

From: [Gccbiost](#) on behalf of [Vladimir Seplyarskiy](#)
To: [Genetic Analysis Center](#)
Subject: [Gccbiost] Fwd: Review TOPMED manuscript
Date: Wednesday, December 4, 2019 12:11:20 PM
Attachments: [Main Text Germline processes last.eb.docx](#)
[ATT00001.txt](#)

Dear DCC,

here the approval from Eric, who represents ARIC. Fro manuscript **Population sequencing data reveal a compendium of mutational processes in human germline.**

----- Forwarded message -----

От: **Boerwinkle, Eric** <Eric.Boerwinkle@uth.tmc.edu>
Date: вт, 29 окт. 2019 г. в 11:42
Subject: RE: Review TOPMED manuscript
To: alicodendrochit@gmail.com <alicodendrochit@gmail.com>

1. Comments attached. Good job, and thanks for including me.
2. My over-arching comment is that it might be seen as too specialized for Science.

Eric.

From: Vladimir Seplyarskiy <alicodendrochit@gmail.com>
Sent: Friday, October 25, 2019 11:08 AM
Subject: Review TOPMED manuscript

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

Dear TOPMed contributor and co-author,

we are happy to present our manuscript entitled "**Population sequencing data reveal a compendium of mutational processes in human germline**"

and linked proposal "Approach to determine germline epigenetic state and mutators alleles based on local mutation rate" for your review.

<https://drive.google.com/drive/folders/1GleyecEOG3huHmjbT6EE31zzbh05QmWp?ths=true>

This study uses SNPs from TOPMed freeze 5 genotype data ,

We are looking forward to apply you comments and submit manuscript to Science.

Best regards,

Vladimir Seplyarskiy, PhD