393000005681 Outgoing Sent

From nlm-support@nlm.nih.gov
To 15202206702@163.com;
Attachment #

You can change the release date of unreleased SRA projects by logging into submission portal and navigating to the "Manage data" view (<https://dataview.ncbi.nlm.nih.gov/>) for your study. Select "Edit" release date and enter the new date. Please allow several hours following release for your data to become indexed and searchable in Entrez. More details are here: <https://www.ncbi.nlm.nih.gov/sra/docs/submitsra/>

<https://dataview.ncbi.nlm.nih.gov/object/PRJNA606875>

Cheers,
Rick Lapoint
SRA Curator

----- Original Message -----

From: 张雅薇;

Received: Wed Mar 25 2020 21:40:32 GMT-0400 (Eastern Daylight Time)

To: NLM/NCBI List sra; SRA Support;

Subject: change the release time of the Bioprojec : PRJNA606875

Dear prof:

I am sorry to trouble you but I would like to change the release time of the Bioproject : PRJNA606875.

Would you please advance the release time of all the seven Biosamples of this project to March 26, 2020, or assoon as possible.

Thank you very much.

Yours Sincerely,

Ya-Wei Zhang

2/18/2020
9:18 AM

Potential SPAM:Change about the bioproject PRJNA606875

Incoming Received

From 907904153@qq.com
To sra@ncbi.nlm.nih.gov; sra@ncbi.nlm.nih.gov;
Attachment #

Dear prof:

This is the second email about the change about the bioproject PRJNA606875. The first email is below this email.

The second email is to add that I also uploaded the new data through FTP to the same folder as before,

“/uploads/907904153_qq.com_CZ9W7QFO/new_folder”.

Address : <ftp-private.ncbi.nlm.nih.gov>

Username :

The new file uploaded via FTP is exactly the same as the attachment in the first email, you can choose to change it in your convenience.

We sincerely hope you could provide us with the numbers of SRA first, as they are urgently requested by the journal editor

who are assessing our study.

Look forward to your reply. Thank you very much.

Yours Sincerely,

Ya-Wei Zhang

Dear prof:

I'm sorry to email you that the files I submitted yesterday (Bioproject: PRJNA606875)has some formatting issues. I'D like to change the data files.

Only the data file needs to be changed, other information does not need to be changed. Could you please help me change the file?

Bioproject: PRJNA606875

Biosample:

SAMN14115940: (SRA:[SRR11093271](#)): use the new file"GX-P2V.fastq"

SAMN14115941: (SRA:[SRR11093270](#)): use the new file"GX-P3B.fastq"

SAMN14115942: (Unaccessioned): use the new file"GX-P4L-1.gz" and "GX-P4L-2.gz". Change from the previous one file to the current two files.

SAMN14115943: (SRA:[SRR11093268](#)):use the new file"GX-P5E-1.gz" and "GX-P5E-2.gz". Change from the previous one file to the current two files.

SAMN14115944: (SRA:[SRR11093267](#)):use the new file"GX-P5L-1.gz" and "GX-P5L-2.gz". Change from the previous one file to the current two files.

SAMN14115945: (SRA:[SRR11093266](#)):use the new file"GX-P1E-1.gz" and "GX-P1E-2.gz". Change from the previous one file to the current two files.

All new data is in the email attachment.I compressed the data, all data should be in fastq format after decompression.

Look forward to your reply. Thank you very much.

Yours Sincerely,
Ya-Wei Zhang

从QQ邮?发???大附?

[CHANGE about bioproject.zip](#) (1.27G, 2020年03月18? 19:41 ?期)

[进?下载页?](#)

2/18/2020 11:09 AM Re: case #CAS-492827-M9B7K9: Potential SPAM:Change about the bioproject PRJNA6... TRACKING:000393000005339 Outgoing Sent

From nlm-support@nlm.nih.gov

To 907904153@qq.com;

Attachment #

The replacement files are now linking to your SRA submission and should be processed by tomorrow.

