

From: James Floyd jfloyd@uw.edu
Subject: Fwd: Record update from chs-nhlbi.org
Date: October 12, 2020 at 3:50 PM
To: Guillaume Lettre Guillaume.Lettre@mhi-humangenetics.org



Dear Guillaume,

CHS has approved the paper below.

Best,

James

Begin forwarded message:

From: chsntlbi@u.washington.edu
Subject: Record update from chs-nhlbi.org
Date: October 12, 2020 at 12:41:08 PM PDT
To: jfloyd@uw.edu
Reply-To: chsntlbi@u.washington.edu

Hello, authors.

Congratulations! Your penultimate draft, find-tfbs: a tool to identify functional non-coding variants associated with complex human traits using open chromatin maps and phased whole-genome sequences, was approved by the Steering Committee. Please see the updated status and details regarding your next steps at the following link: <https://chs-nhlbi.org/node/8435>.

Please let me know if you have any questions.

Best,
Ellen

Ellen Terry
Publications Coordinator
[CHS-NHLBI.org](https://chs-nhlbi.org)
chsntlbi@u.washington.edu

NOTES: Please make sure you are logged on to the CHS website before clicking the above link. This message was sent to the first author and all CHS co-authors.

