

Search Details

Job Title/ Assay Name : VanA(23/7)	Amplicon Sequence: CCACCGGCCTATCATCTTTATTAATAACCCAAAAGGCGGGAGT AGCTATCCCAGCATTTTTTCGCAACGATGTATGTCAACGATTG TCCA
Effective Date : 2025-07-23	Forward: CCACCGGCCTATCATCTTTA
Database : core_nt	Reverse: GATGTATGTCAACGATTTGTCCA
Alignment algorithm : blastN	Probe: AAGGCGGGAGTAGCTATCCCAGCA
Exclusive Organism:	Inclusive Organism:
RID: 80TXMC9H015	Total Result: 1006

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Citrobacter freundii/ 546	1	1	1	1	100%
Escherichia coli/562	1	1	1	1	100%
Staphylococcus/1279	1	1	1	1	100%
	71	69	69	56	81%
Total	1006	967	966	936	97%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Staphylococcus aureus/ 1280					
Staphylococcus haemolyticus/1283	1	1	1	1	100%
Enterococcus/1350	1	1	1	1	100%
Enterococcus faecalis/ 1351	61	60	60	60	100%
Enterococcus faecium/ 1352	806	793	792	791	100%
Enterococcus gallinarum/1353	1	1	1	1	100%
Atopobium minutum/ 1381	1	0	0	0	0%
Niallia circulans/1397	2	2	2	0	0%
Paenibacillus larvae/ 1464	9	0	0	0	0%
Brevibacillus laterosporus/1465	2	2	2	0	0%
	12	11	11	11	100%
Total	1006	967	966	936	97%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Clostridioides difficile/ 1496					
Lentilactobacillus hilgardii/1588	2	0	0	0	0%
Oerskovia turbata/1713	1	1	1	1	100%
Thermoactinomyces vulgaris/2026	2	0	0	0	0%
Enterococcus avium/ 33945	2	2	2	2	100%
Enterococcus casseliflavus/37734	1	1	1	1	100%
Enterococcus cecorum/ 44008	3	3	3	3	100%
Paenibacillus/44249	2	0	0	0	0%
Paenibacillus apiarius/ 46240	3	3	3	0	0%
Paenibacillus thiaminolyticus/49283	5	3	3	0	0%
	3	3	3	3	100%
Total	1006	967	966	936	97%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Enterococcus durans/ 53345					
Paenibacillus sonchi/ 373687	1	0	0	0	0%
Paenibacillus jilunlii/ 682956	1	0	0	0	0%
Brevibacillus aydinogluensis/927786	1	0	0	0	0%
Enterococcus saigonensis/1805431	2	2	2	2	100%
Brevibacillus sedimenti/ 2613334	1	0	0	0	0%
Paenibacillus melissococcoides/ 2912268	6	6	6	0	0%
Total	1006	967	966	936	97%