

## Project: G0266763-1.SQD Contig 1

			102030
			GGGAATTGTGAGCGGATAACAATTCCCCTC
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GGGaATTGTGAGCGGATAaCAATTCCCCTC	
G0266763-1.seq (1>2469)	→	TC	
			405060
			TAGAAATAATTTTGTTTAACTTTAAGAAGG
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TAGAAATAATTTTGTTTAACTTTAAGAAGG	
G0266763-1.seq (1>2469)	→	TAGAAATAATTTTGTTTAACTTTAAGAAGG	
			708090
			AGATATACCATGTCCCCTATACTAGGTTAT
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	AGATATACCATGTCCCCTATACTAGGTTAT	
G0266763-1.seq (1>2469)	→	AGATATACCATGTCCCCTATACTAGGTTAT	
			100110120
			TGGAAAATTAAGGGCCTTGTGCAACCCACT
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TGGAAAATTAAGGGCCTTGTGCAACCCACT	
G0266763-1.seq (1>2469)	→	TGGAAAATTAAGGGCCTTGTGCAACCCACT	
			130140150
			CGACTTCTTTTGGAATATCTTGAAGAAAA
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	CGACTTCTTTTGGAATATCTTGAAGAAAA	
G0266763-1.seq (1>2469)	→	CGACTTCTTTTGGAATATCTTGAAGAAAA	
			160170180
			TATGAAGAGCATTGTATGAGCGCGATGAA
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TATGAAGAGCATTGTATGAGCGCGATGAA	
G0266763-1.seq (1>2469)	→	TATGAAGAGCATTGTATGAGCGCGATGAA	
			190200210
			GGTGATAAATGGCGAAACAAAAAGTTTGAA
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GGTGATAAATGGCGAAACAAAAAGTTTGAA	
G0266763-1.seq (1>2469)	→	GGTGATAAATGGCGAAACAAAAAGTTTGAA	
			220230240
			TTGGGTTTGGAGTTTCCCAATCTTCCTTAT
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TTGGGTTTGGAGTTTCCCAATCTTCCTTAT	
G0266763-1.seq (1>2469)	→	TTGGGTTTGGAGTTTCCCAATCTTCCTTAT	
			250260270
			TATATTGATGGTGATGTTAAATTAACACAG
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TATATTGATGGTGATGTTAAATTAACACAG	
G0266763-1.seq (1>2469)	→	TATATTGATGGTGATGTTAAATTAACACAG	
			280290300
			TCTATGGCCATCATACGTTATATAGCTGAC
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TCTATGGCCATCATACGTTATATAGCTGAC	
G0266763-1.seq (1>2469)	→	TCTATGGCCATCATACGTTATATAGCTGAC	

## Project: G0266763-1.SQD Contig 1

			310320330	AAGCACAAACATGTTGGGTGGTTGTCCAAAA
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	AAGCACAAACATGTTGGGTGGTTGTCCAAAA		
G0266763-1.seq (1>2469)	→	AAGCACAAACATGTTGGGTGGTTGTCCAAAA		
			340350360	GAGCGTGCAGAGATTTCAATGCTTGAAGGA
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GAGCGTGCAGAGATTTCAATGCTTGAAGGA		
G0266763-1.seq (1>2469)	→	GAGCGTGCAGAGATTTCAATGCTTGAAGGA		
			370380390	GCGGTTTTGGATATTAGATACGGTGTTCG
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GCGGTTTTGGATATTAGATACGGTGTTCG		
G0266763-1.seq (1>2469)	→	GCGGTTTTGGATATTAGATACGGTGTTCG		
			400410420	AGAATTGCATATAGTAAAGACTTTGAAACT
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	AGAATTGCATATAGTAAAGACTTTGAAACT		
G0266763-1.seq (1>2469)	→	AGAATTGCATATAGTAAAGACTTTGAAACT		
			430440450	CTCAAAGTTGATTTTCTTAGCAAGCTACCT
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	CTCAAAGTTGATTTTCTTAGCAAGCTACCT		
G0266763-1.seq (1>2469)	→	CTCAAAGTTGATTTTCTTAGCAAGCTACCT		
			460470480	GAAATGCTGAAAATGTTCTGAAGATCGTTTA
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GAAATGCTGAAAATGTTCTGAAGATCGTTTA		
G0266763-1.seq (1>2469)	→	GAAATGCTGAAAATGTTCTGAAGATCGTTTA		
			490500510	TGTCATAAAACATATTTAAATGGTGATCAT
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TGTCATAAAACATATTTAAATGGTGATCAT		
G0266763-1.seq (1>2469)	→	TGTCATAAAACATATTTAAATGGTGATCAT		
			520530540	GTAACCCATCCTGACTTCATGTTGTATGAC
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GTAACCCATCCTGACTTCATGTTGTATGAC		
G0266763-1.seq (1>2469)	→	GTAACCCATCCTGACTTCATGTTGTATGAC		
			550560570	GCTCTTGATGTTGTTTTATACATGGACCCA
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GCTCTTGATGTTGTTTTATACATGGACCCA		
G0266763-1.seq (1>2469)	→	GCTCTTGATGTTGTTTTATACATGGACCCA		
			580590600	ATGTGCCTGGATGCGTTCCCAAATTAGTT
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	ATGTGCCTGGATGCGTTCCCAAATTAGTT		
G0266763-1.seq (1>2469)	→	ATGTGCCTGGATGCGTTCCCAAATTAGTT		

