

AN ABSTRACT OF THE THESIS OF

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Abstract approved: _____

Beta-lactam antibiotics are among the most widely prescribed antibiotics in clinical settings. As a result, it is not uncommon to encounter clinically isolated bacteria that produce beta-lactamase as a means of resistance to these antibiotics. Among the many beta-lactamases currently known, TEM-type beta-lactamase is the most widespread beta-lactamase observed in clinical settings. However, less is known regarding its prevalence among bacteria in natural settings. To determine how widespread the TEM-type beta-lactamase gene is in the environment, experiments were designed to detect this and other beta-lactamase genes from environmentally isolated bacteria. Briefly, soil was collected along the banks of the Niangua River located in west-central Missouri. Diluted soil preparations were plated onto tryptic soy agar plates containing ampicillin and allowed to incubate at room temperature for up to 14 days. Colonies ($n = 58$) growing on the plates were selected and total DNA isolated from each strain. Using 7 groups of beta-lactamase specific DNA primers, the polymerase chain reaction was employed to amplify beta-lactamase genes, if present. In total, 58 isolates were positive for an 800 base pair amplicon indicative of a TEM-type beta-lactamase. DNA sequencing of the

amplified fragment from each strain was sequenced to identify the TEM variant present. The data indicated that TEM-116 beta-lactamases are widespread among natural populations of bacteria from this location. In total, 56 of the genes were TEM-116, 1 was TEM-1, and 1 was TEM-118. These genes were most predominate in the genera *Bacillus* and *Pseudomonas*. Attempts to transfer the resistance genes via transformation and conjugation were unsuccessful.

BETA-LACTAMASE GENES IN SOIL BACTERIA ISOLATED FROM THE BANKS
OF THE NIANGUA RIVER IN MISSOURI

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PREFACE

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Introduction

Antibiotics are medicinal products that inhibit bacterial growth or kill infectious organisms. Beta-lactam antibiotics are among the most widely prescribed antibiotics in clinical settings and include the penicillins, cephalosporins, carbapenems, and monobactams (26). While slightly distinct in their mechanism of action, they all contain a beta-lactam ring structure which inhibits bacterial cell wall biosynthesis. Indeed, 60 percent of all currently used antibiotics belong to the beta-lactam family (48), with the penicillins and cephalosporins predominating (23).

In clinical settings, many infectious organisms have developed resistance to antibiotics because of inappropriate use (17). Oftentimes, bacterial resistance to beta-lactam antibiotics is due to the synthesis of a beta-lactamase, an enzyme that attacks and destroys the beta-lactam ring (25). The constant overuse of beta-lactam antibiotics has caused mutation, resulting in more than 1000 different beta-lactamases being described (22). As a result, it is not uncommon to encounter clinically isolated bacteria that produce beta-lactamase as a means of resistance to these antibiotics.

Beta-lactamases can be divided into several types based on their functional activity and molecular structure (12), with TEM, SHV, CTX-M, and OXA predominating. Among these, TEM beta-lactamases are the most widespread in clinical settings (2, 11). The first TEM beta-lactamase described, TEM-1, was isolated from a penicillin-resistant bacterium in the 1960s (16). To date, 208 TEM-type variants have been reported (see <http://www.lahey.org/Studies/temtable.asp>; May, 2013).

TEM variants differ in amino acid sequence because of genetic mutations, which change their antibiotic resistance spectrum (8, 13). Figure 1 shows the distribution of known mutations in amino acid sequence of TEM-1 beta-lactamase, leading to the different variants. Among all of the 208 beta-lactamases, TEM-10, TEM-12, and TEM-26 are the most common in the United States (10, 28, 37). Clinically, TEM-type beta-lactamases predominate in *Escherichia coli* and *Klebsiella pneumoniae* (42). Because many beta-lactamase genes are plasmid encoded (49), this aids mobilization, leading to increased distribution among other bacterial species (24).

While clinical research regarding antibiotic resistance is on-going, antibiotic resistance research in the natural environment has been overlooked. Although most resistance determinants have been found in either clinical or laboratory settings, antibiotic resistance genes may come directly from the environment (4). A changing environment likely facilitates the emergence of new resistance genes and their transfer to humans. Thus, understanding antibiotic resistance in natural environments may help predict and possibly counteract the emergence and future evolution of antibiotic resistance in clinical settings (32, 43).

To aid our understanding of environmental antibiotic resistance, a study was initiated with the objective of determining how widespread beta-lactamase genes are in the environment. To address this question, soil was collected along the banks of the Niangua River located in west-central Missouri. Ampicillin resistant bacteria were isolated and screened for the presence of beta-lactamase genes. Our data suggests that

K		5	10	14	24	27	28	42	46	47	48	49	61	68	71	72	85	86	92	93	97	101	114	121	132	136	144	145	148	151	152	154	155	156	157	D										
H		11	16	27	28	29	43	61	75	107	109	115	118	132	134	147	V	D	S	V	M	R	S	131	137	I	G	108	V	58	K	173	V	G												
H		12	17	28	29	43	61	75	107	109	115	118	132	134	147	V	D	S	V	M	R	S	131	137	I	G	108	V	58	K	173	V	G													
D		13	18	29	30	44	62	76	108	110	116	122	136	138	151	V	D	S	V	M	R	S	131	137	I	G	108	V	58	K	173	V	G													
D		14	19	30	31	45	63	77	112	114	120	126	149	151	164	V	D	S	V	M	R	S	131	137	I	G	108	V	58	K	173	V	G													
L		15	20	31	32	46	64	78	116	118	124	130	153	155	168	V	D	S	V	M	R	S	131	137	I	G	108	V	58	K	173	V	G													
L		16	21	32	33	47	65	79	117	119	125	131	154	156	169	V	D	S	V	M	R	S	131	137	I	G	108	V	58	K	173	V	G													
TEM-1		Q	F	L	E	K	D	D	Q	L	A	L	M	V	V	G	N	L	E	T	D	R	S	P	H	M	D	H	D	R	W	I	N	D	M	A	T	G	R	A	G	L	A	G	A	H
TEM-1		Q	F	L	E	K	D	D	Q	L	A	L	M	V	V	G	N	L	E	T	D	R	S	P	H	M	D	H	D	R	W	I	N	D	M	A	T	G	R	A	G	L	A	G	A	H
K		4	9	141	152	2	42	60	63	116	157	3	76	59	21	145	164	164	173	175	179	182	184	198	213	218	224	224	237	238	244	244	265	265	275	275	279	T	164	164	163					
K		5	10	142	153	3	43	61	64	117	158	4	77	60	22	146	165	165	174	176	180	183	197	212	217	221	221	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		6	11	143	154	3	44	62	65	118	159	5	78	61	23	147	166	166	175	177	181	184	198	213	218	224	224	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		7	12	144	155	3	45	63	66	119	160	6	79	62	24	148	167	167	176	178	182	185	199	214	219	225	225	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		8	13	145	156	3	46	64	67	120	161	7	80	63	25	149	168	168	177	179	183	186	199	215	220	226	226	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		9	14	146	157	3	47	65	68	121	162	8	81	64	26	150	169	169	178	180	184	187	201	216	221	227	227	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		10	15	147	158	3	48	66	69	122	163	9	82	65	27	151	170	170	179	181	185	188	202	217	222	228	228	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		11	16	148	159	3	49	67	70	123	164	10	83	66	28	152	171	171	180	182	186	189	203	218	223	229	229	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		12	17	149	160	3	50	68	71	124	165	11	84	67	29	153	172	172	181	183	187	190	204	219	224	230	230	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		13	18	150	161	3	51	69	72	125	166	12	85	68	30	154	173	173	182	184	188	191	205	220	225	231	231	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		14	19	151	162	3	52	70	73	126	167	13	86	69	31	155	174	174	183	185	189	192	206	221	226	232	232	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		15	20	152	163	3	53	71	74	127	168	14	87	70	32	156	175	175	184	186	190	193	207	222	227	233	233	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		16	21	153	164	3	54	72	75	128	169	15	88	71	33	157	176	176	185	187	191	194	208	223	228	234	234	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		17	22	154	165	3	55	73	76	129	170	16	89	72	34	158	177	177	186	188	192	195	209	224	229	235	235	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		18	23	155	166	3	56	74	77	130	171	17	90	73	35	159	178	178	187	189	193	196	210	225	230	236	236	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		19	24	156	167	3	57	75	78	131	172	18	91	74	36	160	179	179	188	190	194	197	211	226	231	237	237	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		20	25	157	168	3	58	76	79	132	173	19	92	75	37	161	180	180	189	191	195	198	212	227	232	238	238	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		21	26	158	169	3	59	77	80	133	174	20	93	76	38	162	181	181	190	192	196	199	213	228	233	239	239	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		22	27	159	170	3	60	78	81	134	175	21	94	77	39	163	182	182	191	193	197	200	214	229	234	240	240	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		23	28	160	171	3	61	79	82	135	176	22	95	78	40	164	183	183	192	194	198	201	215	230	235	241	241	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		24	29	161	172	3	62	80	83	136	177	23	96	79	41	165	184	184	193	195	199	202	216	231	236	242	242	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		25	30	162	173	3	63	81	84	137	178	24	97	80	42	166	185	185	194	196	199	203	217	232	237	243	243	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		26	31	163	174	3	64	82	85	138	179	25	98	81	43	167	186	186	195	197	199	204	218	233	238	244	244	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		27	32	164	175	3	65	83	86	139	180	26	99	82	44	168	187	187	196	198	199	205	219	234	239	245	245	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		28	33	165	176	3	66	84	87	140	181	27	100	83	45	169	188	188	197	199	200	205	219	235	240	246	246	237	238	244	244	265	265	275	275	279	T	164	164	163						
K		29	34	166	177	3	67	85	88	141	182	28	101	84	46	170	189	189	198	199	201	206	220	236	242	242	237	238	244	244	265	265	275	275	279	T	164	164	163							
K		30	35	167	178	3	68	86	89	142	183	29	102	85	47	171	190	190	199	200	202	207	221	237	243	243	237	238	244	244	265	265	275	275	279	T	164	164	163							
K		31	36	168	179	3	69	87	90	143	184	30	103	86	48	172	191	191	199	200	203	208	222	238	244	244	237	238	244	244	265	265	275	275	279	T	164	164	163							
K		32	37	169	180	3	70	88	91	144	185	31	104	87	49	173	192	192	199	201	204	209	223	239	245	245	237	238	244	244	265	265	275	275	279	T	164	164	163							
K		33	38	170	181	3	71	89	92	145	186	32	105	88	50	174	193	193	199	202	205	210	224	240	246	246	237	2																		

