

## Metagenomic sequencing of CSF samples - Support #138

### 10 February 2022 meeting update and way forward

02/10/2022 04:02 PM - Ephie Geza

<b>Status:</b>	New	<b>Start date:</b>	02/10/2022
<b>Priority:</b>	Normal	<b>Due date:</b>	
<b>Assignee:</b>	Gerrit Botha	<b>% Done:</b>	0%
<b>Category:</b>		<b>Estimated time:</b>	0.00 hour
<b>Target version:</b>		<b>Spent time:</b>	0.00 hour
<b>Description</b>			
Dear Ruan and Gerrit,			
The following are the updates on the project:			
<ul style="list-style-type: none"><li>1. Ruan will share the 20 samples that are currently available (the other 27 are yet to be sequenced) on ilifu.</li><li>2. The Bioinformatics support team will perform the human analysis, that is, preprocessing, differential expression analysis, sequence-level analysis and predictive modelling, where possible.</li></ul>			
Gerrit, can we create a directory on ilifu under /cbio/projects/ for the raw data or we use one of the existing directories in the same directory?			
Kind regards, Ephie			

#### History

##### #1 - 02/11/2022 01:43 PM - Gerrit Botha

Thank you Ephie

The folder /cbio/projects/017 was created for this project. The folder is shared with the necessary users (you, Ruan and me). Please use that.

Regards,  
Gerrit