## Project: G0266763-1.SQD Contig 1

		610	620	630
		TGTTTTAAAAAACGTATTGAAGCTATCCCA		
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TGTTTTAAAAAACGTATTGAAGCTATCCCA		
G0266763-1.seq (1>2469)	→	TGTTTTAAAAAACGTATTGAAGCTATCCCA		
		640	650	660
		CAAATTGATAAGTACTTGAAATCCAGCAAG		
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	CAAATTGATAAGTACTTGAAATCCAGCAAG		
G0266763-1.seq (1>2469)	→	CAAATTGATAAGTACTTGAAATCCAGCAAG		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	TGATAAGTACTTGAAATCCAGCAAG		
		670	680	690
		TATATAGCATGGCCTTTGCAGGGCTGGCAA		
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TATATAGCATGGCCTTTGCAGGGCTGGCAA		
G0266763-1.seq (1>2469)	→	TATATAGCATGGCCTTTGCAGGGCTGGCAA		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	TATATAGCATGGCCTTTGCAGGGcTgGCAA		
		700	710	720
		GCCACGTTTGGTGGTGGCGACCATCCTCCA		
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	GCCACGTTTGGTGGTGGCGACCATCCTCCA		
G0266763-1.seq (1>2469)	→	GCCACGTTTGGTGGTGGCGACCATCCTCCA		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	GCCACGTTTGGTGGTGGCGACCATCCTCCA		
		730	740	750
		AAATCGGATGCAAGCCTTGTTCCACGTGGT		
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	AAATCGGATGCAAGCCTTGTTCCACGTGGT		
G0266763-1.seq (1>2469)	→	AAATCGGATGCAAGCCTTGTTCCACGTGGT		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	AAATCGGATGCAAGCCTTGTTCCACGTGGT		
		760	770	780
		TCTGTCACAAGTTTGTACAAAAAAGCTGAA		
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	TCTGTCACAAGTTTGTACAAAAAAGCTGAA		
G0266763-1.seq (1>2469)	→	TCTGTCACAAGTTTGTACAAAAAAGCTGAA		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	TCTGTCACAAGTTTGTACAAAAAAGCTGAA		
		790	800	810
		CGAGAAACGTAAATGATATAAATATCAAT		
F02-Y94913-g0266763-1_pET-seqr.ab1 (1>793)	→	CGAGAAACGTAAA		
G0266763-1.seq (1>2469)	→	CGAGAAACGTAAATGATATAAATATCAAT		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	CGAGAAACGTAAATGATATAAATATCAAT		
		820	830	840
		ATATTAAATTAGATTTTGCATAAAAAACAG		
G0266763-1.seq (1>2469)	→	ATATTAAATTAGATTTTGCATAAAAAACAG		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	ATATTAAATTAGATTTTGCATAAAAAACAG		
		850	860	870
		ACTACATAATACTGTAAAACACAACATATC		
G0266763-1.seq (1>2469)	→	ACTACATAATACTGTAAAACACAACATATC		
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	ACTACATAATACTGTAAAACACAACATATC		

