RE: Melbourne MRI data - Project 2

Melbourne, VIC 3004

From	: O'Brien, Terence te.obrien@alfred.org.au	Wednesday, Jun 9, 8:04 PM
	David Wright david.wright@monash.edu, Ryan Cabeen Ryan.Cabeen@loni.usc.edu Pablo Casillas-Espinosa pablo.casillas-espinosa@monash.edu, Idrish Ali idrish.ali@monatolli.grohn@uef.fi, Riikka Immonen riikka.immonen@uef.fi, Neil Harris ngharris@ucla.edu, aristea.galanopoulou@einsteinmed.org, Richard Staba rstaba@mednet.ucla.edu	
Than	nks David and Ryan,	
Сору	ying Aristea and Rick in as well so we are all on the same page.	
Rega	ards,	
Terry		
	om: David Wright I david.wright@monash.edu To: Ryan Cabeen I ryan.cabeen@loni.usc.edu	Thursday, Jun 10, 10:14 AM
	e had a meeting this morning to discuss the epibios project and the MRI data analy ow which data you'd like prioritised to receive from us to progress the analysis?	sis. Can you please let us
Th	anks	
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Ass	ociate Professor David Wright	
Dire	ector of Preclinical Imaging	
Dep	partment of Neuroscience	
Cen	atral Clinical School	
	nash University	
	el 6, Alfred Centre	
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From: David Wright | david.wright@monash.edu

Wednesday, Jun 16, 8:54 PM

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

Cc: Pablo Casillas-Espinosa | pablo.casillas-espinosa@monash.edu, Idrish Ali | idrish.ali@monash.edu, Olli Grohn | olli.grohn@uef.fi, Riikka Immonen | riikka.immonen@uef.fi, Neil Harris | ngharris@ucla.edu, Aristea Galanopoulou | aristea.galanopoulou@einsteinmed.org, Richard Staba | rstaba@mednet.ucla.edu, O'Brien, Terence | te.obrien@alfred.org.au

Hi Ryan,

I've uploaded ~175 datasets from project 2 data to the Loni server. These are full datasets (i.e. include method/acqp/fid/2dseq etc) and include a T2 image you can use to quickly assess lesion severity. They're all compressed as tar.bz2 and have been renamed based on the sequence to hopefully make sorting easier for you.

Best wishes

David

Associate Professor David Wright

Director of Preclinical Imaging

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From: Ryan Cabeen | Ryan.Cabeen@loni.usc.edu

Thursday, Jun 10, 1:04 PM

Hi David,

Thanks for reaching. I'm working on updating the image analysis pipeline to work with project two data, so the main thing is to have a representative sample of cases for testing, e.g. including all the sequences like the multishell diffusion, and also a range of lesion severities. Anything along those lines would be much appreciated.

Cheers,

Ryan

Ryan P. Cabeen, PhD

Chan Zuckerberg Imaging Scientist

Assistant Professor of Research Neurology

Laboratory of Neuro Imaging

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From: Terence I te.obrien@alfred.org.au

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Thanks David and Ryan,

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From: O'Brien, Terence I te.obrien@alfred.org.au

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Thanks David. That is great.

Regards,

Hi Ryan,

Terry

From: David Wright | david.wright@monash.edu To: Ryan Cabeen | ryan.cabeen@loni.usc.edu Thursday, Jun 17, 1:54 PM

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Hi David,

Thanks very much! I received all 175 datasets and pushed one case through to test things out. It's made it through the initial preprocessing steps without issue, so things are looking good so far. I'll work on pushing the rest through and making some visualizations, so we can review them together some time soon.

Cheers, Ryan

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