

From: gb-admin@ncbi.nlm.nih.gov
Date: Wed, 14 Aug 2019 12:16:28 -0400 (Eastern Daylight Time)
To: huben@wh.iov.cn
Subject: GenBank accession MN312240-MN312869
X-Mailer: Smart
MIME-Version: 1.0
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Dear GenBank Submitter:

Please find your assigned Accession Number(s) in the letter below.

Based on your reply and/or submission, some of the sequences may be trimmed at the ends to resolve sequence issues or the following comment may be added to one or more of your record(s):

COMMENT GenBank staff is unable to verify sequence and/or annotation provided by the submitter.

Sequences flagged as unverified will not be included in NCBI BLAST databases.

If you wish to revise these records, please see:

<https://www.ncbi.nlm.nih.gov/genbank/update.html>

for proper update formats and send the updates to:
gb-admin@ncbi.nlm.nih.gov

Sincerely,

The GenBank Submissions Staff
Bethesda, Maryland USA

Dear GenBank Submitter:

Thank you for your direct submission of sequence data to GenBank. We have provided GenBank accession numbers for your nucleotide sequences:

The full list of accession numbers is attached. Please review prior to publishing your accession numbers as submissions may have been rearranged during processing.

file Alpha_CoV_batch_1.sqn:
MN312240 - MN312288

file Alpha_CoV_batch_2.sqn:

MN312289 - MN312362

file Alpha_CoV_batch_3.sqn:
MN312363 - MN312396

file Alpha_CoV_batch_4.sqn:
MN312397 - MN312453

file Alpha_CoV_batch_5.sqn:
MN312454 - MN312537

file Alpha_CoV_batch_6.sqn:
MN312538 - MN312594

file Beta_CoV_batch_1.sqn:
MN312595 - MN312649

file Beta_CoV_batch_2.sqn:
MN312650 - MN312690

file Beta_CoV_batch_3.sqn:
MN312691 - MN312748

file Beta_CoV_batch_4.sqn:
MN312749 - MN312828

file Beta_CoV_batch_5.sqn:
MN312829 - MN312869

The GenBank accession numbers should appear in any publication that reports or discusses these data, as it gives the community a unique label with which they may retrieve your data from our on-line servers. You may prepare and submit your manuscript before your accessions are released in GenBank.

Submissions are not automatically deposited into GenBank after being accessioned. Each sequence record is individually examined and processed by the GenBank annotation staff to ensure that it is free of errors or problems.

You have requested that your data are to be held confidential until:

Dec 31, 2020

They will not be released to the public database until this date, or until the data or accession numbers appear in print, whichever is first.

Since the flatfile record is a display format only and is not an editable format of the data, do not make changes directly to a flatfile. For complete information about different methods to update a sequence record, see: <https://www.ncbi.nlm.nih.gov/Genbank/update.html>

Any inquiries about your submission should be sent to gb-admin@ncbi.nlm.nih.gov

For more information about the submission process or the available submission tools, please contact GenBank User Support at info@ncbi.nlm.nih.gov.

Please reply using the current Subject line.

Sincerely,

Beverly Underwood
Contractor

The GenBank Direct Submission Staff
Bethesda, Maryland USA

gb-admin@ncbi.nlm.nih.gov (for updates/replies to GenBank entries)

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

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