Project: G0266763-1.SQD Contig 1

		<div><div></div><div>880890900</div></div>
		CAGTCACTATGGCGGCCGCATTAGGCACCC
G0266763-1.seq (1>2469)	→	CAGTCACTATGGCGGCCGCATTAGGCACCC
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	CAGTCACTATGGCGGCCGCATTAGGCACCC
		<div><div></div><div>910920930</div></div>
		CAGGCTTTACACTTTATGCTTCCGGCTCGT
G0266763-1.seq (1>2469)	→	CAGGCTTTACACTTTATGCTTCCGGCTCGT
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	CAGGCTTTACACTTTATGCTTCCGGCTCGT
		<div><div></div><div>940950960</div></div>
		ATAATGTGTGGATTTTGAGTTAGGATCCGT
G0266763-1.seq (1>2469)	→	ATAATGTGTGGATTTTGAGTTAGGATCCGT
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	ATAATGTGTGGATTTTGAGTTAGGATCCGT
		<div><div></div><div>970980990</div></div>
		CGAGATTTTCAGGAGCTAAGGAAGCTAAAA
G0266763-1.seq (1>2469)	→	CGAGATTTTCAGGAGCTAAGGAAGCTAAAA
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	CGAGATTTTCAGGAGCTAAGGAAGCTAAAA
		<div><div></div><div>10001010102</div></div>
		TGGAGAAAAAATCACTGGATATAACCACCG
G0266763-1.seq (1>2469)	→	TGGAGAAAAAATCACTGGATATAACCACCG
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	TGGAGAAAAAATCACTGGATATAACCACCG
		<div><div></div><div>10301040105</div></div>
		TTGATATATCCCAATGGCATCGTAAAGAAC
G0266763-1.seq (1>2469)	→	TTGATATATCCCAATGGCATCGTAAAGAAC
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	TTGATATATCCCAATGGCATCGTAAAGAAC
		<div><div></div><div>10601070108</div></div>
		ATTTTGAGGCATTTTCAGTCAGTTGCTCAAT
G0266763-1.seq (1>2469)	→	ATTTTGAGGCATTTTCAGTCAGTTGCTCAAT
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	ATTTTGAGGCATTTTCAGTCAGTTGCTCAAT
		<div><div></div><div>10901100111</div></div>
		GTACCTATAACCAGACCGTTCAGCTGGATA
G0266763-1.seq (1>2469)	→	GTACCTATAACCAGACCGTTCAGCTGGATA
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	GTACCTATAACCAGACCGTTCAGCTGGATA
		<div><div></div><div>11201130114</div></div>
		TTACGGCCTTTTTTAAAGACCGTAAAGAAAA
G0266763-1.seq (1>2469)	→	TTACGGCCTTTTTTAAAGACCGTAAAGAAAA
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	TTACGGCCTTTTTTAAAGACCGTAAAGAAAA
		<div><div></div><div>11501160117</div></div>
		ATAAGCACAAGTTTTATCCGGCCTTTATTC
G0266763-1.seq (1>2469)	→	ATAAGCACAAGTTTTATCCGGCCTTTATTC
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1 (1>793)	→	ATAAGCACAAGTTTTATCCGGCCTTTATTC

