

Setting up a portable RNA-Seq pipeline for CBIO - Issues

#	Tracker	Status	Priority	Subject	Assignee	Updated
30	Support	In Progress	Normal	Update Nextflow to version 0.27.6	Gerrit Botha	04/09/2018 12:56 PM
25	Support	In Progress	Normal	Configure NGI-RNAseq pipeline to run on hex		05/17/2018 02:54 PM
24	Support	New	Normal	Setting up the NGI-RNAseq pipeline on UCT Hex	Katie Lennard	04/09/2018 09:51 AM