

WGS Quality Control Report

Batch Name: 2024-07-04

Experiment Name: 24ARS_EMR_LG788

Isolate No.	Sample ID	Description	ARSRL	WGS
1	23ARS_VSM0511	SAL68	Salmonella species	Salmonella enterica subsp. enterica serovar Enteritidis
2	24ARS_BRH0014	EMR142_2	Pseudomonas aeruginosa	Pseudomonas aeruginosa
3	24ARS_BRT0017	EMR141_2	Pseudomonas aeruginosa	Pseudomonas aeruginosa
4	24ARS_CRH0017	EMR146	Escherichia coli	Escherichia coli
5	24ARS_CRH0030	EMR148	Klebsiella pneumoniae	Klebsiella quasipneumoniae
6	24ARS_DMC0098	EMR149	Klebsiella pneumoniae	Klebsiella pneumoniae
7	24ARS_DMC0102	EMR157	Pseudomonas aeruginosa	Pseudomonas aeruginosa
8	24ARS_EVR0011	EMR155	Pseudomonas aeruginosa	Pseudomonas aeruginosa
9	24ARS_EVR0017	EMR158	Pseudomonas aeruginosa	Pseudomonas aeruginosa
10	24ARS_GMH0029	EMR145	Klebsiella pneumoniae	Klebsiella pneumoniae
11	24ARS_JLM0018	EMR143	Klebsiella pneumoniae	Klebsiella pneumoniae
12	24ARS_JLM0020	EMR144	Klebsiella pneumoniae	Klebsiella quasipneumoniae
13	24ARS_NKI0048	EMR147	Escherichia coli	Escherichia coli
14	24ARS_NKI0050	EMR150	Klebsiella pneumoniae	Klebsiella pneumoniae
15	24ARS_NKI0051	EMR151	Klebsiella pneumoniae	Klebsiella pneumoniae
16	24ARS_SLH0030	EMR154	Pseudomonas aeruginosa	Pseudomonas aeruginosa
17	24ARS_STU0024	EMR152	Klebsiella pneumoniae	Klebsiella pneumoniae
18	24ARS_STU0027	EMR153	Acinetobacter baumannii	Acinetobacter pittii
19	24ARS_ZMC0002	EMR156	Pseudomonas aeruginosa	Pseudomonas aeruginosa
20	UTP_BL_006	tricycle	Escherichia coli	Escherichia coli
21	UTP_ST_031	tricycle	Escherichia coli	Escherichia coli

Legend:

PASS	WARNING	FAILURE	EXCEEDS THRESHOLD METRIC/S	NON-CONCORDANT
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Sample excluded in the analysis

Sample ID	Description	Index reads	Remarks
24ARS_NKI0053	EMR159	0.8429	low read count

Isolate No.	Sample ID	Contamination	Contigs	GC Percent	N50	Total Length
1	23ARS_VSM0511	0	24	52.13	478941	4700439
2	24ARS_BRH0014	0	84	65.97	247991	6878546
3	24ARS_BRT0017	0	122	65.33	220628	7348540
4	24ARS_CRH0017	0	127	50.53	136602	5373741
5	24ARS_CRH0030	0	39	57.75	801312	5379490
6	24ARS_DMC0098	0	44	57.19	479099	5405179
7	24ARS_DMC0102	0	51	66.32	488628	6482852
8	24ARS_EVR0011	0	47	66.47	407140	6291421
9	24ARS_EVR0017	0	131	66.36	151006	6423637
10	24ARS_GMH0029	0	110	56.65	172819	5772594
11	24ARS_JLM0018	0	95	56.79	394359	5742923
12	24ARS_JLM0020	0	77	57.65	229344	5512178
13	24ARS_NKI0048	0	93	50.83	320376	5148168
14	24ARS_NKI0050	0	55	57.20	236088	5425644
15	24ARS_NKI0051	0	81	56.82	238924	5595191
16	24ARS_SLH0030	0	99	65.94	361062	6937344
17	24ARS_STU0024	0	55	57.13	303929	5493891
18	24ARS_STU0027	0	17	38.58	519454	3995354
19	24ARS_ZMC0002	0	89	66.46	197528	6319285
20	UTP_BL_006	0	44	50.61	279880	4940847