			<div><div></div><div>11801190120</div></div>
			ACATTCTTGCCCGCCTGATGAATGCTCATC
G0266763-1.seq(1>2469)	→	ACATTCTTGCCCGCCTGATGAATGCTCATC	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	ACATTCTTGCCCGCCTGATGAATGCTCATC	
			<div><div></div><div>12101220123</div></div>
			CGGAATTCCGTATGGCAATGAAAGACGGTG
G0266763-1.seq(1>2469)	→	CGGAATTCCGTATGGCAATGAAAGACGGTG	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	CGGAATTCCGTATGGCAATGAAAGACGGTG	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	CGGTG	
			<div><div></div><div>12401250126</div></div>
			AGCTGGTGATATGGGATAGTGTTACCCCTT
G0266763-1.seq(1>2469)	→	AGCTGGTGATATGGGATAGTGTTACCCCTT	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	AGCTGGTGATATGGGATAGTGTTACCCCTT	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	AGCTGGTGATATGGGATAGTGTTACCCCTT	
			<div><div></div><div>12701280129</div></div>
			GTTACACCGTTTTCCATGAGCAAAGTGA
G0266763-1.seq(1>2469)	→	GTTACACCGTTTTCCATGAGCAAAGTGA	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	GTTACACCGTTTTCCATGAGCAAAGTGA	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	GTTACACCGTTTTCCATGAGCAAAGTGA	
			<div><div></div><div>13001310132</div></div>
			CGTTTTTCATCGCTCTGGAGTGAATACCACG
G0266763-1.seq(1>2469)	→	CGTTTTTCATCGCTCTGGAGTGAATACCACG	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	CGTTTTTCATCGCTCTGGAGTGAATACCACG	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	CGTTTTTCATCGCTCTGGAGTGAATACCACG	
			<div><div></div><div>13301340135</div></div>
			ACGATTTCCGGCAGTTTCTACACATATATT
G0266763-1.seq(1>2469)	→	ACGATTTCCGGCAGTTTCTACACATATATT	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	ACGATTTCCGGCAGTTTCTACACATATATT	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	ACGATTTCCGGCAGTTTCTACACATATATT	
			<div><div></div><div>13601370138</div></div>
			CGCAAGATGTGGCGTGTTACGGTGAAAACC
G0266763-1.seq(1>2469)	→	CGCAAGATGTGGCGTGTTACGGTGAAAACC	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	CGCAAGATGTGGCGTGTTACGGTGAAAACC	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	CGCAAGATGTGGCGTGTTACGGTGAAAACC	
			<div><div></div><div>13901400141</div></div>
			TGGCCTATTTCCCTAAAGGGTTTATTGAGA
G0266763-1.seq(1>2469)	→	TGGCCTATTTCCCTAAAGGGTTTATTGAGA	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	TGGCCTATTTCCCTAAAGGGTTTATTGAGA	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	TGGCCTATTTCCCTAAAGGGTTTATTGAGA	
			<div><div></div><div>14201430144</div></div>
			ATATGTTTTTCGTCTCAGCCAATCCCTGGG
G0266763-1.seq(1>2469)	→	ATATGTTTTTCGTCTCAGCCAATCCCTGGG	
E10-Y94913-g0266763-1_g0266763-1-seq2.ab1(1>793)	→	ATATGTTTTTCGTCTCAG	
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	ATATGTTTTTCGTCTCAGCCAATCCCTGGG	