Figure 1. Positions and Types of Amino Acid Substitutions in TEM Type Beta-Lactamases (44). This figure illustrates 177 different variants. Since its publication, the number of TEM variants has risen to 208.

TEM type beta-lactamases, specifically TEM-116, are widespread among natural populations of both Gram positive and Gram negative bacteria from this location.

Materials and Methods

Sample Collection

Soil samples were collected along the banks of the Niangua River, located in west-central Missouri, in August 2011. Samples were stored at 4°C for 1 week until processing.

Isolation of Ampicillin Resistant Bacterial Strains

Soil samples were suspended in 10 ml of phosphate buffered saline (PBS). Vortexing was used to mix the sample. Serial 1:10 dilutions were accomplished using PBS until a final dilution of 1:100 was obtained. 100 µl of the 1:100 diluted sample was spread plated onto Luria-Bertani (10 g tryptone, 10 g yeast extract, 5 g NaCl/L) agar (15 g/L) containing 100 µg/ml ampicillin. Plates were incubated at 30 °C for up to 14 days and representative colonies (n = 58) were picked from each plate for further analysis.

Isolation of Total DNA and Plasmid DNA

Bacterial isolates were propagated in 5 ml LB at 37°C with shaking (200 rpm) overnight. Cell pellets obtained after centrifugation in a microcentrifuge at full speed were resuspended in 1ml of TE (10mM Tris, 1mM EDTA; pH8) containing 10 mg/ml lysozyme. After incubation at 37 °C for 30 min, total DNA was obtained using a Zymo Research Quick-gDNA™ MicroPrep kit (Zymo Research, Irvine, CA) according to the manufacturer's instructions. Plasmid DNA was obtained from each isolate using a Zymo Research Plasmid MiniPrep kit (Zymo Research, Irvine, CA) according to the manufacturer's instructions.

Multiplex Polymerase Chain Reaction

DNA from each ampicillin resistant isolate was used as a template to amplify beta-lactamase genes in a multiplex PCR according to the methodology described by Dallenne et al. (15). Primers and amplification conditions are outlined in Table 1. Briefly, reaction mixtures were composed of 100 ng of total DNA, multiplex primers, dNTP's, Phusion High-Fidelity DNA Polymerase, and its buffer (Thermo Scientific, Rockford, IL) in a total volume of 100 µL. Amplification was carried out on a T100 Thermal Cycler (Bio-Rad, Hercules, CA).

16S rRNA Polymerase Chain Reaction

DNA from each ampicillin resistant isolate was used as a template to amplify the 16S rRNA gene (21). Primers consisted of the 8F primer (5'AGAGTTGATCCTGGCTCAG3') and 1492R(L) primer (5'GGTTACCTT GTTACGACTT3'). Reaction mixtures were composed of 100 ng of total DNA, 8F and 1492R(L) primers, 200 µM of dNTP's, 1 U of Phusion High-Fidelity DNA Polymerase (Thermo Scientific) and its buffer in a total volume of 50 µL. Amplification was carried out on a T100 Thermal Cycler (Bio-Rad, Hercules, CA) using the following conditions: initial denaturation at 94 °C for 5 min; 30 cycles of 94 °C for 1 min, 50 °C for 1 min and 72 °C for 1.5 min; and a final elongation step at 72 °C for 7 min.

Table 1. Primers and Amplifications Conditions Used in this Study.*

PCR name	Beta-Lactamase targeted	Primer name	Sequence (5'→3')	Amplicon Size (bp)	Amplification Conditions	
Multiplex I TEM, SHV and OXA-1-like	TEM variants including TEM-1 and TEM-2	MultiTSO-T_for	CATTTCGGTGTGCCCTTATT	800	94 °C for 10 min; 30 cycles of 94 °C for 40 s, 60 °C for 40 s and 72 °C for 1 min; 72 °C for 7 min.	
		MultiTSO-T_rev	CGTTCATCCATAGTTGCCTGAC			
	SHV variants including SHV-1	MultiTSO-S_for	AGCCGCTTGAGCAAATTAAAC	713		
		MultiTSO-S_rev	ATCCCGCAGATAAATCACAC			
	OXA-1, OXA-4 and OXA-30	MultiTSO-O_for	GGCACCAAGATTCAACTTCAAG	564		
		MultiTSO-O_rev	GACCCCAAGTTCCCTGTAAGTG			
Multiplex II CTX-M group 1, group 2 and group 9	Variants of CTX-M group 1 including CTX-M-1, CTX-M-3 and CTX-M-15	MultiCTXMGp1_for	TTAGGAARTGTGCCGCTGYA**	688	94 °C for 10 min; 30 cycles of 94 °C for 40 s, 60 °C for 40 s and 72 °C for 1 min; 72 °C for 7 min.	
		MultiCTXMGp1-2_rev	CGATATCGTTGGTGGTRCCAT**			
	Variants of CTX-M group 2 including CTX-M-2	MultiCTXMGp2_for	CGTTAACGGCACGATGAC	404		
		MultiCTXMGp1-2_rev	CGATATCGTTGGTGGTRCCAT**			
	Variants of CTX-M group 9 including CTX-M-9 and CTX-M-14	MultiCTXMGp9_for	TCAAGCCTGCCGATCTGGT	561		
		MultiCTXMGp9_rev	TGATTCTGCCGCTGAAG			
CTX-M group 8/25	CTX-M-8, CTX-M-25, CTX-M-26 and CTX-M-39 to CTX-M-41	CTX-Mg8/25_for	AACRCRCAGACGCTCTAC**	326	94 °C for 10 min; 30 cycles of 94 °C for 40 s, 60 °C for 40 s and 72 °C for 1 min; 72 °C for 7 min.	
		CTX-Mg8/25_rev	TCGAGCCGGAASGTGTYAT**			
Multiplex III ACC, FOX, MOX, DHA, CIT and EBC	ACC-1 and ACC-2	MultiCaseACC_for	CACCTCCAGCGACTTGTAC	346	94 °C for 10 min; 30 cycles of 94 °C for 40 s, 60 °C for 40 s and 72 °C for 1 min; 72 °C for 7 min.	
		MultiCaseACC_rev	GTTAGCCAGCATCACGATCC			
	FOX-1 to FOX-5	MultiCaseFOX_for	CTACAGTGCAGGTGGTT	162		
		MultiCaseFOX_rev	CTATTGCGGCCAGGTGA			
	MOX-1, MOX-2, CMY-1, CMY-8 to CMY-11 and CMY-19	MultiCaseMOX_for	GCAACAAACGACAATCCATCCT	895		
		MultiCaseMOX_rev	GGGATAGGCAGTAACCTCCCCAA			
	DHA-1 and DHA-2	MultiCaseDHA_for	TGATGGCACAGCAGGATATT	997		
		MultiCaseDHA_rev	GCTTGACTCTTCGGTATT			

