#### **Metagenomics Report**



### **Patient**

**CIDR** 

Lab ID	gstt_control_1	Date:Time	25/06/2024, 17:16
Workflow version	3.7		
Sample Type	none	Sample Class	positive_control
Time Interval	0.5 hrs	Barcode	11
MinKNOW experiment	GSTT_control_sample_01	MinKNOW Sample	GSTT_control_sample_01

# **Quality Control**

Total microbial reads	248	Total reads	250	Total human reads (count/percent of total reads)	0/0%
Total bases	763.50 Kb	Mean read length	3,054.0	Mean read quality	14.2
Internal control (TMV)	NA/NA	Negative Control	☐ Pass ☐ Fail	Unclassified/Below Threshold	1/10

## Taxanomic classification - Organisms above Threshold

Organism	Counts	Percentage
Bordetella parapertussis	44	38.26
Mycoplasma pneumoniae	42	36.52
Bordetella pertussis	27	23.48
Bordetella bronchiseptica	2	1.74

### Viral report

Organism	Counts
Human adenovirus 3	100
Human mastadenovirus C	32
Human mastadenovirus A	1



#### **Metagenomics Report**



## Antimicrobial resistance summary

Species	Gene	Hits	
No genes above threshold	NA	NA	





## Antimicrobial resistance report

No Results



#### **Metagenomics Report**



# Centrifuge Full

Organism	Counts	Percentage	
Bordetella parapertussis	44	38.26	
Mycoplasma pneumoniae	42	36.52	
Bordetella pertussis	27	23.48	
Bordetella bronchiseptica	2	1.74	