		14501460147
		TGAGTTTCACCAGTTTTGATTTAAACGTGG
G0266763-1.seq (1>2469)	→	TGAGTTTCACCAGTTTTGATTTAAACGTGG
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	TGAGTTTCACCAGTTTTGATTTAAACGTGG
		14801490150
		CCAATATGGACAACTTCTTCGCCCCCGTTT
G0266763-1.seq (1>2469)	→	CCAATATGGACAACTTCTTCGCCCCCGTTT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	CCAATATGGACAACTTCTTCGCCCCCGTTT
		15101520153
		TCACTATGGGCAAATATTATACGCAAGGCG
G0266763-1.seq (1>2469)	→	TCACTATGGGCAAATATTATACGCAAGGCG
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	TCACTATGGGCAAATATTATACGCAAGGCG
		15401550156
		ACAAGGTGCTGATGCCGCTGGCGATTCAGG
G0266763-1.seq (1>2469)	→	ACAAGGTGCTGATGCCGCTGGCGATTCAGG
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	ACAAGGTGCTGATGCCGCTGGCGATTCAGG
		15701580159
		TTCATCATGCCGTTTGTGATGGCTTCCATG
G0266763-1.seq (1>2469)	→	TTCATCATGCCGTTTGTGATGGCTTCCATG
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	TTCATCATGCCGTTTGTGATGGCTTCCATG
		16001610162
		TCGGCAGAATGCTTAATGAATTACAACAGT
G0266763-1.seq (1>2469)	→	TCGGCAGAATGCTTAATGAATTACAACAGT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	TCGGCAGAATGCTTAATGAATTACAACAGT
		16301640165
		ACTGCGATGAGTGGCAGGGCGGGGCGTAAA
G0266763-1.seq (1>2469)	→	ACTGCGATGAGTGGCAGGGCGGGGCGTAAA
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	ACTGCGATGAGTGGCAGGGCGGGGCGTAAA
		16601670168
		CGCGTGGATCCGGCTTACTAAAAGCCAGAT
G0266763-1.seq (1>2469)	→	CGCGTGGATCCGGCTTACTAAAAGCCAGAT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	CGCGTGGATCCGGCTTACTAAAAGCCAGAT
		16901700171
		AACAGTATGCGTATTTGCGCGCTGATTTTT
G0266763-1.seq (1>2469)	→	AACAGTATGCGTATTTGCGCGCTGATTTTT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	AACAGTATGCGTATTTGCGCGCTGATTTTT
E04-Y94913-G0266763-1_pET-seqF.ab1 (1>794)	←	TTTGCGCGCTGATTTTT
		17201730174
		GCGGTATAAGAATATATACTGATATGTATA
G0266763-1.seq (1>2469)	→	GCGGTATAAGAATATATACTGATATGTATA
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1 (1>793)	→	GCGGTATAAGAATATATACTGATATGTATA
E04-Y94913-G0266763-1_pET-seqF.ab1 (1>794)	←	GCGGTATAAGAATATATACTGATATGTATA

		17501760177
		CCCGAAGTATGTCAAAAAGAGGTATGCTAT
G0266763-1.seq(1>2469)	→	CCCGAAGTATGTGTCAAAAAGAGGTATGCTAT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	CCCGAAGTATGTGTCAAAAAGAGGTATGCTAT
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	CCCGAAGTATGTGTCAAAAAGAGGTATGCTAT

		17801790180
		GAAGCAGCGTATTACAGTGACAGTTGACAG
G0266763-1.seq(1>2469)	→	GAAGCAGCGTATTACAGTGACAGTTGACAG
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	GAAGCAGCGTATTACAGTGACAGTTGACAG
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	GAAGCAGCGTATTACAGTGACAGTTGACAG

		18101820183
		CGACAGCTATCAGTTGCTCAAGGCATATAT
G0266763-1.seq(1>2469)	→	CGACAGCTATCAGTTGCTCAAGGCATATAT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	CGACAGCTATCAGTTGCTCAAGGCATATAT
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	CGACAGCTATCAGTTGCTCAAGGCATATAT
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	T

		18401850186
		GATGTCAATATCTCCGGTCTGGTAAGCACA
G0266763-1.seq(1>2469)	→	GATGTCAATATCTCCGGTCTGGTAAGCACA
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	GATGTCAATATCTCCGGTCTGGTAAGCACA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	GATGTCAATATCTCCGGTCTGGTAAGCACA
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	GATGTCAATATCTCCGGTCTGGTAAGCACA

		18701880189
		ACCATGCAGAATGAAGCCCGTCGTCTGCGT
G0266763-1.seq(1>2469)	→	ACCATGCAGAATGAAGCCCGTCGTCTGCGT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	ACCATGCAGAATGAAGCCCGTCGTCTGCGT
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	ACCATGCAGAATGAAGCCCGTCGTCTGCGT
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	ACCATGCAGAATGAAGCCCGTCGTCTGCGT

		19001910192
		GCCGAACGCTGGAAAGCGGAAAATCAGGAA
G0266763-1.seq(1>2469)	→	GCCGAACGCTGGAAAGCGGAAAATCAGGAA
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	GCCGAACGCTGGAAAGCGGAAAATCAGGAA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	GCCGAACGCTGGAAAGCGGAAAATCAGGAA
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	GCCGAACGCTGGAAAGCGGAAAATCAGGAA