	LAT-1 to LAT-3, BIL-1, CMY-2 to CMY-7, CMY-12 to CMY-18 and CMY-21 to CMY-23	MultiCaseCIT_for	CGAAGAGGCAATGACCAGAC	538		
		MultiCaseCIT_rev	ACGGACAGGGTTAGGATAGY**			
	ACT-1 and MIR-1	MultiCaseEBC_for	CGGTAAAGCCATGTTGCG	683		
		MultiCaseEBC_rev	AGCCTAACCCCTGATACA			
Multiplex IV VEB, PER and GES	GES-1 to GES-9 and GES-11	MultiGES_for	AGTCGGCTAGACCGGAAAG	399		
		MultiGES_rev	TTTGTCCGTGCTCAGGAT			
	PER-1 and PER-3	MultiPER_for	GCTCCGATAATGAAAGCGT	520		
		MultiPER_rev	TTCGGCTTGACTCGGCTGA			
	VEB-1 to VEB-6	MultiVEB_for	CATTTCGGATGCAAAGCGT	648		
		MultiVEB_rev	CGAAGTTCTTGACTCTG			
Multiplex V GES and OXA-48-like	GES-1 to GES-9 and GES-11	MultiGES_for	AGTCGGCTAGACCGGAAAG	399	94 °C for 10 min; 30 cycles of 94 °C for 40 s, 57 °C for 40 s and 72 °C for 1 min; 72 °C for 7 min.	
		MultiGES_rev	TTTGTCCGTGCTCAGGAT			
	OXA-48-like	MultiOXA-48_for	GCTTGATGCCCTCGATT	281		
		MultiOXA-48_rev	GATTGCTCCGTGGCCGAAA			
Multiplex VI IMP, VIM and KPC	IMP variants except IMP-9, IMP-16, IMP-18, IMP-22 and IMP-25	MultiIMP_for	TTGACACTCCATTACDG**	139	94 °C for 10 min; 30 cycles of 94 °C for 40 s, 55 °C for 40 s and 72 °C for 1 min; 72 °C for 7 min.	
		MultiIMP_rev	GATYGAGAATTAAGGCCACYCT**			
	VIM variants including VIM-1 and VIM-2	MultiVIM_for	GATGGTGTGCTGCATA	390		
		MultiVIM_rev	CGAATGCGCAGCACCAAG			
	KPC-1 to KPC-5	MultiKPC_for	CATTCAAGGGCTTCTTGCTGC	538		
		MultiKPC_rev	ACGACGGCATAGTCATTGCG			

*(Dallenne et al., 2010)

**Y=T or C; R=A or G; S=G or C; D=A or G or T.

Agarose Gel Electrophoresis

To effectively separate DNA fragments of various sizes, agarose gel electrophoresis was employed according to standard conditions (46). Briefly, 30 ml of 1X TAE buffer prepared from a 50X TAE stock (242 g Tris, 57.1 ml acetic acid, and 4 ml 0.5 M EDTA in 1 L of H₂O), 0.3 g of agarose, and 2 µl of 10 mg/ml ethidium bromide (EtBr) were mixed and heated in the microwave until the agarose dissolved. The mixture was poured into a gel casting tray containing a gel comb. After solidification, the comb was removed and 1X TAE buffer was added to completely cover the gel. Subsequently, DNA samples were mixed with loading buffer, loaded into the wells, and electrophoresed using a Bio-Rad model 250/2.5 power supply (Bio-Rad), at ~100 volts for 30-40 min. DNA was visualized after electrophoresis using a UV Intensity Transilluminator (Fisher Scientific™; St. Louis, MO).

Purification of DNA from Agarose Gels

DNA was excised from electrophoresed gels using a razor blade and purified using a Zymoclean™ Gel DNA Recovery Kit (Zymo Research) according to the manufacturer's instructions.

Determination of DNA Concentration

DNA concentration was measured using a nanodrop spectrophotometer (Thermo Scientific, Rockford, IL). Briefly, according to the manufacturer's instructions, after blanking the instrument using sterile water at $\lambda=260$ nm, the concentration of the DNA sample was determined.

DNA Sequencing and BLAST Analysis

Purified DNA amplicons were submitted to the DNA Sequencing Facility at the University of Arkansas for Medical Sciences (Dept. Microbiology/Immunology, Little Rock, AR) for DNA sequence determination. The Basic Local Alignment Search Tool (BLAST) (5) was used to analyze the sequenced DNA.

Preparation of Competent Cells

Escherichia coli TG1 was grown in 2X LB medium at 30 °C overnight with shaking at 250 rpm. Subsequently, 0.5 ml of overnight culture was used to inoculate 200 ml of fresh 2X LB in a flask at 30 °C with shaking. When the OD₆₀₀ of the culture reached 0.3, 4 ml of 1M MgCl₂ was added and incubation continued until an OD₆₀₀ of 0.45-0.55 was obtained. The culture was placed on ice for 2 hrs prior to centrifugation at 3000 rpm at 4 °C for 5 min in a J2-HS centrifuge (Beckham Coulter Inc; Brea, CA). The cell pellet was resuspended in 100 mM CaCl₂ media (0.05 M CaCl₂, 0.04 M MnCl₂, and 0.02 M CH₃COON, 4°C, pH7.5). After incubation on ice for 40 min, cells were pelleted again by centrifugation and resuspended in fresh ice-cold 100 mM CaCl₂ media containing 15% glycerol. Competent cells were stored in 100 µl aliquots at -80 °C.

Transformation

Transformation was used to transfer ampicillin resistance into competent *E. coli* TG1. Briefly, 100 ng of plasmid DNA isolated from an ampicillin resistant isolate was mixed with 100 µl competent *Escherichia coli* TG1 cells and put on ice for 15 min. Subsequently, incubation at 42 °C for 90 s followed by ice for 60 s was performed. 900

μ l LB liquid media was added to each culture and after incubation for 45 min at 37 °C, 100 μ l was spread plated onto a LB/ampicillin (100 μ g/ml) agar plate and incubated overnight at 37 °C.

Conjugation

Conjugation was used in an attempt to transfer ampicillin resistance from an ampicillin resistant isolate to recipient *Escherichia coli* HWS174. Briefly, environmental isolates (donor) and *Escherichia coli* HWS174 (recipient) were grown overnight at 37 °C with shaking at 200 rpm. A 50 μ l aliquot was transferred to fresh LB media and incubation continued until a 0.5 McFarland turbidity standard was obtained. 500 μ l of the donor strain was mixed with 500 μ l of the recipient strain followed by incubation at 37 °C for 90 min. Subsequently, 100 μ l of the conjugated culture was plated on LB agar containing ampicillin (100 μ g/ml) and rifampicin (100 μ g/ml) followed by incubation overnight at 37°C.

Results

Amplification of Beta-lactamase Genes via Multiplex PCR

Using seven multiplex PCRs (15), total DNA isolated from 58 different ampicillin resistant bacterial strains was analyzed for the presence of beta-lactamase genes as outlined in Table 2. The TEM multiplex reaction yielded an amplification product in all 58 strains (Figure 2). All other multiplex reactions were negative (data not shown).

DNA Sequencing and BLAST Analysis of TEM-type Beta-Lactamase Genes

The TEM-type beta-lactamase gene amplified from all 58 strains was subjected to DNA sequencing. Each gene was sequenced three times in both directions. Data obtained was subjected to the Basic Local Alignment Search Tool (BLAST) (5) to identify each TEM-type beta-lactamase gene. Our data indicated 56 of the genes were TEM-116, 1 was TEM-1, and 1 was TEM-118. This data is summarized in Table 2 and Appendix 1.

Amplification of 16S rRNA Gene

Using primers 8F and 1492R (L) (21), the 16S rRNA gene was amplified from chromosomal DNA templates prepared from the 58 ampicillin resistant strains via PCR. Subsequently, DNA sequencing followed by BLAST analysis were used to make a tentative identification at the genus level. This data is summarized in Table 2. The DNA sequence of the 16S rRNA gene used for analysis is presented in Appendix 2.

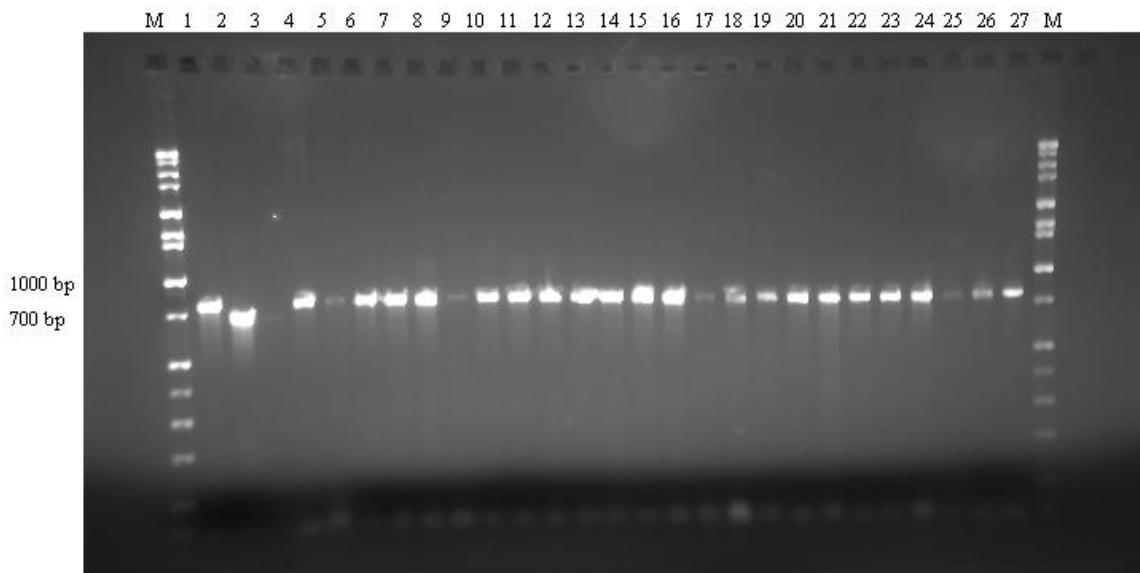


Figure 2. Representative Agarose Gel Showing Amplification of the TEM-type Beta-Lactamase Gene. DNA was amplified using Multiplex I TEM, SHV and OXA-1-like primers. M - DNA Marker, Lane 1, 2 - Positive control (1. *Escherichia coli* 5667, TEM positive; 2. *Klebsiella pneumoniae* 5585, SHV positive), Lane 3 - Blank, Lane 4-27 - Total DNA from bacteria isolated in this study. Gel shown is representative of 3 additional agarose gels containing amplified TEM - type beta-lactamases from the remaining bacteria isolated in this study.

