

Search Details

Job Title/ Assay Name : VanC1(29/7)	Amplicon Sequence: ACTGCTTGAAC TAATGAACCTGCCTTATGTTGGTTGCCATGTCGCTG CCTCCGCATTATGTATGAACAAATGGCTCTTGCATCAACTTGCTGA
Effective Date : 2025-07-29	Forward: ACTGCTTGAAC TAATGAACCTG
Database : core_nt	Reverse: TCAGCAAGTTGATGCAAGAG
Alignment algorithm : blastN	Probe: TATGTTGGTTGCCATGTCGCTGCCTC
Exclusive Organism:	Inclusive Organism:
RID: 8HDVNNGGJ013	Total Result: 195
App Version: 1.0	

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Enterococcus gallinarum/1353	36	33	32	32	100%
Enterococcus faecalis/1351	8	7	7	6	86%
Enterococcus/1350	21	21	21	4	19%
Enterococcus faecium/1352	6	3	3	3	100%
Enterococcus casseliflavus/37734	49	41	41	0	0%
Enterococcus innesii/2839759	1	1	1	0	0%
Bacillus altitudinis/293387	50	0	0	0	0%
Bacillus/1386	17	0	0	0	0%
Meleagris gallopavo/9103	1	0	0	0	0%
Ureibacillus thermosphaericus/51173	1	0	0	0	0%
Ureibacillus/160795	1	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%
Mesobacillus/2675231	1	0	0	0	0%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Total	195	106	105	45	43%