		19301940195
		GGGATGGCTGAGGTCGCCCCGTTTATTGAA
G0266763-1.seq(1>2469)	→	GGGATGGCTGAGGTCGCCCCGTTTATTGAA
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	GGGATGGCTGAGGTCGCCCCGTTTATTGAA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	GGGATGGCTGAGGTCGCCCCGTTTATTGAA
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	GGGATGGCTGAGGTCGCCCCGTTTATTGAA

		19601970198
		ATGAACGGCTCTTTTGCTGACGAGAACAGG
G0266763-1.seq(1>2469)	→	ATGAACGGCTCTTTTGCTGACGAGAACAGG
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	ATGAACGGCTCTTTTGCTGACGAGAACAGG
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	ATGAACGGCTCTTTTGCTGACGAGAACAGG
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	ATGAACGGCTCTTTTGCTGACGAGAACAGG



Project: G0266763-1.SQD Contig 1

		<div><div></div><div>19902000201</div></div>
		GGCTGGTGAAATGCAGTTTAAGGTTTACAC
G0266763-1.seq(1>2469)	→	GGCTGGTGAAATGCAGTTTAAGGTTTACAC
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	GGCTGGTGAAATGCAGTTTAAGGTTTACAC
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	GGCTGGTGAAATGCAGTTTAAGGTTTACAC
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	GGCTGGTGAAATGCAGTTTAAGGTTTACAC
		<div><div></div><div>20202030204</div></div>
		CTATAAAAGAGAGAGCCGTTATCGTCTGTT
G0266763-1.seq(1>2469)	→	CTATAAAAGAGAGAGCCGTTATCGTCTGTT
E11-Y94913-g0266763-1_g0266763-1-seq3.ab1(1>793)	→	CTATAAAA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	CTATAAAAGAGAGAGCCGTTATCGTCTGTT
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	CTATAAAAGAGAGAGCCGTTATCGTCTGTT
		<div><div></div><div>20502060207</div></div>
		TGTGGATGTACAGAGTGATATTATTGACAC
G0266763-1.seq(1>2469)	→	TGTGGATGTACAGAGTGATATTATTGACAC
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	TGTGGATGTACAGAGTGATATTATTGACAC
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	TGTGGATGTACAGAGTGATATTATTGACAC
		<div><div></div><div>20802090210</div></div>
		GCCCCGGGCGACGGATGGTGATCCCCCTGGC
G0266763-1.seq(1>2469)	→	GCCCCGGGCGACGGATGGTGATCCCCCTGGC
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	GCCCCGGGCGACGGATGGTGATCCCCCTGGC
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	GCCCCGGGCGACGGATGGTGATCCCCCTGGC
		<div><div></div><div>21102120213</div></div>
		CAGTGCACGTCTGCTGTCAGATAAAGTCTC
G0266763-1.seq(1>2469)	→	CAGTGCACGTCTGCTGTCAGATAAAGTCTC
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	CAGTGCACGTCTGCTGTCAGATAAAGTCTC
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	CAGTGCACGTCTGCTGTCAGATAAAGTCTC
		<div><div></div><div>21402150216</div></div>
		CCGTGAACTTTACCCGGTGGTGCATATCGG
G0266763-1.seq(1>2469)	→	CCGTGAACTTTACCCGGTGGTGCATATCGG
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	CCGTGAACTTTACCCGGTGGTGCATATCGG
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	CCGTGAACTTTACCCGGTGGTGCATATCGG
		<div><div></div><div>21702180219</div></div>
		GGATGAAAGCTGGCGCATGATGACCACCGA
G0266763-1.seq(1>2469)	→	GGATGAAAGCTGGCGCATGATGACCACCGA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	GGATGAAAGCTGGCGCATGATGACCACCGA
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	GGATGAAAGCTGGCGCATGATGACCACCGA
		<div><div></div><div>22002210222</div></div>
		TATGGCCAGTGTGCCGGTTTCCGTTATCGG
G0266763-1.seq(1>2469)	→	TATGGCCAGTGTGCCGGTTTCCGTTATCGG
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←	TATGGCCAGTGTGCCGGTTTCCGTTATCGG
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→	TATGGCCAGTGTGCCGGTTTCCGTTATCGG



