



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

<b>some key</b>	some val
<b>some key 0</b>	some val
<b>some key 1</b>	some val
<b>some key 2</b>	some val
<b>some key 3</b>	some val

Report generated on 2021-11-07, 23:45 based on data in:  
/work

## Lane Statistics 3

This is in a different section

Lane	Total # Reads	Total # PF Reads	Undetermined
1.0	196 298 213	186 009 926	5.45%

## This data is grouped

### Lane Statistics 2

This is a second description for a module sub-section

Lane	Total # Reads	Total # PF Reads	Undetermined
1.0	196 298 213	186 009 926	5.45%

## Lane Statistics 4

This is in a different other section

Lane	Total # Reads	Total # PF Reads	Undetermined
1.0	196 298 213	186 009 926	5.45%

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# Plasmid Detection

Plasmid	Circular	Number of Contigs	Predicted Host Range
pX01	Yes	1	Bacillaceae
pX02	Yes	1	Bacillus

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[MultiQC v1.12.dev0](#) - Written by [Phil Ewels](#), available on [GitHub](#).

This report uses [HighCharts](#), [jQuery](#), [jQuery UI](#), [Bootstrap](#), [FileSaver.js](#) and [clipboard.js](#).