Please refer to this data using the accession PRJNA606875 in future publications. Unfortunately there is currently no way to allow selective, pre-release access to SRA sequence data for colleagues and reviewers only. A link can be generated that will direct reviewers to the metadata for your SRA submission including submission structure, sample metadata, number of bases and spots loaded, etc.

To create this link please log into submission portal once your submission is complete and go to the "Manage Data" page for PRJNA606875 (<https://dataview.ncbi.nlm.nih.gov/object/PRJNA606875>) and click the "create reviewer link" button at the top of the page.

Cheers,

Rick Lapoint
SRA Curator

* PLEASE DO NOT MODIFY THE SUBJECT LINE OF THIS EMAIL WHEN RESPONDING TO ENSURE CORRECT TRACKING *

Case Information:

Case #: CAS-492827-M9B7K9

Customer Name: 907904153

Customer Email: 907904153@qq.com

Case Created: 2020-02-18T14:18:31Z

Summary: Potential SPAM:Change about the bioproject PRJNA606875

Details:

Dear prof:

This is the second email about the change about the bioproject PRJNA606875. The first email is below this email.

The second email is to add that I also uploaded the new data through FTP to the same folder as before,

“/uploads/907904153_qq.com_CZ9W7QFO/new_folder”.

Address : ftp-private.ncbi.nlm.nih.gov

Username :

The new file uploaded via FTP is exactly the same as the attachment in the first email, you can choose to change it in your convenience.

We sincerely hope you could provide us with the numbers of SRA first, as they are urgently requested by the journal editor who are assessing our study.

Look forward to your reply. Thank you very much.

Yours Sincerely,

Ya-Wei Zhang

Dear prof:

I'm sorry to email you that the files I submitted yesterday (Bioproject: PRJNA606875)has some formatting issues. I'd like to change the data files.

Only the data file needs to be changed, other information does not need to be changed. Could you please help me change the file?

Bioproject: PRJNA606875

Biosample:

SAMN14115940: (SRA:[SRR11093271](#)): use the new file"GX-P2V.fastq"

SAMN14115941: (SRA:[SRR11093270](#)): use the new file"GX-P3B.fastq"

SAMN14115942: (Unaccessioned): use the new file"GX-P4L-1.gz" and "GX-P4L-2.gz". Change from the previous one file to the current two files.

SAMN14115943: (SRA:[SRR11093268](#)):use the new file"GX-P5E-1.gz" and "GX-P5E-2.gz". Change from the previous one file to the current two files.

SAMN14115944: (SRA:[SRR11093267](#)):use the new file"GX-P5L-1.gz" and "GX-P5L-2.gz". Change from the previous one file to the current two files.

SAMN14115945: (SRA:[SRR11093266](#)):use the new file"GX-P1E-1.gz" and "GX-P1E-2.gz". Change from the previous one file to the current two files.

All new data is in the email attachment.I compressed the data, all data should be in fastq format after decompression.

Look forward to your reply. Thank you very much.

Yours Sincerely,
Ya-Wei Zhang

从QQ邮?发???大附?

CHANGE about bioproject.zip (1.27G, 2020年03月18? 19:41 ?期)

[进?下载页?](#)

2/21/2020 10:07 AM Undeliverable: Case CAS-492827-M9B7K9 - National Library of Medicine Customer Service confirmation TRACKING:000435000381360 Incoming Received

From MAILER-DAEMON@nihcesxwayst06.hub.nih.gov

To 907904153@qq.com;

Attachment # 1

The following message to <907904153@qq.com> was undeliverable.

The reason for the problem:

5.4.7 - Delivery expired (message too old) 'timeout'

2/21/2020 11:34 AM Undeliverable: case #CAS-492827-M9B7K9: Potential SPAM:Change about the bioproject PRJNA6... TRACKING:000393000005339 Incoming Received

From MAILER-DAEMON@nihcesxwayst03.hub.nih.gov

To 907904153@qq.com;

Attachment # 1

The following message to <907904153@qq.com> was undeliverable.