## Project: G0266763-1.SQD Contig 1

			22302240225
			GGAAGAAGTGGCTGATCTCAGCCACCGCGA
G0266763-1.seq(1>2469)	→		GGAAGAAGTGGCTGATCTCAGCCACCGCGA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		GGAAGAAGTGGCTGATCTCAGCCACCGCGA
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		GGAAGAAGTGGCTGATCTCAGCCACCGCGA
			22602270228
			AAATGACATCAAAAACGCCATTAACCTGAT
G0266763-1.seq(1>2469)	→		AAATGACATCAAAAACGCCATTAACCTGAT
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		AAATGACATCAAAAACGCCATTAACCTGAT
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		AAATGACATCAAAAACGCCATTAACCTGAT
			22902300231
			GTTCTGGGGAATATAAATGTCAGGCTCCCT
G0266763-1.seq(1>2469)	→		GTTCTGGGGAATATAAATGTCAGGCTCCCT
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		GTTCTGGGGAATATAAATGTCAGGCTCCCT
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		GTTCTGGGGAATATAAATGTCAGGCTCCCT
			23202330234
			TATACACAGCCAGTCTGCAGGTCGACCATA
G0266763-1.seq(1>2469)	→		TATACACAGCCAGTCTGCAGGTCGACCATA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		TATACACAGCCAGTCTGCAGGTCGACCATA
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		TATACACAGCCAGTCTGCAGGTCGACCATA
			23502360237
			GTGACTGGATATGTTGTGTTTTACAGTATT
G0266763-1.seq(1>2469)	→		GTGACTGGATATGTTGTGTTTTACAGTATT
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		GTGACTGGATATGTTGTGTTTTACAGTATT
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		GTGACTGGATATGTTGTGTTTTACAGTATT
			23802390240
			ATGTAGTCTGTTTTTTATGCAAATCTAAT
G0266763-1.seq(1>2469)	→		ATGTAGTCTGTTTTTTATGCAAATCTAAT
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		ATGTAGTCTGTTTTTTATGCAAATCTAAT
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		ATGTAGTCTGTTTTTTATGCAAATCTAAT
			24102420243
			TTAATATATTGATATTTATATCATTTTACG
G0266763-1.seq(1>2469)	→		TTAATATATTGATATTTATATCATTTTACG
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		TTAATATATTGATATTTATATCATTTTACG
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		TTAATATATTGATATTTATATCATTTTACG
			24402450246
			TTTCTCGTTCAGCTTTCTTGACAAAGTGG
G0266763-1.seq(1>2469)	→		TTTCTCGTTCAGCTTTCTTGACAAAGTGG
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		TTTCTCGTTCAGCTTTCTTGACAAAGTGG
E03-Y94913-G0266763-1_G0266763-1-SEQ4.ab1(1>795)	→		TTTCTCGTTCAGCTTTCTTGACAAAGTGG
			24702480249
			TTCATCACCACCATCATCACTAAGTGATTA
G0266763-1.seq(1>2469)	→		TTCATCACCACCATCATCACTAAGTGATTA
E04-Y94913-G0266763-1_pET-seqF.ab1(1>794)	←		TTCATCACCACCATCATCACTAAGTGA

E03-Y94913-G0266763-1\_G0266763-1-SEQ4.ab1 (1>795) → TTCATCACCACCATCATCACTAAGTGATTA

G0266763-1.seq(1>2469) → ACCTAGG

E03-Y94913-G0266763-1\_G0266763-1-SEQ4.ab1 (1>795) → ACCTAGGCTGCTGCCACCGCTGAGCAATAA

E03-Y94913-G0266763-1\_G0266763-1-SEQ4.ab1 (1>795) → CTAGCATAACCCCTTGGGGCCTCTAAACGG

E03-Y94913-G0266763-1\_G0266763-1-SEQ4.ab1 (1>795) → GTCTTGAGGGGTTTTTTGCTGAAAGGAGGA

E03-Y94913-G0266763-1\_G0266763-1-SEQ4.ab1 (1>795) → ACTATATCCGATTGGCGAATGGGACGCGC

E03-Y94913-G0266763-1\_G0266763-1-SEQ4.ab1 (1>795) → CCTGTAGCGGCGCA