Table 2. Summary of BLAST Analysis Using DNA Sequencing from 16S rRNA and TEM-type Beta-Lactamase Genes.

Isolate	Strain Name	Genus	GenBank	Identity	TEM-
1	NRCJ-1	<i>Pseudomonas</i>	NR_025228.1	98%	116
2	NRCJ-2	<i>Pseudomonas</i>	NR_025228.1	98%	116
3	NRCJ-3	<i>Pseudomonas</i>	NR_025228.1	98%	116
4	NRCJ-4	<i>Stenotrophomonas</i>	NR_028930.1	99%	116
5	NRCJ-5	<i>Pseudomonas</i>	NR_043314.1	99%	116
6	NRCJ-6	<i>Nocardoides</i>	NR_040938.1	96%	116
7	NRCJ-7	<i>Pseudomonas</i>	NR_043314.1	99%	116
8	NRCJ-8	<i>Pseudomonas</i>	NR_028906.1	99%	116
9	NRCJ-9	<i>Pseudomonas</i>	NR_024918.1	99%	116
10	NRCJ-10	<i>Stenotrophomonas</i>	NR_028930.1	99%	116
11	NRCJ-11	<i>Curtobacterium</i>	NR_025467.1	97%	181
12	NRCJ-12	<i>Curtobacterium</i>	NR_042315.1	95%	116
13	NRCJ-13	<i>Pseudomonas</i>	NR_041953.1	97%	116
14	NRCJ-14	<i>Pseudomonas</i>	NR_074834.1	99%	1
15	NRCJ-15	<i>Pseudomonas</i>	NR_074834.1	95%	116
16	NRCJ-16	<i>Pseudomonas</i>	NR_041953.1	97%	116
17	NRCJ-17	<i>Bacillus</i>	NR_043403.1	99%	116
18	NRCJ-18	<i>Pseudomonas</i>	NR_025227.1	84%	116
19	NRCJ-19	<i>Pseudomonas</i>	NR_028906.1	99%	116
20	NRCJ-20	<i>Stenotrophomonas</i>	NR_074875.1	99%	116
21	NRCJ-21	<i>Pseudomonas</i>	NR_040859.1	99%	116
22	NRCJ-22	<i>Stenotrophomonas</i>	NR_041577.1	86%	116
23	NRCJ-23	<i>Pseudomonas</i>	NR_041953.1	99%	116
24	NRCJ-24	<i>Pseudomonas</i>	NR_041953.1	99%	116
25	NRCJ-25	<i>Pseudomonas</i>	NR_028906.1	97%	116
26	NRCJ-26	<i>Pseudomonas</i>	NR_024902.1	94%	116
27	NRCJ-27	<i>Pseudomonas</i>	NR_024662.1	99%	116
28	NRCJ-28	<i>Pseudomonas</i>	NR_025228.1	94%	116
29	NRCJ-29	<i>Pseudomonas</i>	NR_028906.1	99%	116
30	NRCJ-30	<i>Lysobacter</i>	NR_036925.1	94%	116
31	NRCJ-31	<i>Bacillus</i>	NR_043403.1	88%	116
32	NRCJ-32	<i>Bacillus</i>	NR_043403.1	97%	116
33	NRCJ-33	<i>Bacillus</i>	NR_043403.1	98%	116
34	NRCJ-34	<i>Bacillus</i>	NR_043403.1	99%	116
35	NRCJ-35	<i>Lysinibacillus</i>	NR_042072.1	95%	116
36	NRCJ-36	<i>Bacillus</i>	NR_043403.1	98%	116
37	NRCJ-37	<i>Bacillus</i>	NR_043403.1	98%	116
38	NRCJ-38	<i>Bacillus</i>	NR_043403.1	98%	116
39	NRCJ-39	<i>Pseudomonas</i>	NR_041953.1	99%	116

40	NRCJ-40	<i>Sphingobacterium</i>	NR_040953.1	96%	116
41	NRCJ-41	<i>Pseudomonas</i>	NR_041953.1	99%	116
42	NRCJ-42	<i>Cupriavidus</i>	NR_028766.1	88%	116
43	NRCJ-43	<i>Pseudomonas</i>	NR_043935.1	99%	116
44	NRCJ-44	<i>Pseudomonas</i>	NR_025228.1	84%	116
45	NRCJ-45	<i>Bacillus</i>	NR_043403.1	99%	116
46	NRCJ-46	<i>Isoptericola</i>	NR_025439.1	99%	116
47	NRCJ-47	<i>Bacillus</i>	NR_043403.1	97%	116
48	NRCJ-48	<i>Pseudomonas</i>	NR_025881.1	95%	116
49	NRCJ-49	<i>Pseudomonas</i>	NR_024662.1	99%	116
50	NRCJ-50	<i>Pseudomonas</i>	NR_024662.1	99%	116
51	NRCJ-51	<i>Bacillus</i>	NR_043403.1	97%	116
52	NRCJ-52	<i>Stenotrophomonas</i>	NR_041577.1	99%	116
53	NRCJ-53	<i>Pseudomonas</i>	NR_026532.1	98%	116
54	NRCJ-54	<i>Pseudomonas</i>	NR_025228.1	99%	116
55	NRCJ-55	<i>Curtobacterium</i>	NR_042315.1	97%	116
56	NRCJ-56	<i>Pseudomonas</i>	NR_024662.1	98%	116
57	NRCJ-57	<i>Shigella</i>	NR_026331.1	99%	116
58	NRCJ-58	<i>Pseudomonas</i>	NR_028906.1	98%	116

*Isolates harboring TEM-1 and TEM-118 genes are in bold.

Discussion

Since antibiotic usage is commonplace throughout all aspects of human society, it is inevitable that residual amounts will be present in the environment and continuously work as selective agents for the development of environmental antibiotic resistance (31).

The first antibiotic was found in 1928. In the 80 plus years since, numerous antibiotics have been discovered for which resistance soon followed. To aid our understanding of environmental antibiotic resistance, it is important to determine what type of resistance is present. As a step toward this goal, this study has focused on beta-lactamase resistance.

Ampicillin is a beta-lactam antibiotic belonging to the penicillin family and is the successor of penicillin, one of the more predominating antibiotics in today's world. Utilizing bacteriological media containing ampicillin, 58 bacterial isolates were obtained from soil located along the banks of the Niangua River in west-central Missouri. Beta-lactamases are produced at high concentrations by certain Gram-positive organisms, and in particular *Bacillus cereus* (30), *Bacillus licheniformis* (7) and *Staphylococcus aureus* (34). The enzyme is also widely distributed in Gram-negative organisms and many different variations of beta-lactamases can be recognized on the grounds of activity towards different beta-lactams (33). The molecular classification of beta-lactamases is based on the amino acid sequence of these enzymes. To date, four classes are recognized (A-D), correlating with a functional classification. Classes A, C, and D act by a serine-based mechanism, whereas class B (also called metallo-beta-lactamases) need zinc for their action (6).

Resistance to beta-lactam antibiotics is an increasing problem, commonly due to the production of beta-lactamases (20). To investigate the mechanism of resistance observed in the ampicillin resistant isolates obtained in this study, it was hypothesized that beta-lactamase genes could be detected. Following the assay described by Dallenne et al. (15), seven groups of primers were utilized to amplify the most frequently encountered beta-lactamases genes. All 58 isolates were positive for TEM type beta-lactamases genes (Figure 2).

Several TEM type beta-lactamases have been identified in laboratory studies before their identification in clinical isolates (9, 51). Figure 1 shows all the amino acid substitutions in TEM type beta-lactamase genes. One or more amino acid substitutions around the active site result in extended-spectrum beta-lactamases, which demonstrate broad resistance to penicillins, aztreonam, and cephalosporins (38). Salverda et al. (45) overviewed the frequency of substitutions found at different positions within the TEM-1 amino acid sequence of clinical and laboratory isolates. The results demonstrated not all of the substitutions necessarily confer increased resistance and some substitutions, such as Alanine to Valine at position 184, are frequent in the laboratory but not currently present in clinical isolates. Investigating beta-lactamases in the environment or laboratory has direct clinical relevance because they may reveal novel phenotypes that could arise in clinical isolates. In this study, three types of TEM type beta-lactamases genes were identified; TEM-116 ($n = 56$), TEM-1 ($n = 1$), and TEM-181 ($n = 1$). Table 3 shows the amino acid positions of the substitutions among these three types of beta-lactamase.

TEM-1 beta-lactamase is the most prevalent beta-lactamase found in Gram-negative bacteria and is capable of hydrolyzing both penicillin and cephalosporin (6). Indeed, up to 90% of all ampicillin resistance noted in *E. coli* is due to the production of TEM-1 (14). TEM-116 was first and most prevalently identified from clinical isolates of *Klebsiella pneumoniae* and *Escherichia coli* from a Korean nationwide survey (29). The amino acid replacement at position 84 and 184 have not been observed in other TEM type beta-lactamases. In a study of resistance analyzing extended-spectrum cephalosporins in clinical *Escherichia coli* isolates from companion animals in the United States, Shaheen et al. (47) analyzed the structural difference between wild-type TEM-1 and TEM-181 beta-lactamases. DNA sequence analysis results suggested there might be a scheme for *in vivo* evolution: from TEM-1 to TEM-181, and from TEM-181 to TEM-116. The current study did not attempt to address this issue. (Table 3).

