

## Identification of biomarkers in keloids and folliculitis keloidalis nuchae (FKN) - Support #80

### Renaming all files according to the experiments

05/13/2019 10:58 AM - Jon Ambler

<b>Status:</b>	Resolved	<b>Start date:</b>	05/13/2019
<b>Priority:</b>	High	<b>Due date:</b>	
<b>Assignee:</b>	Jon Ambler	<b>% Done:</b>	100%
<b>Category:</b>		<b>Estimated time:</b>	0.00 hour
<b>Target version:</b>		<b>Spent time:</b>	2.50 hours
<b>Description</b>			
Relebohile asked if the fastq files could be renamed according to the experimental conditions.			

#### History

##### #1 - 05/14/2019 02:58 PM - Jon Ambler

- Status changed from New to In Progress
- % Done changed from 0 to 90

To make sure files are named correctly, a testing set was created where the files contained their original names to make sure they are not incorrectly labeled.

I wrote a little python script to go through all folders and rename accordingly, keeping information on the lane, R1/R2, and making sure all names are unique before running the rename.

It passed the testing stage, now I am waiting for the new files to be copied to an external to give to Relebohile before combining all the samples and renaming them.

##### #2 - 05/15/2019 09:28 AM - Jon Ambler

- Status changed from In Progress to Resolved
- % Done changed from 90 to 100

Files uploaded and renamed successfully.