21	UTP_ST_031	0	109	50.42	166206	5380880
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Legend:

PASS	WARNING	FAILURE	EXCEEDS THRESHOLD METRIC/S	*Isolates were tagged with warning due to uncertain results of species identification using bactinspector or sequence identification levels.
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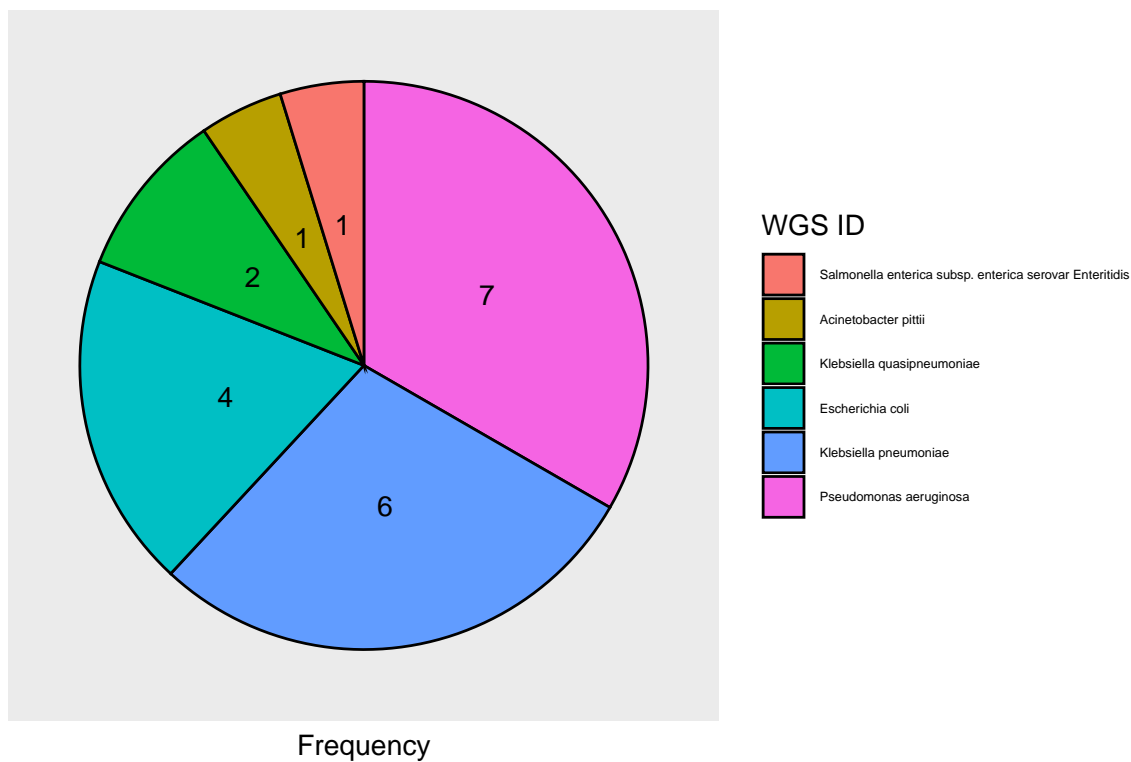
List of samples above/below QC threshold metrics

Sample ID	Remarks
No QC failures found.	

