# **RE: Cortical Thickness Update**

From: Rachael Garner | Rachael.Garner@loni.usc.edu

Thursday, Jun 3, 8:29 AM

To: Riikka Immonen | riikka.immonen@uef.fi

Cc: Ryan Cabeen | Ryan.Cabeen@loni.usc.edu, Aubrey Martinez | aubreylmartinez@gmail.com, Sweta Bhagavatula | sbhagava@usc.edu

Hi Riikka,

We spent the last month testing the cortical thickness pipeline on Project 2 imaging modalities and wanted to share an update. As the pipeline was optimized for MT data, we found that MGE or MGE derived images did not produce reliable cortical thickness estimation outputs (including t2star, r2star and various extractions over the 13 echoes including base/biased/mean images produced by Ryan). Given this, we believe that it makes most sense to move forward with submitting a manuscript for the pipeline as is, and we can concurrently work on reoptimizing the pipeline for MGE data (e.g. the denoised MGE) for a second manuscript.

We plan to work on an MT manuscript draft, using your gold standard thickness measurements as validation, and can send to you to review for co-authorship. Let us know how this sounds – we are happy to set up a call to walk through any of the outputs or discuss next steps with the manuscript. Thank you again for all your guidance and collaboration.

Warm regards,

#### **Rachael Garner**

**Project Specialist** 

Laboratory of Neuro Imaging

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Email: Rachael.Garner@loni.usc.edu

From: Riikka Immonen | riikka.immonen@uef.fi

Friday, Jun 4, 3:52 AM

To: Rachael Garner | Rachael.Garner@loni.usc.edu

Cc: Ryan Cabeen | Ryan.Cabeen@loni.usc.edu, Aubrey Martinez | aubreylmartinez@gmail.com, Sweta Bhagavatula | sbhagava@usc.edu

Hi Racheal,

#### Thanks for the update!

Excellent that you have done the testing with other modalities – did you get it to work on ANY cortical areas / how

far off are we there? (no need to go through all details – if you have a pic of a representative case I could eye-ball it to get the idea) Anyhow, now you have the answers for reviewers and EpiBios crew.

Yes, I agree with the plan – go ahead with the manuscript, and I'll review for my part. If you plan to keep it as methodology paper and (only) refer to the biomarker aspect/potential & explain the EpiBios context, rather than using it to epileptic / non-epileptic grouping, then there is no need to more extensive discussions beforehand. But if you wish to claim about some predictive value in posttraumatic epilepsy (I.e. use the epilepsy outcome classifications), then we need to arrange a wider discussion including Asla Pitkänen and Olli Gröhn.

Do you actually need still some 2days postTBI thickness measures from me? (I recall not to have completed those..?) Let me know if you do!

Best Regards, Riikka

### Riikka Immonen, PhD

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From: Rachael Garner | Rachael.Garner@loni.usc.edu To: Riikka Immonen | riikka.immonen@uef.fi Thursday, Jun 3, 6:29 PM Hi Riikka,

We spent the last month testing the cortical thickness pipeline on Project 2 imaging modalities and wanted to share an update. As the pipeline was optimized for MT data, we found that MGE or MGE derived images did not produce reliable cortical thickness estimation outputs (including t2star, r2star and various extractions over the 13 echoes including base/biased/mean images produced by Ryan). Given this, we believe that it makes most sense to move forward with submitting a manuscript for the pipeline as is, and we can concurrently work on reoptimizing the pipeline for MGE data (e.g. the denoised MGE) for a second manuscript.

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