

Search Details

Job Title/ Assay Name : GES	Amplicon Sequence: CGTTCAAGTTTCCGCTAGCCGCGCTGGTCTTTGAAAGAATTGACTC AGGCACCGAGCGGGGGGATCGAAAAC TTTCATATGGGCC
Effective Date : 2025-11-19	Forward: CGTTCAAGTTTCCGCTAG
Database : core_nt	Reverse: GGCCCATATGAAAGTTTTTC
Alignment algorithm : blastN	Probe: AATTGACTCAGGCACCGAGCG
Exclusive Organism:	Inclusive Organism:
RID: HUKZNK57014	Total Result: 409
App Version: 1.0	

Species/tax-id Detected	# Accessions Returned	# Meeting Inclusion Criteria	# Accessions Analyzed	# Accession Detected	% Detected
Pseudomonas aeruginosa/287	196	193	193	193	100%
Acinetobacter baumannii/470	41	41	41	41	100%
Klebsiella pneumoniae/573	28	28	28	28	100%
Aeromonas caviae/648	18	18	18	18	100%
Klebsiella quasipneumoniae/1463165	16	16	16	16	100%
Enterobacter cloacae/550	13	13	13	13	100%
Escherichia coli/562	13	13	13	13	100%
Serratia marcescens/615	9	9	9	9	100%
Klebsiella michiganensis/1134687	8	8	8	8	100%
Citrobacter freundii/546	7	7	7	7	100%
Citrobacter portucalensis/1639133	6	6	6	6	100%
Aeromonas/642	5	5	5	5	100%
Aeromonas hydrophila/644	5	5	5	5	100%
Klebsiella oxytoca/571	4	4	4	4	100%
Aeromonas veronii/654	4	4	4	4	100%
Raoultella ornithinolytica/54291	4	4	4	4	100%
Enterobacter kobei/208224	4	4	4	4	100%
Enterobacter roggenskampi/1812935	4	4	4	4	100%
Klebsiella variicola/244366	3	3	3	3	100%

Species/tax-id Detected	# Accessions Returned	# Meeting Inclusion Criteria	# Accessions Analyzed	# Accession Detected	% Detected
Acinetobacter/469	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Raoultella planticola/575	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Citrobacter braakii/57706	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Enterobacter asburiae/61645	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Klebsiella grimontii/2058152	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Pseudomonas putida/303	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Citrobacter/544	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Morganella morganii/582	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Salmonella/590	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Aeromonas media/651	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Stenotrophomonas maltophilia/40324	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Pandoraea/93217	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Undibacterium/401469	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Fluviibacter phosphoraccumulans/1751046	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Cloning vector pET28A-blaGES-9/2573024	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Proteus mirabilis/584	<u>1</u>	<u>0</u>	<u>0</u>	<u>0</u>	0%
Total	409	405	405	405	100%