The reason for the problem:

5.4.7 - Delivery expired (message too old) 'timeout'

6/19/2020 1:24 AM

Re:BioProject ID PRJNA607174

Incoming Received

From xiaokpcc@163.com
To bioprojecthelp@ncbi.nlm.nih.gov;

Attachment #

Dear Curator:

The SRA data in the bioproject PRJNA607174 had been released few days ago, but I want upload one more SRA data in this bioproject.

Can I update a new SRA data under this project? If allowed, please tell me how to do it.

I look forward to your reply as soon as possible.

Thank you very much!

Kangpeng Xiao

At 2020-04-22 17:01:32, bioprojecthelp@ncbi.nlm.nih.gov wrote:

>Dear Kangpeng Xiao,

>

>This is an automatic acknowledgment that your submission:

>

>SubmissionID: SUB6982752

>BioProject ID: PRJNA607174

>Title:

>

>has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for

>each linked BioSample are included in the locustagprefix.txt file that can accessed

>from this BioProject in the submission portal:

>

><https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6982752/overview>

>

>In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

>

><https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

>

>Please use the BioProject ID PRJNA607174 with your correspondence and your data submissions.

>

>Send questions to bioprojecthelp@ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

>

>Regards,

>

>NCBI BioProject Submissions Staff

>Bethesda, Maryland USA

>*****

>(301) 496-2475

>(301) 480-2918 (Fax)

>bioprojecthelp@ncbi.nlm.nih.gov (for BioProject questions/replies)

>info@ncbi.nlm.nih.gov (for general questions regarding NCBI)

>*****

6/19/2020 3:16 PM Re: case #CAS-557246-T9N1F3: Re:BioProject ID PRJNA607174 TRACKING:000300000004709

Outgoing Sent

From nlm-support@nlm.nih.gov

To xiaokpcc@163.com;

Attachment #

Dear Kangpeng Xiao,

Thank you for your email. You can submit additional Biosamples and/or sequence data for an existing BioProject at any time. Simply initiate a new submission in the appropriate submission portal wizard. You can enter the BioProject ID for your existing BioProject, and the new submission will be linked automatically.

If you have other comments or questions, please reply to bioprojecthelp@ncbi.nlm.nih.gov.

Best regards,
Carol DeWeese Scott, Ph.D.
BioProject Curation Staff

* PLEASE DO NOT MODIFY THE SUBJECT LINE OF THIS EMAIL WHEN RESPONDING TO ENSURE CORRECT TRACKING *

Case Information:

Case #: CAS-557246-T9N1F3

Customer Name: 肖康鹏

Customer Email: xiaokpcc@163.com

Case Created: 2020-06-19T05:24:04Z

Summary: Re:BioProject ID PRJNA607174

Details:

Dear Curator:

The SRA data in the bioproject PRJNA607174 had been released few days ago, but I want upload one more SRA data in this bioproject.

Can I update a new SRA data under this project? If allowed, please tell me how to do it.

I look forward to your reply as soon as possible.

Thank you very much!

Kangpeng Xiao

At 2020-04-22 17:01:32, bioprojecthelp@ncbi.nlm.nih.gov wrote:

>Dear Kangpeng Xiao,

>

>This is an automatic acknowledgment that your submission:

>

>SubmissionID: SUB6982752

>BioProject ID: PRJNA607174

>Title:

>

>has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for

>each linked BioSample are included in the locustagprefix.txt file that can accessed

