

Raffield, Laura

From: Raffield, Laura
Sent: Thursday, June 29, 2023 12:18 PM
To: Heard-Costa, Nancy L; Ann Walsh
Cc: Mutalik, Karen M; Jiang, Minzhi
Subject: RE: TOPMed inflammation paper draft for review- proposal # 1854 Whole Genome Sequencing Based Analysis of Inflammation Biomarkers

Wonderful thank you Nancy!

From: Heard-Costa, Nancy L <nheard@bu.edu>
Sent: Thursday, June 29, 2023 11:49 AM
To: Raffield, Laura <laura_raffield@unc.edu>; Ann Walsh <walshac@nhlbi.nih.gov>
Cc: Mutalik, Karen M <kmutalik@bu.edu>; Jiang, Minzhi <minzhi@live.unc.edu>
Subject: RE: TOPMed inflammation paper draft for review- proposal # 1854 Whole Genome Sequencing Based Analysis of Inflammation Biomarkers

Hi Laura,

The proposal looks good and I see that FHS related personnel including Emelia Benjamin and Honghuang Lin are listed as co-authors, so there is no further action needed at this time.

Nancy

From: Raffield, Laura <laura_raffield@unc.edu>
Sent: Thursday, June 29, 2023 10:51 AM
To: Ann Walsh <walshac@nhlbi.nih.gov>; Heard-Costa, Nancy L <nheard@bu.edu>
Cc: Mutalik, Karen M <kmutalik@bu.edu>; Jiang, Minzhi <minzhi@live.unc.edu>
Subject: RE: TOPMed inflammation paper draft for review- proposal # 1854 Whole Genome Sequencing Based Analysis of Inflammation Biomarkers

Hi Nancy! Just wanted to confirm nothing more we need to do for FHS before submitting to TOPMed portal.
Thanks so much,
Laura

From: Raffield, Laura
Sent: Tuesday, June 20, 2023 3:58 PM
To: Walsh, Ann (NIH/NHLBI) [E] <walshac@nhlbi.nih.gov>; Nancy Heard-Costa <nheard@bu.edu>
Cc: kmutalik@bu.edu
Subject: RE: TOPMed inflammation paper draft for review- proposal # 1854 Whole Genome Sequencing Based Analysis of Inflammation Biomarkers

Hi Ann! Yes, this one had non NHLBI FHS representatives...I will certainly submit to TOPMed portal once I have heard back from and incorporated coauthor comments and gotten study specific approvals.

Thanks everyone!
Laura

draft of our freeze 8 findings. We hope to submit to Human Molecular Genetics (alternate journal suggestions welcome as well).

We welcome your comments on how to best present our findings and improve the paper, and any needed edits, but would appreciate circulation of the paper to relevant cohort P&Ps at this time. Please double check your name and affiliation and cohort acknowledgments. Let Paul, Sheila, Min-Zhi and I know any questions you may have!

Best wishes,
Laura

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