Many resistance genes are carried on transposable elements, such as transposons, integrons, and plasmids, and can be transferred to other bacteria (1, 39, 40, 50) through horizontal gene transfer. Murray and Mederski-Samaroj (35) reported penicillin resistance and beta-lactamase activity could be transferred by conjugation at a high frequency to an enterococcal laboratory recipient strain, demonstrating the emergence of plasmid-mediated penicillin resistance in the genus *Streptococcus*. Elwell et al. (24) also showed that R plasmids encoding beta-lactamase activity contained about 40% of all transposable ampicillin resistance. Recently, Dierikx et al. (24) tested the transfer of resistance plasmids by conjugation and transformation experiments and found IncI1

Table 3. Amino Acid Substitutions among the Three Types of Beta-Lactamase Genes Identified in this Study.

β -Lactamase	Residue (coding triplet) at amino acid	
	84	184
TEM-1	Val (GTA)	Ala (GCA)
TEM-116	Ile (ATT)	Val (GTA)
TEM-181	Val (GTA)	Val (GTA)

plasmids containing blaCTX-M-1 can spread in *E. coli* and *Salmonella enterica* isolates.

Doi et al. (19) confirmed the contribution of horizontal gene transmission of beta-lactamase genes among *E. coli* and *K. pneumonia*.

To determine the genetic location of the TEM beta-lactamase resistance in the 58 ampicillin resistant isolates from this study, transformation and conjugation experiments were performed. For conjugation, the recipient cell was *E. coil* HWS174 (rifampicin resistance), while the donor cell (environmental isolates) is resistant to ampicillin. After co-incubation of the donor and recipient, if beta-lactamase genes are located on a conjugatable plasmid of the donor cells, it will be transferred to *E. coil* HWS174, which is noted by its growth on bacteriological media containing both rifampicin and ampicillin. However, no colonies were found on the plates, suggesting beta-lactamase genes are not located on the conjugatable plasmid of the donor cells. Alternatively, plasmids with TEM genes may have been transferred, but the beta-lactamase genes were not expressed. Control experiments demonstrated *E. coil* HWS174 was well suited as a recipient cell in conjugation assays (data not shown).

For transformation experiments, competent *E. coli* TG1 served as the recipient strain and isolated plasmid DNA from each ampicillin resistant bacterium was used as transforming DNA. If beta-lactamase genes are located on the plasmid of the donor cells and can be transformed into *E. coli*, colonies will form on ampicillin containing medium. No colonies were found. As with the conjugation studies, no attempt was made to determine if plasmids were transferred but TEM genes not expressed. Control

experiments showed competent *E. coli* TG1 was a suitable recipient cell in transformation experiments (data not shown). Taken together, the data suggest a chromosomal location of the TEM genes or a lack of plasmid gene expression. If the TEM genes are not expressed, other mechanisms may account for the beta-lactam resistance, such as efflux pumps (36); inactivation of the antibiotic itself (41), inhibition of antibiotic uptake (3) or down-regulation of the cell target molecule (27).

In conclusion, beta-lactam resistance is widespread among soil bacteria from the banks of the Niangua River. All 58 organisms resistant to ampicillin contained TEM-type beta-lactamases, with TEM-116 predominating from the genera *Bacillus* and *Pseudomonas*. Attempts to transfer the resistance genes via transformation and conjugation were unsuccessful. This study highlights the presence of antibiotic resistance in the environment without direct antibiotic exposure and reinforces the need for continued surveillance of antibiotic resistance in natural settings.

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Appendix 1

DNA Sequence of TEM-type Beta-Lactamase Genes Identified in This Study

(5' → 3')

TEM-1

CATTCCGTGTCGCCCTTATTCCCTTTGCGGCATTTGCCTCCTGTTTG
 CTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGC
 ACGAGTGGGTTACATCGAACCTGGATCTCAACAGCGGTAAGATCCTGAGAGTT
 TTCGCCCCGAAGAACGTTTCCAATGATGAGCAGCTTAAAGTTCTGCTATGTG
 GCGCGGTATTATCCCCTGTTGACGCCGGCAAGAGCAACTCGGTCGCCGCATA
 CACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTCACAGAAAAGCATCT
 TACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAACCATGAGTG
 ATAACACTGCGGCCAACTTACTTCTGACAACGATCGGAGGACCAGAAGGAGCT
 AACCGCTTTTGACAAACATGGGGATCATGTAACTCGCCTGATCGTTGGG
 AACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCCACGATGCC
 TGCAGCAATGGCAACAAACGTTGCGCAAACATTAACTGGCGAACTACTTACTC
 TAGCTTCCGGCAACAATTAAATAGACTGGATGGAGGCGGATAAAGTTGCAGGA
 CCACTCTCGCCTGCCCTCCGGCTGGCTGGTTATTGCTGATAATCTGGA
 GCCGGTAGCGTGGGTCTCGCGGTATCATTGCACTGGGCCAGATGGTAA
 GCCCTCCGTATCGTAGTTACACGACGGGAGTCAGGCAACTATGGATG
 AACG

TEM-116

CATTCCGTGTCGCCCTTATTCCCTTTGCGGCATTTGCCTCCTGTTTG
 CTCACCCAGAAACGCTGGTGAAAGTAAAAGATGCTGAAGATCAGTTGGGTGC
 ACGAGTGGGTTACATCGAACCTGGATCTCAACAGCGGTAAGATCCTGAGAGTT
 TTCGCCCCGAAGAACGTTTCCAATGATGAGCAGCTTAAAGTTCTGCTATGTG
 GCGCGGTATTATCCCCTGTTGACGCCGGCAAGAGCAACTCGGTCGCCGCATA
 CACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTCACAGAAAAGCATCT
 TACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAACCATGAGTG
 ATAACACTGCGGCCAACTTACTTCTGACAACGATCGGAGGACCAGAAGGAGCT
 AACCGCTTTTGACAAACATGGGGATCATGTAACTCGCCTGATCGTTGGG
 AACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCCACGATGCC
 TGTAGCAATGGCAACAAACGTTGCGCAAACATTAACTGGCGAACTACTTACTC
 TAGCTTCCGGCAACAATTAAATAGACTGGATGGAGGCGGATAAAGTTGCAGGA
 CCACTCTCGCCTGCCCTCCGGCTGGTTATTGCTGATAATCTGGA

GCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAA
GCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATG
AACG

TEM-181

CATTCCCGTGTGCCCTTATTCCCTTTGCGGCATTTGCCTCCTGTTTG
CTCACCCAGAAACGCTGGTGAAGTAAAAGATGCTGAAGATCAGTTGGGTGC
ACGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTT
TTCGCCCCGAAGAACGTTTCCAATGATGAGCACTTTAAAGTTCTGCTATGTG
GCGCGGTATTATCCCGTGTGACGCCGGCAAGAGCAACTCGGTCGCCGCATA
CACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTCACAGAAAAGCATCT
TACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAACCATGAGTG
ATAACACTGCGGCCAACTTACTTCTGACAACGATCGGAGGACCGAAGGAGCT
AACCGCTTTTGACAAACATGGGGATCATGTAACTCGCCCTGATCGTTGGG
AACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCCACGATGCC
TGTAGCAATGGCAACAAACGTTGCGCAAACATTAACTGGCGAACTACTTACTC
TAGCTTCCCGCAACAATTAAATAGACTGGATGGAGGCGGATAAAGTTGCAGGA
CCACTTCTGCGCTCGGCCCTCCGGCTGGCTGGTTATTGCTGATAAATCTGGA
GCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAA
GCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATG
AACG

Appendix 2

16S rRNA Gene Sequences (5'→3') Used for BLAST Analysis

Strain NRCJ-1

TCGAGCGGATGAAAGGAGCTTGCTCCTGGATTCA
 ATGCCTAGGAATCTGCCTGGTAGTGGGGACAACGTTGAAAGGAACGCTA
 ATACCGCATACGTCCCTACGGGAGAAAGCAGGGACCTCGGGCCTGCGCTA
 TCAGATGAGCCTAGGTGGATTAGCTAGTTGGTAGGTAATGGCTACCAAG
 GCGACGATCCGTAACTGGTCTGAGAGGATGATCAGTCACACTGGAAC TGAGA
 CACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATATTGGACAATGGC
 GAAAGCCTGATCCAGCCATGCCCGTGTGAAGAAGGTCTCGGATTGTA
 AGCACTTAAGTGGAGGAAGGGTTGTAGATTAATACTCTGCAATTGAC
 GTTACCGACAGAATAAGCACCGGCTAACCTGTGCCAGCAGCCCGTAATA
 CAGAGGGTGCAAGCGTTAACCGAATTACTGGCGTAAAGCGCGCGTAGGTG
 GTTGTAAAGTGGATGTGAAATCCCCGGGCTAACCTGGAACTGCATCCA
 AAACTGGCAAGCTAGAGTATGGTAGAGGGTGGAAATTCCCTGTGTAGCGG
 TGAAATGCGTAGATATAGGAAGGAACACCAGTGGCGAAAGCGACCACTGC
 ACTGATACTGACACTGATGTGCAAAGCGTGGGAGCAAACAGGATTAGATA
 CCCTGGTAGTCCAGGCCGTAACACTATATCAACTAACCTTGGAGCCTGAAC
 TATTATTGGGCCAGCTCAGCAAATCATTACTTCCCGGGAGGGAAAGACC
 GCAGGGGGAGGCTCAAATGTATATTAAAGTGGCCGGCGCACCCACGCC
 CTGCAGCTTGTCTAAACTACGAGCAAACCGAAAAACCCCTCGAGGACAG
 ACGAACCTAAAGACATTCCATTAGTGGTTGTGCACTTGAGAAAAATT
 AACACAGTCGCACAGTGGACTT