>from this BioProject in the submission portal:
>
><https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6982752/overview>
>
>In addition, you can view the locustagprefix.txt files for all of your
BioProjects from the BioProject submission page,
>
><https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.
>
>Please use the BioProject ID PRJNA607174 with your correspondence and your
data submissions.
>
>Send questions to bioprojecthelp@ncbi.nlm.nih.gov, and include the
BioProject ID and organism name.
>
>Regards,
>
>NCBI BioProject Submissions Staff
>Bethesda, Maryland USA
>*****
>(301) 496-2475
>(301) 480-2918 (Fax)
>bioprojecthelp@ncbi.nlm.nih.gov (for BioProject questions/replies)
>info@ncbi.nlm.nih.gov (for general questions regarding NCBI)
>*****

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov

Wed Apr 22 05:01:32 EDT 2020

Dear Kangpeng Xiao,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6982752

BioProject ID: PRJNA607174

Title:

has been updated, eg by the linkage of one or more BioSamples. The locus_tag prefixes for each linked BioSample are included in the locustagprefix.txt file that can accessed from this BioProject in the submission portal:

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/SUB6982752/overview>

In addition, you can view the locustagprefix.txt files for all of your BioProjects from the BioProject submission page,

<https://submit.ncbi.nlm.nih.gov/subs/bioproject/>.

Please use the BioProject ID PRJNA607174 with your correspondence and your data submissions.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff

Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov

Tue Feb 18 04:46:21 EST 2020

Dear Kangpeng Xiao,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB6982752

BioProject ID: PRJNA607174

Title:

has been successfully registered with the BioProject database. After review by the database staff, your project information will be accessible with the following link, usually within a few days of the release date that you set (or the release of linked data, whichever is first):

<http://www.ncbi.nlm.nih.gov/bioproject/607174>

Please use the BioProject ID PRJNA607174 with your correspondence and your data submissions.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff

Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)

sra at ncbi.nlm.nih.gov sra at ncbi.nlm.nih.gov
Wed Apr 22 01:41:16 EDT 2020

Dear Kangpeng Xiao,

This email confirms that NCBI has completed your request to release the data associated with the BioProject on 2020-04-22. Please allow four hours for the data to be indexed and searchable in Entrez.

Note that the following linked data has also been released to the public with the BioProject's release.

BioProject: PRJNA607174

Please contact sra at ncbi.nlm.nih.gov if you need to suppress this data from the public view, make other modifications, or if it has been 2 weeks and you do not see all your data public at <https://www.ncbi.nlm.nih.gov/bioproject/PRJNA607174>.

Thank you!

sra at ncbi.nlm.nih.gov sra at ncbi.nlm.nih.gov
Wed Apr 22 01:40:46 EDT 2020

Dear Kangpeng Xiao,

We've received your request to update the release date for the BioProject below to 2020-04-22.

Note that the following linked data will also be released on the same date:

BioProject: PRJNA607174

BioSamples:

SAMN14126321

SAMN14126322

SAMN14126323

SAMN14126324

SAMN14126325

SAMN14126326

SAMN14126329

If you have requested that Sequence Read Archive (SRA) data's release be deferred, you will receive a reminder of your upcoming release 2 weeks prior to the release date.

A final email will be generated to confirm the successful release of your data. Please contact sra at ncbi.nlm.nih.gov with any urgent questions or concerns.

Thank you!

3/10/2020 5:47 AM Request to delete two SRA terms of from my submission Incoming Received

From xiaokpcc@163.com
To sra@ncbi.nlm.nih.gov;sra@ncbi.nlm.nih.gov;

Attachment #

Dear administrator:

I uploaded two unrelated files in my Bioproject, can you help me to delete them?

The BioProject ID is PRJNA607174.

The submission ID is SUB6988522.

The BioSample accession of two objects need to deletet: SAMN14126327 and SAMN14126328.

Please help me to delete these two BioSample (SAMN14126327 and SAMN14126328) and their related SRA files.

Thank you very much.

