

Identification of biomarkers in keloids and folliculitis keloidalis nuchae (FKN) - Issues

#	Tracker	Status	Priority	Subject	Assignee	Updated
104	Feature	Resolved	Normal	Summary table of mRNA and proteomics analysis for identified gene targets, protein level mean abundance per group added.	Thys Potgieter	11/12/2019 11:12 AM
103	Feature	Resolved	Normal	Median group intensity added to output for visualization of proteomics target expression profiles	Thys Potgieter	11/12/2019 11:10 AM
101	Support	New	Normal	Secondary analysis: Transcription factor and pathway enrichment	Jon Ambler	10/04/2019 01:29 PM
100	Support	Resolved	High	Time seres analysis	Jon Ambler	10/04/2019 01:25 PM
93	Feature	Resolved	High	Improve normalization and imputaiton of tissue sample proteomics data and visualization	Thys Potgieter	08/05/2019 02:06 PM
92	Support	Resolved	Urgent	Differential expression analysis	Jon Ambler	10/04/2019 01:19 PM
90	Support	Resolved	High	QC of aligned reads	Jon Ambler	10/04/2019 01:30 PM
86	Support	Resolved	High	Analyze keloid proteomics data with mqmetaproteomics pipeline on ilifu	Thys Potgieter	07/02/2019 09:22 AM
85	Support	Resolved	High	MaxQuant run of raw proteomics data	Thys Potgieter	07/02/2019 09:16 AM
82	Support	Resolved	High	Preliminary analysis of Proteome identification data	Thys Potgieter	05/21/2019 12:57 PM
81	Support	Resolved	High	Full run on ilifu of RNA seq pipeline	Jon Ambler	06/19/2019 12:53 PM
80	Support	Resolved	High	Renaming all files according to the experiments	Jon Ambler	05/15/2019 09:28 AM
79	Support	Resolved	High	Uploading new data from BGI	Jon Ambler	05/13/2019 09:09 AM
78	Support	Resolved	Normal	Proteomics raw data uploaded to Blackburn Lab researchdata	Thys Potgieter	05/14/2019 10:30 AM
77	Support	Resolved	High	Testing pipeline on subset of data	Jon Ambler	05/13/2019 09:06 AM
74	Support	Resolved	Normal	Uploading of data	Jon Ambler	05/03/2019 09:39 AM