

M. smegmatis variant calling pipeline - Support #114

Preparation of the reusable pipeline

02/14/2020 01:12 PM - Jon Ambler

Status:	In Progress	Start date:	02/14/2020
Priority:	Normal	Due date:	
Assignee:	Jon Ambler	% Done:	60%
Category:		Estimated time:	0.00 hour
Target version:		Spent time:	8.00 hours
Description			
The group requested that the pipeline be reusable.			
For this, I am modifying existing Nextflow pipelines to provide a extensive set of analyses in one pipeline.			
This pipeline will be related to the pipeline being developed for analysis of Pneumonia genomes.			

History

#1 - 06/04/2020 05:54 PM - Jon Ambler

- Status changed from New to In Progress
- % Done changed from 0 to 60

Overlap in development with the S. pneumoniae pipeline allowed both to be developed together.

Time spent getting the pipeline working specific to this project.