Yours

Kangpeng Xiao

20200310

3/10/2020 10:49 AM Re: case #CAS-503400-N7S9S2: Request to delete two SRA terms of from my submi... TRACKING:000393000005514 Outgoing Sent

From nlm-support@nlm.nih.gov
To xiaokpcc@163.com;

Attachment #

I have withdrawn SAMN14126327 and SAMN14126328, along with the associated SRA data.

Cheers,

Rick Lapoint

SRA Curator

* PLEASE DO NOT MODIFY THE SUBJECT LINE OF THIS EMAIL WHEN RESPONDING TO ENSURE CORRECT TRACKING *

Case Information:

Case #: CAS-503400-N7S9S2

Customer Name: 肖康鹏

Customer Email: xiaokpcc@163.com

Case Created: 2020-03-10T09:47:39Z

Summary: Request to delete two SRA terms of from my submission

Details:

Dear administrator:

I uploaded two unrelated files in my Bioproject, can you help me to delete them?

The BioProject ID is PRJNA607174.

The submission ID is SUB6988522.

The BioSample accession of two objects need to deletet: SAMN14126327 and SAMN14126328.

Please help me to delete these two BioSample (SAMN14126327 and SAMN14126328) and their related SRA files.

Thank you very much.

Yours

Kangpeng Xiao
20200310

3/14/2020 8:57 AM Re:Re: case #CAS-503400-N7S9S2: Request to delete two SRA terms of from my submi... TRACKING:000393000005514 Incoming Received

From xiaokpcc@163.com

To nlm-support@nlm.nih.gov;nlm-support@nlm.nih.gov;nlm-support@nlm.nih.gov;

Attachment #

Dear Curator:

I can see the two withdrew data from the reviewer link of PRJNA607174.

Will these withdrew data be downloaded by others when the project is released?

Thank you.

At 2020-03-10 22:49:54, "NLM Support" <nlm-support@nlm.nih.gov> wrote:

I have withdrawn SAMN14126327 and SAMN14126328, along with the associated SRA data.

Cheers,

Rick Lapoint
SRA Curator

* PLEASE DO NOT MODIFY THE SUBJECT LINE OF THIS EMAIL WHEN RESPONDING TO ENSURE CORRECT TRACKING *

Case Information:

Case #: CAS-503400-N7S9S2

Customer Name: 肖康鹏

Customer Email: xiaokpcc@163.com

Case Created: 2020-03-10T09:47:39Z

Summary: Request to delete two SRA terms of from my submission

Details:

Dear administrator:

I uploaded two unrelated files in my Bioproject, can you help me to delete them?

The BioProject ID is PRJNA607174.

The submission ID is SUB6988522.

The BioSample accession of two objects need to deletet: SAMN14126327 and SAMN14126328.

Please help me to delete these two BioSample (SAMN14126327 and SAMN14126328) and their related SRA files.

Thank you very much.

Yours
Kangpeng Xiao
20200310

3/16/2020 5:49 PM Re:Re: case #CAS-503400-N7S9S2: Request to delete two SRA terms of from my submi... TRACKING:000393000005514 Outgoing Sent

From nlm-support@nlm.nih.gov
To xiaokpcc@163.com;
Attachment #

No, SAMN14126327 and SAMN14126328, and the associated SRA data will not be available for download when the rest of the submission is released.

Cheers,
Rick Lapoint
SRA Curator

----- Original Message -----

From: 肖康鹏;
Received: Sat Mar 14 2020 08:55:55 GMT-0400 (Eastern Daylight Time)
To: nlm-support@nlm.nih.gov; Inbound - NLM Support; Triage Team;
Subject: Re:Re: case #CAS-503400-N7S9S2: Request to delete two SRA terms of from my submi... TRACKING:000393000005514

Dear Curator:
I can see the two withdrew data from the reviewer link of PRJNA607174.
Will these withdrew data be downloaded by others when the project is released?
Thank you.