Strain NRCJ-2

AGGCCTAACACATGCAAGTCGAGCGGATGAAAGGAGCTTGCTCCTGGATTCA
 CGGGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGACAAC
 GTTCGAAAGGAACGCTAACACCGCATACGTCCCTACGGGAGAAAGCAGGG
 ACCTTCGGGCCTGCGCTATCAGATGAGCCTAGGTGGATTAGCTAGTTGGT
 AGGTAATGGCTACCAAGGCAGCGATCCGTAACGGTCTGAGAGGATGATCA
 GTCACACTGGAACGTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTGG
 GGAATATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCCGTGTGAA
 GAAGGTCTCGGATTGTAAAGCACTTAAGTGGAGGAAGGGTTGTAGATT

AATACTCTGCAATTGACGTTACCGACAGAATAAGCACC GGCTAACTCTGTG
 CCAGCAGCCGCGGTAAATACAGAGGGT GCAAGCGTTAAC TCGGAATTACTGGC
 GTAAAGCGCGCTAGGTGGTTGTTAACAGTTGGATGTGAAATCCCCGGGCTCA
 ACCTGGGAACTGCATCCAAA ACTGGCAAGCTAGAGTATGGTAGAGGGGGGG
 CGGAA

Strain NRCJ-3

GCAGGCCTAACACATGCAAGTCGAGCGGATGAAAGGAGCTTGCTCCTGGATT
 CAGCGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGGACA
 ACGTTCGAAAGGAACGCTAATACCGCATACGCTCACGGGAGAAAGCAGGG
 GACCTTCGGGCCTGCGCTATCAGATGAGCCTAGGTGGATTAGCTAGTTGGT
 GAGGTAATGGCTACCAAGGCAGATCCGTAACTGGTCTGAGAGGATGATC
 AGTCACACTGGAACTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTG
 GGGAAATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCCGTGTGTA
 AGAAGGTCTCGGATTGTAAGCACTTTAACGGAGGAAGGGTTGTAGAT
 TAATACTCTGCAATTGACGTTACCGACAGAATAAGCACC GGCTAACTCTGT
 GCCAGCAGCCGCGTAATACAGAGGGT GCAAGCGTTAAC TCGGAATTACTGGG
 CGTAAAGCGCGCTAGGTGGTTGTTAACAGTTGGATGTGAAATCCCCGGGCTC
 AACCTGGGAACTGCATCCAAA ACTGGCAAGCTAGAGTATGGTAGAGGGTGGT
 GGAATT CCTGTGTAGCGGTGAAATGCCTAGATATAGGAAGGAACACCCTGT
 GGCCCAACGCCACCACCTGGACTGATTCTGACACTGCCGTGCCAAAGCGTGG
 GTAGCAATCAGGCTTAGATCCCCGGGCTGCTACCCCTCCATCCAACCCAA
 CTACCG

Strain NRCJ-4

TCGAACGGCAGCACAGTAAGAGCTT GCTCTTATGGGTGGCGAGTGGCGGACG
 GGTGAGGAATACATCGGAATCTACCTT CGTGGGGATAACGTAGGGAAAC
 TTACGCTAATACCGCATACGACCTCGGGT GAAAGCAGGGACCTTCGGGCC
 TTGCGCGGATAGATGAGCCGATGTCGGATTAGCTAGTTGGCGGGGTAAAGGC
 CCACCAAGGCAGCGATCCGTAGCTGGTCTGAGAGGATGATCAGCCACACTGG
 AACTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATATTGG
 ACAATGGCGCAAGCCTGATCCAGCCATACCGCGTGGGTGAAGAAGGCCTTC
 GGGTTGTAAGCCCTTTGTTGGAAAGAAAAGCAGTCGGCTAACCTCGTGCCAGCAGCC
 TGTTCTGACGGTACCCAAAGAATAAGCACC GGCTAACCTCGTGCCAGCAGCC

GCGGTAATACGAAGGGTGCAAGCGTTACTCGGAATTACTGGGCGTAAAGCGT
 GCGTAGGTGGTTTTAAGTCTGTTGAAAGCCCTGGGCTAACCTGGGAAT
 TGCAGTGGATACTGGGCGACTAGAGTGTGGTAGAGGGTAGTGGAAATTCCCGG
 TGTAGCAGTGAAATGCGTAGAGATCGGGAGGAACATCCATGGCGAAAGGCA
 GCTACCTGGACCAACACTGACACTGAGGCACGAAAGCGTGGGAGCAAACA
 GGATTAGATACCCTGGTAGTCCACGCCCTAAAACGATGCGAActGGATGTTG
 GGTGCAATTGGCACGCA

Strain NRCJ-5

TACACCGTGGTAACCGTCCTCCGAAGGTTAGACTAGCTACTTCTGGTGCAC
 CCACTCCCATGGTGTGACGGGCGGTGTACAAGGCCGGAACGTATTAC
 CGCGACATTCTGATTCGCGATTACTAGCGATTCCGACTTCACGCAGTCGAGTT
 GCAGACTGCGATCCGGACTACGATCGGTTATGGGATTAGCTCCACCTCGCG
 GCTTGGCAACCCTTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGTA
 AGGGCCATGATGACTTGACGTACCCCCACCTCCTCCGGTTGTCACCGGCA
 GTCTCCTTAGAGTGCCCACCATAACGTGCTGGTAACTAAGGACAAGGGTTGC
 GCTCGTTACGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCCA
 TGCAGCACCTGTCTCAATGTTCCGAAGGCACCAATCCATCTCTGGAAAGTTC
 ATTGGATGTCAAGGCCTGGTAAGGTTCTCGCGTTGCTCGAATTAAACCACA
 TGCTCCACCGCTTGTGCGGGCCCCGTCAATTGAGTTAACCTTGCG
 GCCGTACTCCCCAGGCGGTCAACTTAATGCGTTAGCTGCGCCACTAAGAGCT
 CAAGGCTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGGTA
 TCTAATCCTGTTGCTCCCCACGCTTCGCACCTCAGTGTCACTGAGTCCA
 GGTGGTCGCCCTCGCCACTGGTGTCCCTATATCTACGCATTCAACCGCTA
 CACAGGAAATTCCACCACCCCTTACCATACTCTAGCTGCCAGTTGGATGC
 AGTTCCCAGGTTGAGCCCCGGGATTTCACATCCAACCTAACAAACCACCTAC
 GCGCGCTTACGCCAGTAATTCCGATTAACGCTTGACCCCTGTATTACC
 GCGGCTGCTGGCACAGAGTTAGCCCG

Strain NRCJ-6

TGCTTAAACATGCAAGTCGAGCGGAAAGGCTCCTCGGGGGTACTCGAGCGG
 CGAACGGGTGAGTAACACGTGAGTAATCTGCCCTGTGCTCTGGGATAGCCAC
 CGGAAACGGTGATTAATACCGGATATGACTGCTACCCGCATGGGTTGGTGGT
 GGAAAGTTTCGGCGCAGGATGTGCTCGCGGCCATCAGCTGTTGGTGGAG

GTAATGGCTCACCAAGGCTTGACGGGTAGCCGCCCTGAGAGGGGTGACCGGT
 CACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGG
 AATATTGGACAATGGCGGAAGCCTGATCCAGCAACGCCGCGTGAGGGATG
 ACGGCCTCGGGTTGTAACCTCTTCAGTACCGACGAAGCGAAAGTGACGG
 TAGGTACAGAAGAAGGACCGGCCAACTACGTGCCAGCAGCCGCGTAATAC
 GTAGGGTCCGAGCGTTGCCGAATTATTGGCGTAAAGGGCTCGTAGGC
 TTTGTCGCGTCGGGAGTGAAAACAGCCAGCTTAACCTGTTGCTGCTTCGAT
 ACGGGCAGACTAGAGGTATGCAGGGGAGAATGGAATTCTCTGGTAGCGGT
 GAAAATGCGCAGATATCAGGAGGAACACCGTGGCGAAGGCGTTCTCTGG
 GCATTACCTGACGCTGAGGAGCGAAAGTGTGGGGAGCGAACAAAGATTATAT
 ACCCTGGTAGTCCACACCGTAAACGTTGGCGCTAGGTGTGGGACCAAATC
 CATTGGGTTCCGTGCCGTACTAACACAAATAACCACCCCCCCTGGGAGTG
 CAGCGGCGGGCTAAAAGAATAGCAGGGGAC

