

Search Details

Job Title/ Assay Name : 450-072-LMP: EcloaXX-F1/ Ecloa 496R/ Ecloa 441senP-FAM ; Syn_E.cloa_NC014121	Amplicon Sequence: ACAAAGGAGTCGGGATGAGTTCGCAATCCCAGGCCAAATCACCGG AAGCCTTACGAGCAATGGTCGCCGGGACGCTGGCTAACTTTCAGCA TCCAA
Effective Date : 2025-08-15	Forward: ACAAAGGAGTCGGGATGAGTTC
Database : core_nt	Reverse: TTGGATGCTGAAAGTTAGCC
Alignment algorithm : blastN	Probe: CCGGAAGCCTTACGAGCAATGGTC
Exclusive Organism:	Inclusive Organism:
RID: 9ZAGNT2C015	Total Result: 5000
App Version: 1.0	

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Enterobacter hormaechei/158836	534	534	534	533	100%
Enterobacter asburiae/61645	66	66	66	66	100%
Enterobacter roggenskampi/1812935	61	61	61	61	100%
Enterobacter cloacae/550	53	53	53	52	98%
Enterobacter ludwigii/299767	46	46	46	46	100%
Enterobacter/547	47	47	47	43	91%
Enterobacter kobei/208224	38	38	38	38	100%
Enterobacter cloacae complex/354276	33	33	33	33	100%
Enterobacter bugandensis/881260	22	22	22	22	100%
Enterobacter mori/539813	9	9	9	9	100%
Enterobacter cancerogenus/69218	8	8	8	8	100%
Enterobacter chuandaensis/2497875	6	6	6	6	100%
Enterobacter sichuanensis/2071710	4	4	4	4	100%
Enterobacter chengduensis/2494701	3	3	3	3	100%
Enterobacter pseudoroggenskampii/2996112	3	3	3	3	100%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Lelliottia/1330545	<u>3</u>	<u>3</u>	<u>3</u>	<u>2</u>	67%
Enterobacter soli/885040	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Enterobacter oligotrophicus/2478464	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Enterobacter quasiroggenkampii/2497436	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Enterobacter wuhouensis/2529381	<u>2</u>	<u>2</u>	<u>2</u>	<u>2</u>	100%
Shigella flexneri/623	<u>63</u>	<u>63</u>	<u>63</u>	<u>1</u>	2%
Lelliottia amnigena/61646	<u>8</u>	<u>8</u>	<u>8</u>	<u>1</u>	13%
Lelliottia nimipressuralis/69220	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Enterobacter huaxiensis/2494702	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Enterobacter vonholyi/2797505	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Enterobacter dykesii/2797506	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Enterobacter pasteurii/3029761	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Enterobacter intestinihominis/3133180	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Enterobacter adelaidei/3142537	<u>1</u>	<u>1</u>	<u>1</u>	<u>1</u>	100%
Escherichia coli/562	<u>3623</u>	<u>3623</u>	<u>3623</u>	<u>0</u>	0%
Klebsiella aerogenes/548	<u>93</u>	<u>93</u>	<u>93</u>	<u>0</u>	0%
Klebsiella oxytoca/571	<u>65</u>	<u>65</u>	<u>65</u>	<u>0</u>	0%
Shigella sonnei/624	<u>42</u>	<u>42</u>	<u>42</u>	<u>0</u>	0%
Leclercia adecarboxylata/83655	<u>23</u>	<u>23</u>	<u>23</u>	<u>0</u>	0%
Shigella dysenteriae/622	<u>20</u>	<u>20</u>	<u>20</u>	<u>0</u>	0%
Shigella boydii/621	<u>15</u>	<u>15</u>	<u>15</u>	<u>0</u>	0%
Kosakonia cowanii/208223	<u>15</u>	<u>15</u>	<u>15</u>	<u>0</u>	0%
Escherichia marmotae/1499973	<u>15</u>	<u>15</u>	<u>15</u>	<u>0</u>	0%
Klebsiella pneumoniae/573	<u>13</u>	<u>13</u>	<u>13</u>	<u>0</u>	0%
Leclercia/83654	<u>11</u>	<u>11</u>	<u>11</u>	<u>0</u>	0%
Leclercia pneumoniae/2815358	<u>5</u>	<u>5</u>	<u>5</u>	<u>0</u>	0%

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Raoultella terrigena/577	<u>4</u>	<u>4</u>	<u>4</u>	<u>0</u>	0%
Kosakonia sacchari/1158459	<u>4</u>	<u>4</u>	<u>4</u>	<u>0</u>	0%
Kosakonia/1330547	<u>4</u>	<u>4</u>	<u>4</u>	<u>0</u>	0%
Siccibacter colletis/1505757	<u>3</u>	<u>3</u>	<u>3</u>	<u>0</u>	0%
Kosakonia pseudosacchari/1646340	<u>3</u>	<u>3</u>	<u>3</u>	<u>0</u>	0%
Escherichia/561	<u>2</u>	<u>2</u>	<u>2</u>	<u>0</u>	0%
Pseudescherichia vulneris/566	<u>2</u>	<u>2</u>	<u>2</u>	<u>0</u>	0%
Klebsiella/570	<u>2</u>	<u>2</u>	<u>2</u>	<u>0</u>	0%
Siccibacter turicensis/357233	<u>2</u>	<u>2</u>	<u>2</u>	<u>0</u>	0%
Franconibacter helveticus/357240	<u>2</u>	<u>2</u>	<u>2</u>	<u>0</u>	0%
[Enterobacter] lignolyticus/1334193	<u>2</u>	<u>2</u>	<u>2</u>	<u>0</u>	0%
Superficieibacter/2303321	<u>2</u>	<u>2</u>	<u>2</u>	<u>0</u>	0%
unidentified bacterial endosymbiont/2355	<u>1</u>	<u>1</u>	<u>1</u>	<u>0</u>	0%
Triticum aestivum/4565	<u>1</u>	<u>1</u>	<u>1</u>	<u>0</u>	0%
Kosakonia oryzendophytica/1005665	<u>1</u>	<u>1</u>	<u>1</u>	<u>0</u>	0%
Escherichia ruysiae/2608867	<u>1</u>	<u>1</u>	<u>1</u>	<u>0</u>	0%
Kosakonia calanthes/3139408	<u>1</u>	<u>1</u>	<u>1</u>	<u>0</u>	0%
Klebsiella pneumoniae complex/3390273	<u>1</u>	<u>1</u>	<u>1</u>	<u>0</u>	0%
Total	5000	5000	5000	946	19%