At 2020-03-10 22:49:54, "NLM Support" <nlm-support@nlm.nih.gov> wrote:

I have withdrawn SAMN14126327 and SAMN14126328, along with the associated SRA data.
Cheers,
Rick Lapoint
SRA Curator

* PLEASE DO NOT MODIFY THE SUBJECT LINE OF THIS EMAIL WHEN RESPONDING TO ENSURE CORRECT TRACKING *

Case Information:
Case #: CAS-503400-N7S9S2

Customer Name: 肖康鹏

Customer Email: xiaokpcc@163.com

Case Created: 2020-03-10T09:47:39Z

Summary: Request to delete two SRA terms of from my submission

Details:

Dear administrator:

I uploaded two unrelated files in my Bioproject, can you help me to delete them?

The BioProject ID is PRJNA607174.

The submission ID is SUB6988522.

The BioSample accession of two objects need to deletet: SAMN14126327 and SAMN14126328.

Please help me to delete these two BioSample (SAMN14126327 and SAMN14126328) and their related SRA files.

Thank you very much.

Yours

Kangpeng Xiao

20200310

6/19/2020 7:39
PM

Request to release data

Incoming Received

From

xiaokpcc@163.com

To

bioprojecthelp@ncbi.nlm.nih.gov;

Attachment #

Dear Curator:

I have uploaded a new sequence data under bioproject PRJNA607174, the new SRA accession ID is SRR12053850.

Please help me release this new sequence data (SRR12053850) as soon as possible.

Thank you very much.

Kangpeng Xiao

6/22/2020 12:15
PM

Re: case #CAS-557677-B6K7R5: Request to release data
TRACKING:000414000006953

Outgoing Sent

From

nlm-support@nlm.nih.gov

To

xiaokpcc@163.com;

Attachment #

Dear Kangpeng ,

I released this run. Please allow 24-48 hrs for propagation of the data to the public site.

For making updates please refer to the Update
guide: <https://www.ncbi.nlm.nih.gov/sra/docs/submitupdate/>

Best regards,

Svetlana

If you have any questions or concerns regarding your **SRA** submission please don't hesitate to contact sra@ncbi.nlm.nih.gov (applies to new questions). We *normally* respond *within 2 business days*.

11/8/2020 4:29 AM Request update data Incoming Received

From xiaokpcc@163.com
To sra@ncbi.nlm.nih.gov;sra@ncbi.nlm.nih.gov;
Attachment #

Dear Curator:

I submitted the amplicon sequencing data to SRA on October 30 (SRA accession no. SRR12950102) under the Bioproject PRJNA607174. Due to negligence, an amplicon sequence was missing from this data, the correct file should contain 33 sequences, but only 32 sequences were included in the submitted file. Can I resubmit a file to overwrite the original data? This is very important to us and we look forward to hearing from you.

Thank you very much.

11/9/2020 9:32 AM Re: case #CAS-630560-S5D6W3: Request update data TRACKING:000414000008360 Outgoing Sent

From nlm-support@nlm.nih.gov
To xiaokpcc@163.com;
Attachment #

Hello,

I had withdrawn this SRA records and you can submit a new one by making new SRA submission and using existing bioproject's and sample's accessions for SRA metadata.

Please note, however, that judging by your description there are too few sequences in your file to qualify for SRA submission. SRA fastq files usually contain thousands or millions of sequences.

Please refer to the File Format guide: <https://www.ncbi.nlm.nih.gov/sra/docs/submitformats/>

Best regards,

Svetlana

If you have any questions or concerns regarding your **SRA** submission please don't hesitate to contact sra@ncbi.nlm.nih.gov (applies to new questions). We *normally* respond *within 2 business days*.

Svetlana lazvovskaia
The NCBI SRA database submission staff

=====

Dear Curator:

I submitted the amplicon sequencing data to SRA on October 30 (SRA accession no. SRR12950102) under the Bioproject PRJNA607174. Due to negligence, an amplicon sequence was missing from this data, the correct file should contain 33 sequences, but only 32 sequences were included in the submitted file. Can I resubmit a file to overwrite the original data? This is very important to us and we look forward to hearing from you. Thank you very much.