Strain NRCJ-7

CACCGTGGTAACCGTCCTCCGAAGGTTAGACTAGCTACTTCTGGTGCAACCC
 ACTCCCATGGTGTGACGGCGGTGTACAAGGCCGGAACGTATTACCG
 CGACATTCTGATTGCGATTACTAGCGATTCCGACTTCACGCAGTCGAGTTGC
 AGACTGCGATCCGGACTACGATCGTTTATGGGATTAGCTCCACCTCGCGGC
 TTGGCAACCCTTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGTAAAG
 GGCCATGATGACTTGACGTCATCCCCACCTCCTCCGGTTGTACCGGCAGT
 CTCCCTAGAGTGCCCACCATAACGTGCTGGTAACTAAGGACAAGGGTGC
 TCGTTACGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCCATG
 CAGCACCTGTCTCAATGTTCCGAAGGCACCAATCCATCTGGAAAGTTCAT
 TGGATGTCAAGGCCTGGTAAGGTTCTCGCGTTGCTTCGAATTAAACCACATG
 CTCCACCGCTTGTGCGGGCCCCCGTCAATTGAGTTAACCTTGC
 CGTACTCCCCAGGCAGGTCAACTTAATGCGTTAGCTGCCACTAAGAGCTCA
 AGGCTCCAAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGGTATC
 TAATCCTGTTGCTCCCCACGCTTCGCACCTCAGTGTCACTAGTCCAGG
 TGGTCGCCCTCGCCACTGGTGTCCCTATCTACGCATTACCGCTACA
 CAGGAAATTCCACCAACCCTTACCATACCTAGCTGCCAGTTGGATGCA
 GTTCCCAGGTTGAGCCCAGGATTTCACATCCAACCTAACAAACCACCTACGC
 GCGCTTACGCCAGTAATTCCGATTAACGCTTGCACCCCTTGTATTACCG

Strain NRCJ-8

GCAGGCCTAACACATGCAAGTCGAGCGGCAGCACGGTACTTGTACCTGGTG
 GCGAGCGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGG
 ATAACGCTCGAAACGGACGCTAATACCGCATACTGCCTACGGGAGAAAGC
 AGGGGACCTTCGGGCCTGCGCTATCAGATGAGCCTAGGTCGGATTAGCTAG
 TTGGTGAGGTAATGGCTACCAAGGCACGATCCGTAACGGTCTGAGAGGA
 TGATCAGTCACACTGGAACGTGAGACACGGTCCAGACTCCTACGGGAGGCAGC
 AGTGGGAATATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCCGTGT
 GTGAAGAAGGTCTCGGATTGTAAAGCACTTAAGTTGGGAGGAAGGGCATT
 TACCTAATACGTAAGTGTGTTGACGTTACCGACAGAATAAGCACCGGCTAAC
 TCTGTGCCAGCAGCCCGGTAATACAGAGGGTGCAAGCGTTAACGGGATT
 CTGGCGTAAAGCGCGTAGGTGGTCGTTAAGTTGGATGTGAAATCCCCG
 GGCTCAACCTGGGAACGTGATTCAAAACTGTCGAGCTAGAGTATGGTAGAGG
 GTGGTGGGAATTTCTGGGTTCCCAGGGGAAATTCTCATCTTGAAAGGG
 GAAAACCTTGAGGGAAGTCAAACCCGGTCCCACCCCGCCTCCCACCG
 CCAAAGCCAGGTGAACCCCAAGTGGTCACATCCCCTGTTGGTCTCCACTCCC
 AAATCTACCCATTACCCGGTGCACGTAACGGGAAATCGCCTCATATA
 ATGCATGAGATCATCTTGGGGGGAGTCCCAAGGAGGCAAAGCTTATATCA
 TTTGATGGGCCCCCAACCGCGGGACCTGGGCATAATTGAAAGCACCC
 CGAAGACCCCCCTGAGTCTGGCCGCCAGCGGACCTACAAGAAGTGGGATT
 GTTGCCTGGGGAAACTAAAAACAAGGTTGCAGGAATGGAGGAACAAAC
 CCCCTCCGTTCCCCAGGTAAAGTAGGGGTTAAGTTCCCCAAAAAAACGCT
 GCAACCCCAATCTCCCCTCAGATTG

Strain NRCJ-9

ATCACCGTGGTAACCGTCTCCGAAGGTTAGACTAGCTACTTCTGGTGAAC
 CCACTCCATGGTGTGACGGCGGTGTACAAGGCCGGAACGTATTCAC
 CGCGACATTCTGATTGCGATTACTAGCGATTCCGACTTCACGCAGTCGAGTT
 GCAGACTGCGATCCGGACTACGATCGTTATGGGATTAGCTCCACCTCGCG
 GCTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGT
 AAGGCCATGACTGACGTACCCACCTTCCCGGTTGTACCG
 AGTCTCCTAGAGTGCACCGATTACGTGCTGGTAACTAAGGACAAGGGTTG
 CGCTCGTTACGGACTAACCCAACATCTCACGACACGAGCTGACGACAGCC
 ATGCAGCACCTGTCTCAATGTTCCGAAGGCACCAATCCATCTGGAAAGTT

CATTGGATGTCAAGGCCTGGTAAGGTTCTCGCGTGCTTCGAATTAAACCAC
 ATGCTCCACCGCTTGTGCGGGCCCCGTCAATTGAGTTAACCTTGC
 GGCGTACTCCCCAGGCAGGTCAACTTAATGCGTAGCTGCCACTAAGAGC
 TCAAGGCTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGGT
 ATCTAATCCTGTTGCTCCCCACGCTTCGCACCTCAGTGTCACTACGTC
 AGGTGGTCGCCACTGGTGTCCCTATATCTACGCATTACCGCT
 ACACAGGAAATTCCACCAACCCCTTACCATACTCTAGCTTCAGTTGAATG
 CAGTTCCCAGGTTGAGCCGGGGCTTCACATCCAACTTAACAAACCACCTA
 CGCGCGCTTACGCCAGTAATTCCGATTAACGCTTGACCCCTCT

Strain NRCJ-10

CCGCACACCGTGGCAAGCGCCCTCCGAAGGTTAACGCTACCTGCTCTGGTG
 CAACAAACTCCCAGGTGTGACGGCGGTGTACAAGGCCGGAACGTAT
 TCACCGCAGCAATGCTGATCTGCGATTACTAGCGATTCCGACTTCATGGAGTC
 GAGTTGCAGACTCCAATCCGGACTGAGATAGGGTTCTGGGATTGGCTTGCC
 CTCGCGGGTTGCAGCCCTGTCCCTACCATTGTAGTACGTGTAGCCCTG
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 CCGCGGGCTCCTTAGAGTTCCCACCATTACGTGCTGGCAACTAAGGACAAG
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 TCTTGCACCGTACTCCCCAGGCGGAACCTAACGCGTAGCTCGATAACTG
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 CAGGGTATCTAATCCTGTTGCTCCCCACGCTTCGTGCCTCAGTGTCACTGTT
 GGTCCAGGTAGCTGCCTTCGCATGGATGTTCTCCGATCTACGCATTTC
 ACTGCTACACCGGGAAATTCCACTACCCTCTACCAACTCTAGTCGCCAGTAT
 CCACTGCAATTCCCAGGTTGAGCCCAGGGCTTCACAACAGACTAAACAAAC
 CACCTACGCACGCTTACGCCAGTAATTCCGAGTAACGCTTGACCCCTCGT
 ATTACCGCGGCTGCTGGCACGAAGTTAGCCC

Strain NRCJ-11

CTGCGGCGTGCTTAACACATGCAAGTCGAACGATGATGCCAGCTGCTGGG
 TGGATTAGTGGCGAACGGGTGAGTAACACGTGAGTAACCTGCCCTGACTCT

GGGATAAGCGTTGGAAACGACGTCTAATACTGGATATGATGCCGCCGCAT
 GGTCTGGTGGTGGAAAGATTGGTGGGATGGACTCGCGGCCTATCAG
 CTTGTTGGTGGAGGTAATGGCTACCAAGGCGACGACGGTAGCCGGCTGAG
 AGGGTGACCGGCCACACTGGACTGAGACACGGCCCAGACTCCTACGGGAG
 GCAGCAGTGGGAATATTGCACAATGGCGAAAGCCTGATGCAGCAACGCC
 CGGTGAGGGATGACGGCCTCGGGTTGAAACCTCTTAGTAGGGAAAGAAG
 CGAAAGTACGGTACCTGCAGAAAAAGCACCGGCTAACTACGTGCCAGCAG
 CCGCGGTAATACGTAGGGTGCAAGCGTTGTCCGGAATTATTGGCGTAAAGA
 GCTCGTAGGCCTTGTCGCTGCTGTGAAATCCCCAGGCTAACATCTCG
 CTTGCAGTGGGTACGCGCGGACTATAGTCGGCGGGGAGATTATTGGAAT
 TTCCTGGAGGAA