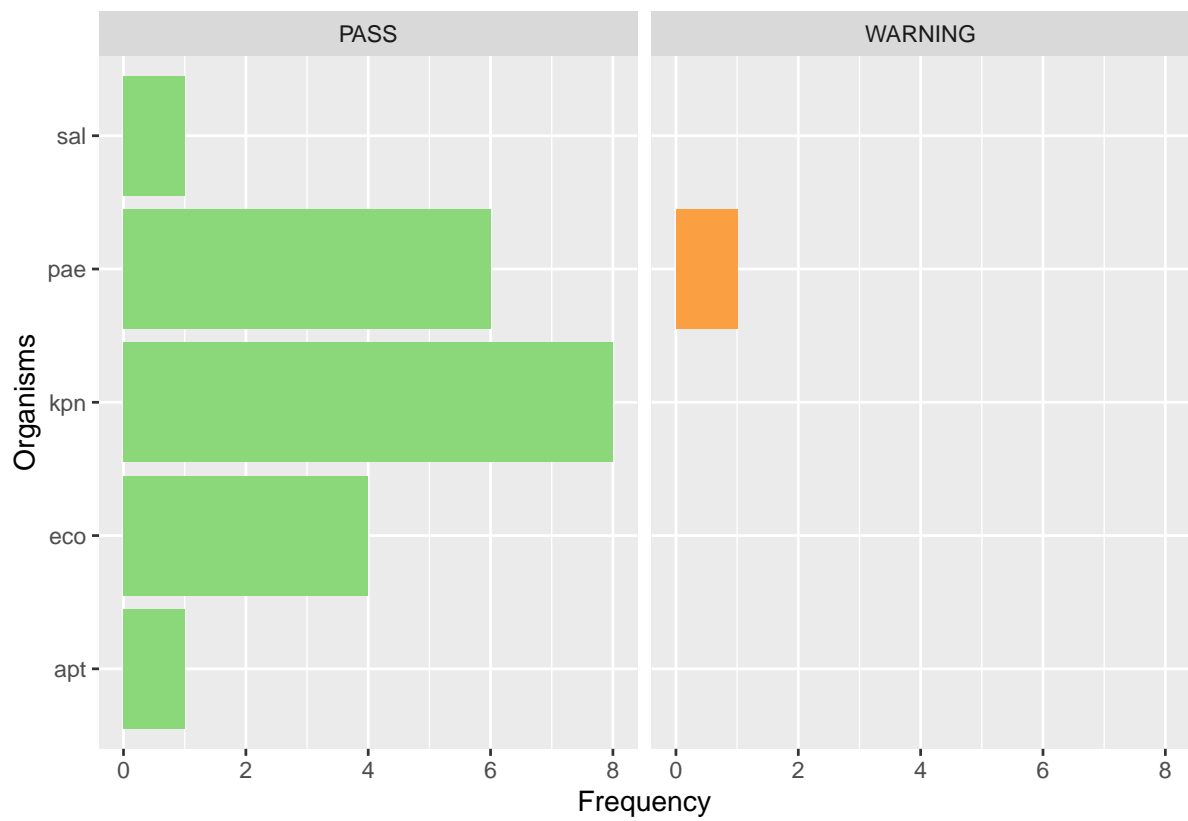
WGS_ID	Number
Pseudomonas aeruginosa	7
Klebsiella pneumoniae	6
Escherichia coli	4
Klebsiella quasipneumoniae	2
Acinetobacter pittii	1
Salmonella enterica subsp. enterica serovar Enteritidis	1

- 6 distinct species were identified among 21 isolates.
- 95.24 % (n=20) of the isolates passed the QC, while 4.76 % (n=1) were tagged with warning.
- Concordance between ARSRL and WGS species report was 100.00 %.