11/9/2020 9:51 AM Re:Re: case #CAS-630560-S5D6W3: Request update data TRACKING:000414000008360 Incoming Received
From xiaokpcc@163.com
To nlm-support@nlm.nih.gov;nlm-support@nlm.nih.gov;nlm-support@nlm.nih.gov;
Attachment #

Dear Curator:

Please do not withdraw my original data for the time being (SRA accession no. SRR12950102) . Our data is under editor review. We will update the data after the review results come out. I am very sorry about my previous request was not clearly stated, can you help me release the withdrew original data as soon as possible ? Thank you very much.

At 2020-11-09 22:32:18, "NLM Support" <nlm-support@nlm.nih.gov> wrote:

Hello,

I had withdrawn this SRA records and you can submit a new one by making new SRA submission and using existing bioproject's and sample's accessions for SRA metadata.

Please note, however, that judging by your description there are too few sequences in your file to qualify for SRA submission. SRA fastq files usually contain thousands or millions of sequences.

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Best regards,

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Svetlana lazvovskaia
The NCBI SRA database submission staff

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Thank you very much.

11/9/2020 9:54 AM Re:Re: case #CAS-630560-S5D6W3: Request update data TRACKING:000414000008360 Outgoing Sent
From nlm-support@nlm.nih.gov
To xiaokpcc@163.com;
Attachment #

I reinstated this run.

Best regards,

Svetlana

If you have any questions or concerns regarding your **SRA** submission please don't hesitate to contact sra@ncbi.nlm.nih.gov (applies to new questions). We *normally* respond *within 2 business days*.

Svetlana lazvovskaia
The NCBI SRA database submission staff

----- Original Message -----

From: 肖康鹏;

Received: Mon Nov 09 2020 09:51:06 GMT-0500 (Eastern Standard Time)

To: nlm-support@nlm.nih.gov; Inbound - NLM Support; Triage Team;

Subject: Re:Re: case #CAS-630560-S5D6W3: Request update data TRACKING:000414000008360

Dear Curator:

Please do not withdraw my original data for the time being (SRA accession no. SRR12950102) . Our data is under editor review. We will update the data after the review results come out. I am very sorry about my previous request was not clearly stated, can you help me release the withdrew original data as soon as possible ? Thank you very much.

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Please note, however, that judging by your description there are too few sequences in your file to qualify for SRA submission. SRA fastq files usually contain thousands or millions of sequences.

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Svetlana

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Svetlana lazvovskaia

The NCBI SRA database submission staff

=====

Dear Curator:

I submitted the amplicon sequencing data to SRA on October 30 (SRA accession no. SRR12950102) under the Bioproject PRJNA607174. Due to negligence, an amplicon sequence was missing from this data, the correct file should contain 33 sequences, but only 32 sequences were included in the submitted file. Can I resubmit a file to overwrite the original data? This is very important to us and we look forward to hearing from you. Thank you very much.

11/13/2020 9:53 AM

Re:Error in SRA submission SUB8543863

Incoming Received

From xiaokpcc@163.com
To sra@ncbi.nlm.nih.gov;sra@ncbi.nlm.nih.gov;
Attachment #

Dear Curator:

I uploaded two files (SUB8543863) to replace the previously withdrawn SRA data (SRA accession no. SRR12950102). One file is fastq format (converted from sanger sequencing data), and the other one is a tar ball file include all the raw amplicon chromatograms data.

I previously received enthusiastic NCBI staff reply that '*I recommend to upload fastq file with Sanger sequences and additionally tar ball with ab1 files. We'll load it provisionally for now and when ab1 loader will be available we will load them in a standard way.*'

Please help us reload SRR12950102 with this two files.

Thank you very much!

Best wishes.

