

Search Details

Job Title/ Assay Name : VanC1(24/7)	Amplicon Sequence: ACTGCTTGAAC TAATGAACCTGCCTTATGTTGGTTGCCATGTC GCTGCCTCCGCATTATGTATGAACAAATGGCTCTTGCATCAAC TTGCTGA
Effective Date : 2025-07-24	Forward: ACTGCTTGAAC TAATGAACCTG
Database : core_nt	Reverse: CTCTTGCATCAACTTGCTGA
Alignment algorithm : blastN	Probe: TATGTTGGTTGCCATGTCGCTGCCTC
Exclusive Organism:	Inclusive Organism:
RID: 843FDSVF015	Total Result: 195

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Enterococcus/1350	21	4	4	4	100%
Enterococcus faecalis/ 1351	8	6	6	6	100%
Enterococcus faecium/ 1352	6	3	3	3	100%
Total	195	46	45	45	100%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Enterococcus gallinarum/1353	36	33	32	32	100%
Bacillus/1386	17	0	0	0	0%
Meleagris gallopavo/ 9103	1	0	0	0	0%
Enterococcus casseliflavus/37734	49	0	0	0	0%
Ureibacillus thermosphaericus/ 51173	1	0	0	0	0%
Ureibacillus/160795	1	0	0	0	0%
Bacillus altitudinis/ 293387	50	0	0	0	0%
Bacillus aerius/293388	1	0	0	0	0%
Bacillus xiamenensis/ 1178537	1	0	0	0	0%
Candidatus Enterococcus mangumiae/2230878	1	0	0	0	0%
Total	195	46	45	45	100%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Mesobacillus/2675231	1	0	0	0	0%
Enterococcus innesii/ 2839759	1	0	0	0	0%
Total	195	46	45	45	100%