# Fwd: COMMENT - Project 1 Abstract Submission

From: **Dominique Duncan** | Dominique.Duncan@loni.usc.edu Tuesday, Jun 15, 11:40 PM

To: **Ryan Cabeen** | Ryan.Cabeen@loni.usc.edu

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> From: **Asla Pitkänen** | asla.pitkanen@uef.fi To: **Akul Sharma** | Akul.Sharma@loni.usc.edu Tuesday, Jun 15, 9:34 AM
>
> Deal Akul,
>
> I got the eMail below from Olli. I have to say, I am shocked.
>
> 1. You used project 1 data without consulting me, Terry and Rick who are the PIs in project 1
> 2. You used UEF data without consulting me and the UEF team
> 3. You assigned animals to PTE vs non-PTE groups without knowledge who is who as you can not know that. Only I currently know the most updated list (that we are about to finalize) and that has been on purpose to avoid these kind of strange publications to be submitted.
> 4. You completely disregard the work of all Project 1 postdocs and researcher not including they as co-authors (if they wanted to be)
> 5. Lastly, as I have no idea what parameters you have put into the algorithm we can not know if your analysis makes any sense
>
> I hope I made myself clear. This can not happen again!
>
> Any data that is prepared from project 1 material MUST go through Project 1 PIs!

I suggest that you withdraw the abstract.

asla

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*If you receive this eMail outside your working hours, you are not expected to respond untill you return to work.*

From: **Akul Sharma** | Akul.Sharma@loni.usc.edu To: **Olli Gröhn** | olli.grohn@uef.fi Tuesday, Jun 15, 12:00 AM

Good afternoon,

After our call last week, we quickly began some preliminary analyses on the Project 1 structural data. We started with a machine learning-based classification of PTE vs. non-PTE TBI using diffusion features at the 4 time points. Specifically, we used Ryan's structural pipeline to extract FA values in 48 white matter bundles. We found that the results were quite interesting: with a sample of 32 UEF rodents, the highest prediction accuracy was 75% (78% specificity, 66% sensitivity) with the 30-day data. We were excited to see these results, so we went ahead and drafted an abstract to AES, adding you all as coauthors. We would like to discuss the data further and detail our methodology on our next call, but we wanted to make sure we submitted an abstract by the deadline last night so we hope this is okay with you. Please find the abstract attached to this email.

Thank you

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