At 2020-11-13 22:10:04, sra@ncbi.nlm.nih.gov wrote:

>Dear Kangpeng Xiao,

>

>Your submission SUB8543863 could not be processed due to an error(s).

>To review your submissions and see the cause of the error, please log in to the NCBI Submission Portal at <https://submit.ncbi.nlm.nih.gov/subs/>

>Please contact SRA at sra@ncbi.nlm.nih.gov for guidance in fixing your submission. You will not receive an accession number until the submission is complete.

>

>Regards,

>

>NCBI SRA Submissions Staff

>Bethesda, Maryland USA

11/17/2020 8:58 AM

Error in SRA submission SUB8543863

Incoming Received

From xiaokpcc@163.com
To sra@ncbi.nlm.nih.gov;sra@ncbi.nlm.nih.gov;

Attachment #

Dear Curator:

Since the journal editor wanted to release raw Sanger sequences, we submitted the fastq file of the amplified sequence and the original ab1 file. We very much hope that these data will be released as soon as possible. Could you please tell us what we need to do further? Do I need to resubmit the sequence?

Thank you very much, looking forward to your reply!

Kangpeng Xiao

At 2020-11-13 22:51:18, "Kangpeng Xiao" <xiaokpcc@163.com> wrote:

Dear Curator:

I uploaded two files (SUB8543863) to replace the previously withdrawn SRA data (SRA accession no. SRR12950102). One file is fastq format (converted from sanger sequencing data), and the other one is a tar ball file include all the raw amplicon chromatograms data.

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>

>Regards,

>

>NCBI SRA Submissions Staff

>Bethesda, Maryland USA

11/17/2020 2:47 PM case #CAS-633322-C2V3B7: RE: Error in SRA submission SUB8543863 TRACKING:000422000003768 Outgoing Sent

From nlm-support@nlm.nih.gov

To xiaokpcc@163.com;

Attachment #

Dear Kangpeng Xiao,

Files have been processed and released, they should be available for download soon.

Please use BioProject accession PRJNA607174 to reference whole set in your publication and update it with publication details when you'll have them:

<https://dataview.ncbi.nlm.nih.gov/object/PRJNA607174>

Best,

Inna

==

Inna Belaia
SRA curator

----- Original Message -----

From: 肖康鹏;

Received: Tue Nov 17 2020 08:55:31 GMT-0500 (Eastern Standard Time)

To: NLM/NCBI List sra; SRA Support;

Subject: Error in SRA submission SUB8543863

Dear Curator:

Since the journal editor wanted to release raw Sanger sequences, we submitted the fastq file of the amplified sequence and the original ab1 file. We very much hope that these data will be released as soon as possible. Could you please tell us what we need to do further? Do I need to resubmit the sequence?

Thank you very much, looking forward to your reply!

Kangpeng Xiao

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>Regards,

>

>NCBI SRA Submissions Staff

>Bethesda, Maryland USA

bioprojecthelp at ncbi.nlm.nih.gov bioprojecthelp at ncbi.nlm.nih.gov
Mon Aug 10 08:22:17 EDT 2020

Dear Weifeng Shi,

This is an automatic acknowledgment that your submission:

SubmissionID: SUB7897952
BioProject ID: PRJNA656060
Title:

has been successfully registered with the BioProject database. After review by the database staff, your project information will be accessible with the following link, usually within a few days of the release date that you set (or the release of linked data, whichever is first):

<http://www.ncbi.nlm.nih.gov/bioproject/656060>

Please use the BioProject ID PRJNA656060 with your correspondence and your data submissions.

Send questions to bioprojecthelp at ncbi.nlm.nih.gov, and include the BioProject ID and organism name.

Regards,

NCBI BioProject Submissions Staff
Bethesda, Maryland USA

(301) 496-2475

(301) 480-2918 (Fax)

bioprojecthelp at ncbi.nlm.nih.gov (for BioProject questions/replies)

info at ncbi.nlm.nih.gov (for general questions regarding NCBI)
