


From: Guillaume Lettre guillaume.lettre@mhi-humangenetics.org 

Subject: find-tfbs manuscript

Date: October 2, 2020 at 11:40 AM

To: Alex Reiner apreiner@u.washington.edu, Thibord, Florian (NIH/NHLBI) [F] florian.thibord@nih.gov, Paul William Livermore Auer pauer@uwm.edu, James Floyd jfloyd@uw.edu, RASIKA Mathias rmathias@jhmi.edu, Lewis, Joshua jlewis2@som.umaryland.edu, Coban Akdemir, Zeynep H Zeynep.H.CobanAkdemir@uth.tmc.edu, Myriam.Fornage@uth.tmc.edu, john.blangero@utrgv.edu, North, Kari E kari_north@unc.edu, LESLIE.LANGE@cuanschutz.edu LESLIE.LANGE@CUANSCHUTZ.EDU, Matt Conomos mconomos@uw.edu

Cc: sebastian.meric.de.bellefon@umontreal.ca, Sébastien Méric de Bellefon arnaudpourseb@gmail.com

GL

Dear co-authors,

Sébastien and I have finalized the find-tfbs manuscript. Thanks to all for agreeing to be author and sending your comments and acknowledgments. You can use the attached version for your study level P&P review/approval.

@Alex, to submit to TOPMed P&P, I need a letter (I guess email is fine) from you as chair of the H&H working group.

@Matt, I need guidance from you for one last point. On the TOPMed manuscript submission webpage, I am asked to attach: "all necessary study-specific approval documentation". What does that mean exactly? Do I need to wait until I hear back from each study before submitting to TOPMed? Or is this email sufficient?

Thanks,
Guillaume



findtfbs_2Oct20
20_forPandP.pdf



findtfbs_v2_supT
ables_...20.xlsx

Parent Study short name	Person contacted on October 2nd 2020	Response
Amish	Joshua Lewis	Waiting
ARIC	Zeynep Coban-Akdemir	Waiting
CARDIA	Myriam Fornage	Approved
CHS	James Floyd	Approved
FHS	Florian Thibord	Waiting
GeneSTAR	R. Mathias	Approved
HCHS_SOL	Jee-Young Moon	Waiting
JHS	A. Reiner	Approved (A. Reiner, convener)
MESA	L. Lange	Waiting
SAFS	John Blangero	Waiting
WHI	Paul Auer	Waiting

From: Fornage, Myriam Myriam.Fornage@uth.tmc.edu
Subject: Re: find-tfbs manuscript
Date: October 8, 2020 at 2:42 PM
To: Guillaume Lettre Guillaume.Lettre@mhi-humangenetics.org

MF

Hi Guillaume:
You have the green light from CARDIA.
Best wishes,
Myriam

From: Guillaume Lettre <Guillaume.Lettre@mhi-humangenetics.org>
Sent: Thursday, October 8, 2020 1:34 PM
To: Alex Reiner; Thibord, Florian (NIH/NHLBI) [F]; Paul William Livermore Auer; James Floyd; RASIKA Mathias; Lewis, Joshua; Coban Akdemir, Zeynep H; Fornage, Myriam; john.blangero@utrgv.edu; North, Kari E; LESLIE.LANGE@cuanschultz.edu; Matt Conomos
Cc: sebastian.meric.de.bellefon@umontreal.ca; Sébastien Méric de Bellefon
Subject: Re: find-tfbs manuscript

**** **EXTERNAL EMAIL** ****

Hi everybody,

I just wanted to provide an update about the status of this manuscript. I appreciate your patience with me as I am learning how to navigate the TOPMed process.

I sent this email on October 2nd, asking if you could submit to your study P&P. If you have heard back already and gotten approval, send me the email. If you have not submitted the manuscript for review, please do it now. As per the TOPMed rules, I can send the paper for TOPMed review on October 16 (2 weeks after my original request) even if I haven't heard back, but it would be great to get all approvals in beforehand.

Thanks,
Guillaume

On Oct 2, 2020, at 11:40 AM, Guillaume Lettre <Guillaume.Lettre@mhi-humangenetics.org> wrote:

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Thanks,
Guillaume

<findtfbs_2Oct2020_forPandP.pdf>

<findtfbs_v2_supTables_22Sept2020.xlsx>

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: chsnhlbi@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: chsnhlbi@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)
chsnhlbi@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.

From: RASIKA Mathias rmathias@jhmi.edu
Subject: Re: find-ifbs manuscript
Date: October 9, 2020 at 10:25 AM
To: Guillaume Lettre Guillaume.Lettre@mhi-humangenetics.org
Cc: sebastian.meric.de.bellefon@umontreal.ca, Sébastien Méric de Bellefon arnaudpourseb@gmail.com

RM

Guillaume,

Here are the acknowledgements that need to go in for GeneSTAR:

GeneSTAR was supported by the National Institutes of Health/National Heart, Lung, and Blood Institute (U01 HL72518, HL087698, HL112064) and by a grant from the National Institutes of Health/National Center for Research Resources (M01-RR000052) to the Johns Hopkins General Clinical Research Center.

For sequencing center, based on the N's in Table 1, it looks like only MacroGen and Illumina data were included. This seems very odd to me, if using freeze 8, why not use the Broad data too? Did you exclude some Genestar samples?

If only MacroGen and Illumina, then sequencing center text needs to be changed to

"WGS for "NHLBI TOPMed: GeneSTAR" (phs001218) was performed at Psomagen (3R01HL112064-04S1) and Illumina (HL112064)."

If Broad too, then should say

"WGS for "NHLBI TOPMed: GeneSTAR" (phs001218) was performed at Psomagen (3R01HL112064-04S1), Illumina (HL112064), and the Broad Institute of MIT and Harvard (HHSN268201500014C)"

You have the green light on submission from GeneSTAR.

Rasika

Rasika Mathias, ScD.

Associate Professor of Medicine,
Johns Hopkins University.

Director of Genomics, GeneSTAR Research program, General Internal Medicine.

Director, Data Analysis Unit, Allergy and Clinical Immunology.

5501 Hopkins Bayview Circle, 3A.62A

Baltimore, MD 21224

Tel: 410-550-2487

From: Guillaume Lettre <Guillaume.Lettre@mhi-humangenetics.org>

Sent: Friday, October 2, 2020 11:40 AM

To: Alex Reiner <apreiner@u.washington.edu>; Thibord, Florian (NIH/NHLBI) [F] <florian.thibord@nih.gov>; Paul William Livermore Auer <pauer@uwm.edu>; James Floyd <jfloyd@uw.edu>; RASIKA Mathias <rmathias@jhmi.edu>; Lewis, Joshua

<jlewis2@som.umaryland.edu>; Coban Akdemir, Zeynep H
<Zeynep.H.CobanAkdemir@uth.tmc.edu>; Myriam.Fornage@uth.tmc.edu
<Myriam.Fornage@uth.tmc.edu>; john.blangero@utrgv.edu <john.blangero@utrgv.edu>;
North, Kari E <kari_north@unc.edu>; LESLIE.LANGE@cuanschultz.edu
<LESLIE.LANGE@CUANSCHUTZ.EDU>; Matt Conomos <mconomos@uw.edu>
Cc: sebastian.meric.de.bellefon@umontreal.ca
<sebastian.meric.de.bellefon@umontreal.ca>; Sébastien Méric de Bellefon
<arnaudpourseb@gmail.com>
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