

Epigenetic Clock GWAS draft MS

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 1 attachments (3 MB)

McCartney_et_al.zip;

Dear all,

Thank you again for your help with our GWAS meta-analysis of epigenetic ageing biomarkers. We are delighted to share the first draft of the paper with you!

There are a couple of minor analyses that remain ongoing: polygenic predictions into 4 additional cohorts along with colocalisation analyses for mQTL/eQTL SNPs. However, the main message from the paper is not going to change. We are also waiting for cohort summary and descriptive data from a few studies.

We are aiming to submit the manuscript by the first week of March so if you are able to get comments back to us by Friday 28th Feb that would be much appreciated. Also, given the number of authors on the paper, please could the cohort analysts collate comments from their respective team members to minimise the number of files we have to cross reference with our master copy.

Many thanks and best wishes,

Daniel and Riccardo

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