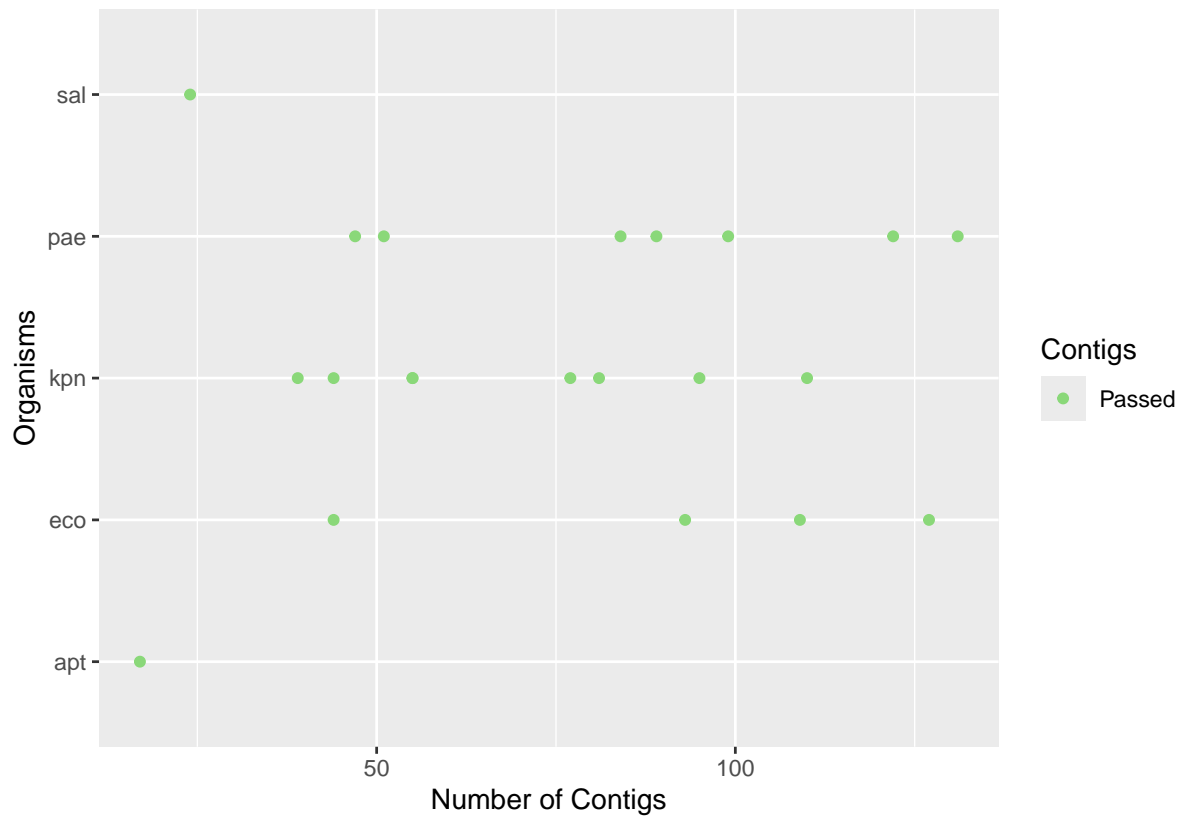
GRAPHS



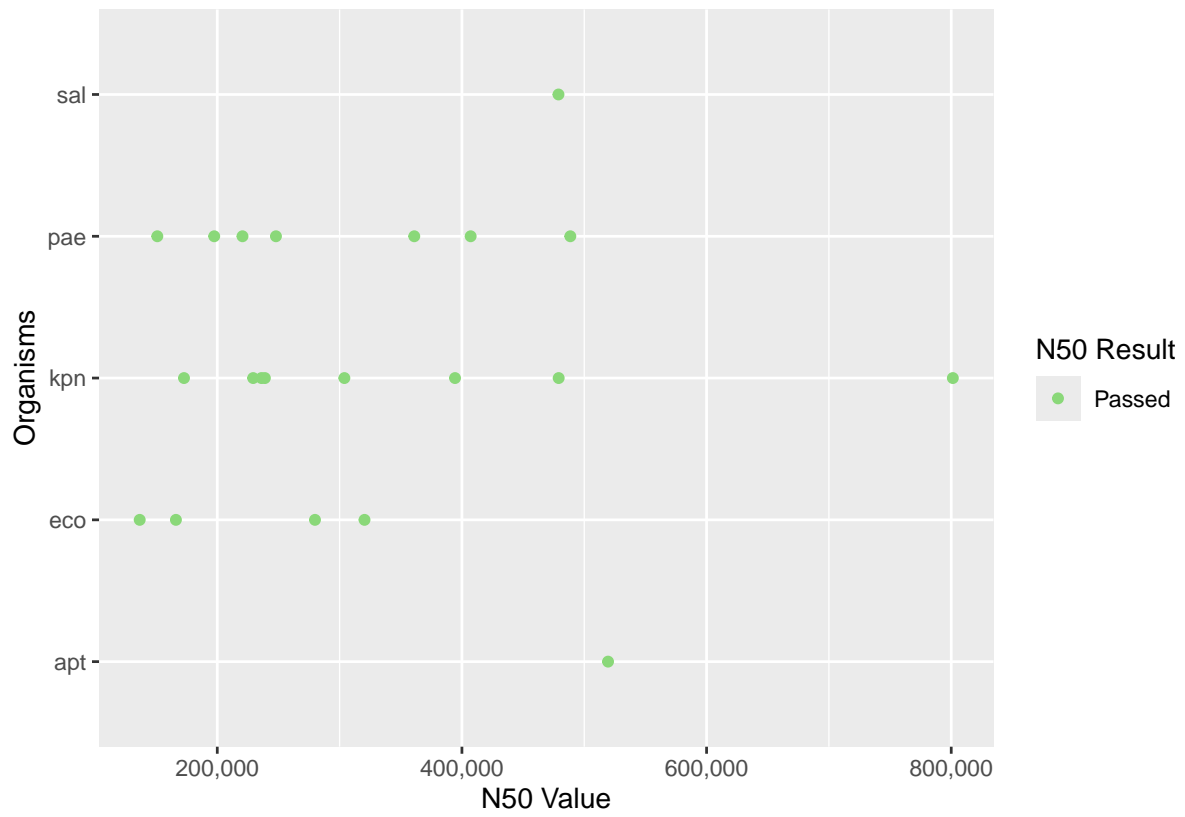
Result Classification



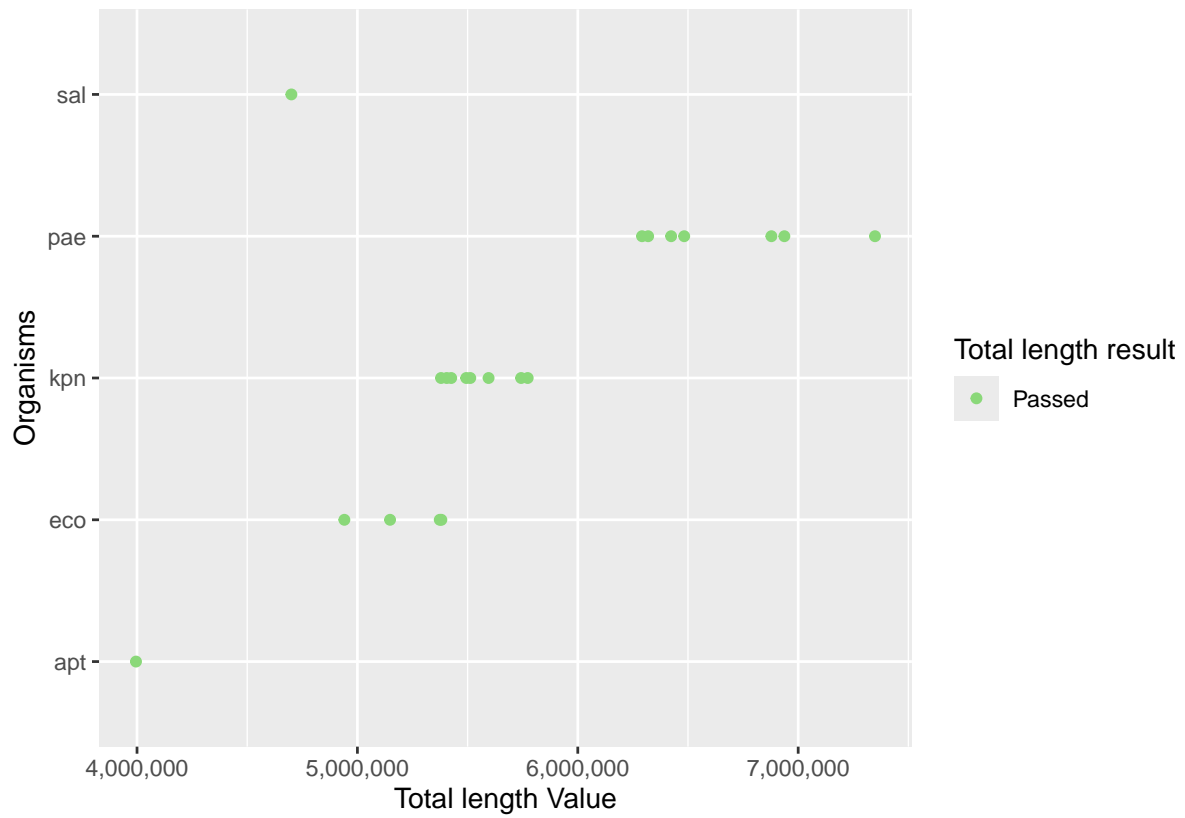
Number of contigs



N50 Value



Total Length



RECOMMENDATION:

Sample ID	Action	Reason
No further action required for this batch.		

MLST AND AMR GENE RESULTS:

sample_id	species	MLST	aroC	dnaN	hemD	hisD	purE	sucA	thrA
23ARS_VSM0511	Salmonella enterica subsp. enterica serovar Enteritidis	-	5,638	2	3	7	6	6	11

sample_id	species	MLST	aroC	dnaN	hemD	hisD	purE	sucA	thrA
24ARS_BRH0014	Pseudomonas aeruginosa	639	11	19	19	3	4	4	7
