

## (http://multiqc.info)

A modular tool to aggregate results from bioinformatics analyses across many samples into a single report.

some key 0 some val some val some val some val some val some key 3 some val

Report generated on 2021-11-07, 23:46 based on data in: /work

## Lane Statistics 3

This is in a different section

Lane	Total # Reads	Total # PF Reads	Undetermined	
1.0				

## This data is grouped

Lane Statistics 2

This is a second description for a module sub-section

1.0

Lane Statistics 4

This is in a different other section

Lane	Total # Reads	Total # PF Reads	Undetermined
1.0			

## **Plasmid Detection**

Plasmid	Circular	Number of Contigs	Predicted Host Range
pXO1	Yes		
pXO2	Yes		

<u>MultiQC v1.12.dev0 (http://multiqc.info)</u> - Written by <u>Phil</u> <u>Ewels (http://phil.ewels.co.uk)</u>, available on <u>GitHub</u> (<u>https://github.com/ewels/MultiQC</u>).



(http://www.scilifelab.se/)

This report uses <u>HighCharts (http://www.highcharts.com/)</u>, jQuery (<u>https://jquery.com/</u>), jQuery UI (<u>https://jqueryui.com/</u>), <u>Bootstrap (http://getbootstrap.com/</u>), <u>FileSaver.js (https://github.com/eligrey/FileSaver.js)</u> and <u>clipboard.js</u> (<u>https://clipboardjs.com/</u>).