Hi Sarah,

I cc'd you on this since I'm not sure how to proceed. I contacted FHS over a month ago for study-specific review of the phenotype harmonization manuscript and have heard *nothing* from Ann Walsh. I think I need their approval to submit the manuscript on the TOPMed website, right? Can I submit to the website if I've contacted them numerous times and gotten no response? Do you have any suggestions about what I can or should do?

Best,

Adrienne

On May 4, 2020, at 10:45 AM, Adrienne M Stilp <amstilp@uw.edu> wrote:

Hi Ann and Karen,

Do you have any updates about FHS review of this TOPMed manuscript? This is the one requirement that is holding up this manuscript - we've heard back from all the other studies, but can't submit until we get approval from FHS.

Thanks, Adrienne

On Apr 24, 2020, at 11:41 AM, Mutalik, Karen M <kmutalik@bu.edu> wrote:

Hi Adrienne and Ann,

Adrienne: I will need to defer to Ann Walsh. She would be able to answer the question. Ann: Should this be forwarded to any other BU person?

Karen

-----Original Message-----From: Adrienne M Stilp <amstilp@uw.edu> Sent: Friday, April 24, 2020 12:31 PM To: Ann Walsh <walshac@nhlbi.nih.gov>; Mutalik, Karen M <kmutalik@bu.edu> Cc: Cathy C. Laurie <cclaurie@uw.edu> Subject: Re: FHS review request for TOPMed manuscript

Hi Ann and Karen,

I still haven't heard from you about the TOPMed DCC phenotype harmonization manuscript. Has it been reviewed by FHS yet?

Thanks, Adrienne

On Apr 13, 2020, at 9:29 AM, Adrienne M Stilp <amstilp@uw.edu> wrote:

Hi Ann and Karen,

I haven't heard from you about this manuscript, so I wanted to check in about its status. Can you let me know if it's being reviewed by FHS, or if I need to provide additional information?

Best, Adrienne

On Apr 1, 2020, at 9:35 AM, Adrienne M Stilp <amstilp@uw.edu> wrote:

Dear Ann Walsh and Karen Mutalik,

I am submitting the following TOPMed manuscript for review by FHS. All co-authors have reviewed the manuscript and approve of it. The paper details the process that the TOPMed DCC used to harmonize phenotypes across 17 different TOPMed studies, one of which was FHS. Please let me know when the manuscript has been approved, or if any changes are requested.

AS

Best, Adrienne Stilp

Adrienne Stilp, PhD Research Scientist TOPMed DCC/Genetic Analysis Center Department of Biostatistics University of Washington

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