24ARS_BRT0017	Pseudomonas aeruginosa	-	28,347	5	36	3	3	13	7
24ARS_DMC0102	Pseudomonas aeruginosa	2617	84	3	20	71	4	7	1
24ARS_EVR0011	Pseudomonas aeruginosa	-	15,345	5	11	3	15	42	9
24ARS_EVR0017	Pseudomonas aeruginosa	641	6	5	6	5	4	4	7
24ARS_SLH0030	Pseudomonas aeruginosa	-	11,345	20	1	65	4	4	10
24ARS_ZMC0002	Pseudomonas aeruginosa	3014	16	5	12	3	3	1	18

sample_id	species	MLST	aroC	dnaN	hemD	hisD	purE	sucA	thrA
24ARS_CRH0017	Escherichia coli	2083	6	322	5	16	11	8	7
24ARS_NKI0048	Escherichia coli	131	53	40	47	13	36	28	29
UTP_ST_031	Escherichia coli	38	4	26	2	25	5	5	19

sample_id	species	MLST	aroC	dnaN	hemD	hisD	purE	sucA	thrA
24ARS_CRH0030	Klebsiella quasipneumoniae	-	18	22	74	22	123	20	99
24ARS_DMC0098	Klebsiella pneumoniae	17	2	1	1	1	4	4	4
24ARS_GMH0029	Klebsiella pneumoniae	147	3	4	6	1	7	4	38
24ARS_JLM0018	Klebsiella pneumoniae	39	2	1	2	4	9	1	14
24ARS_JLM0020	Klebsiella quasipneumoniae	6403	18	22	26	59	92	13	192
24ARS_NKI0050	Klebsiella pneumoniae	2050	2	3	2	37	10	1	15
24ARS_NKI0051	Klebsiella pneumoniae	15	1	1	1	1	1	1	1
24ARS_STU0024	Klebsiella pneumoniae	39	2	1	2	4	9	1	14

sample_id	species	MLST	aroC	dnaN	hemD	hisD	purE	sucA	thrA
24ARS_STU0027	Acinetobacter pittii	-	~101	~233	46	29	~124	~59	119