Strain NRCJ-12

CGAACCTCGCGTGCTTAAACATGCAAGTCGAACGATGATGCCAGCTTGC
 TGGGTGGATTAGTGGCGAACGGGTGAGTAACACGTGAGTAACCTGCCCTGA
 CTCTGGATAAGCGTTGGAAACGACGTCTAATACTGGATATGATCACCGGCC
 GCATGGTCTGGTGGAAAGATTGGTGGGATGGACTCGCGGCCTAT
 CAGCTTGGTGGAGGTAATGGCTACCAAGGCGACGACGGTAGCCGGCCT
 GAGAGGGTGACCGGCCACACTGGACTGACACACGGCCCAGACTCCTACGG
 GAGGCAGCAGTGGGAATATTGCACAATGGCGAAAGCCTGATGCAGCAAC
 GCCCGTGAGGGATGACGGCCTCGGGTTGAAACCTCTTAGTAGGGAAAG
 AAGCGAAAGTACGGTACCTGCAGAAAAAGCACCGGCTAACTACGTGCCAG
 CAGCCCGGTAATACGTAGGGTGCAAGCGTTGTCCGGAATTATTGGCGTAA
 AGAGCTCGTAGGCCCTTGTGCGCTGCTGTGAAATCCGAGGCTAACCTC
 GGGCTGCAGTGGTACGGCAGACTAGAGTGCAGGAGGGAGATTGGAATT
 CCTGGTGTAGCGGTGGAAATGCGCAGATATCATGAGGAACACCGATGGCGAAG
 GCAGATCTCTGGCGTAACTGACGCTGAGGAGCGAAAGCGTGGGAGCGA
 ACAGGATTAGATAACCTGGTAGTCCACGCCGAAACGTTGGCGCTAGATGT
 AGGGACCTTCCACGGTTCTGTGCGTAGCTAACGCATTAAGCGCCCCGCCT
 GGGAGTACGCCGAAAGCTAAACTCAAAGGAATGACGGGCCGACAACCG
 GCGGAGCATGCGATTAATTGATGCAACCGAAGAACCTTACAAAGCTTGAC
 ATACACCCGAAACGCTAGAGATGGTCGCCCCCTGTGTCGGGTACAGGTGTGC
 ATGGTGTGTCCTGCTGGATGTTAGGTAAGTCCACACAGCCAACGCTCG
 TTCATATTGCCAGCCCTAGCAGGACTCATAGAAACTACAGGTCACTCGGAG
 AAGTGGGATGAGTGAACACGACCACTAGCTGGCTTAGCAGCTACCAGGCC

GTCAAGGGTCGCGTAACCGTAGTAAGCGATAGCGAGAGAACCGAGTCAATCG
ATAGAGTTCTGGACACTATCACTCATAGCTACACCTG

Strain NRCJ-13

AGAGAGCTTGCTCTGATTCA CGGGGGACGGGTGAGTAATGCCTAGGAAT
CTGCCTGGTAGTGGGGACAACGTCTCGAAAGGGACGCTAATACCGCATACG
TCCTACGGGAGAAAGCAGGGACCTCAGGGCTTGCCTGCGCTATCAGATGAGCCT
AGGTCGGATTAGCTAGTTGGTGAGGTAATGGCTACCAAGGCAGCAGATCCGT
AACTGGTCTGAGAGGATGATCAGTCACACTGGAAC TGAGACACGGTCCAGAC
TCCTACGGGAGGCAGCAGTGGGAATATTGGACAATGGCGAAAGCCTGATC
CAGCCATGCCCGTGTGAAGAAGGTCTCGGATTGTAAAGCACTTTAAGT
TGGGAGGAAGGGTTGTAGATTAATACTCTGCAATTGACGTTACCGACAGA
ATAAGCACCGGCTAACTCTGTGCCAGCAGCCCGGTAATACAGAGGGTGCAA
GCGTTAACCGAACATTACTGGCGTAAAGCGCGCTAGGTGGTTGTTAAGTT
GGATGTGAAAGCCCCGGCTAACCTGGGAACTGCATTCAAAACTGACAAGC
TAGAGTATGGTAGAGGGTGGTGAATTCTGTGAGCGGTGAAAATGCGTA
GATATAGGAAGGGACACCAGTGGCGAAGGCGACCACCTGGCTGATACTGA
CACTGAGGTGCGAACGTGGGGAGCAAACACGAGTAGATACCCCTGGTAGT
CGACGCCGCATACGATGTCAACTAGCCGTTGGGGCCTGAGCTCTTATTGGC
GCCCTCACCACTTAATTGATAGCCTGTGGGAGCGGCCGGCGGTGGAA
CACACACGTATTAATTGCGCGCGCACAAACACTGGGCGAGTCAGTGTATT
TAATGCAACGCACGAACAAGAACAGACTGGCCTC

Strain NRCJ-14

TACGATAACACCGTGGTAACCGTCTCCGAAGGTTAGACTAGCTACTTCTGGT
GCAACCCACTCCATGGTGTGACGGGCGGTGTACAAGGCCGGAACGTA
TTCACCGCGACATTCTGATTGCGATTACTAGCGATTCCGACTCACGAGTC
GAGTTGCAGACTGCGATCCGGACTACGATCGTTTCTGGATTAGCTCCACC
TCGCGGCTGGCAACCCCTGTACCGACCATTGTAGCACGTGTAGCCCAG
GCCGTAAGGCCATGACTGACGTCATCCCCACCTCCTCCGGTTGTCA
CCGGCAGTCTCCTTAGAGTGCCCACCATTACGTGCTGGTAACTAAGGACAAG
GGTTCGCTCGTTACGGACTTAACCCAACATCTCACGACACGAGCTGACGA
CAGCCATGCAGCACCTGTCTCAATGTTCCGAAGGCACCAATCCATCTGG
AAAGTTCATGGATGTCAAGGCCTGGTAAGGTTCTCGCGTTGCTCGAATTA

AACCACATGCTCCACCGCTTGTGCGGGCCCCGTCAATTGAGTTAA
 CCTTGCAGGCCGTACTCCCCAGGCGGTCAACTTAATGCGTTAGCTGCGCCACTA
 AGAGCTCAAGGCTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACC
 AGGGTATCTAATCCTGTTGCTCCCCACGCTTCGACCTCAGTGTCACTATC
 AGTCCCAGGTGGTCGCCTCGCCACTGGTGTTCCTCCTATATCTACGCATTTC
 ACCGCTACACAGGAAATTCCACCACCCCTTACCATACTCTAGCTGACAGTTT
 TGAATGCAGTTCCCAGTTGAGCCGGGATTCACATCCAACCTAACGAACC
 ACCTACGCGCGCTTACGCCAGTAATTCCGATTAACGCTTGACCCCTGTAA
 TTACCGCGGCTGCTGGCAC

Strain NRCJ-15

TCACCGTGGTAACCGCCTCCGAAGGTTAGACTAGCTACTTCTGGTGCAACC
 CACTCCCATGGTGTGACGGCGGTGTACAAGGCCGGAACGTATTCA
 GCGACATTCTGATTGCGATTACTAGCGATTCCGACTTCACGCAGTCGAGTTG
 CAGACTGCGATCCGGACTACGATCGGTTCTGGGATTAGCTCCACCTCGCGG
 CTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGTA
 AGGGCCATGATGACTTGACGTACATCCCCACCTCCTCCGGTTGTCACCGGCA
 GTCTCCTTAGAGTGCACCATTACGTGCTGGTAACTAAGGACAAGGGTTGC
 GCTCGTTACGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCCA
 TGCAACCTGTCTCAATGTTCCGAAGGCACCAATCCATCTGGAAAGTTC
 ATTGGATGTCAAGGCCTGGTAAGGTTCTCGCGTTGCTCGAATTAAACCACA
 TGCTCCACCGCTTGTGCGGGCCCCGTCAATTGAGTTAACCTTGCG
 GCCGTACTCCCCAGGCGGTCAACTTAATGCGTTAGCTGCGCCACTAACAGAGCT
 CAAGGCTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGGTA
 TCTAATCCTGTTGCTCCCCACGCTTCGACCTCAGTGTCACTACGCTTACCGCT
 GGTGGTCGCCTCGCCACTGGTGTCCCTATATCTACGCATTACCGCT
 ACACAGGATAATTCCACCCACCCCTACCCACTCTAGCTGACAGTTTG
 AATGCAGGTTCCCAGGTTGAGCCGGGATTCACATCCTACTAACGAACC
 ACCCTACGCGCGCCTTACGCCAGTAATTGATTAACGCTTGACCCCTC
 CTGTATTACCGCTGCTGCCACAGAGGTTAGTCCGGTGCTTATTCTG
 TTCGCTACCGGAAAAACATCTACGTTAGGTAAAATGCCTTCCCTTC
 CCTCACTAAAACAGGTCTTACACTCCGTAAGAACCTTCTTCACAACACT
 GCCGACATACTGG

Strain NRCJ-16

AGAGCTTGCTCTGATTCA CGGGCGGACGGGTGAGTAATGCCTAGGAATCT
 GCCTGGTAGTGGGGACAACGTCTCGAAAGGGACGCTAATACCGCATACTC
 CTACGGGAGAAAGCAGGGACCTCGGGCCTGCGCTATCAGATGAGCCTAG
 GTCGGATTAGCTAGTTGGT GAGGTAATGGCTACCAAGGCACGATCCGTA
 CTGGTCTGAGAGGATGATCAGTCACACTGGA ACTGAGACACGGTCCAGACTC
 CTACGGGAGGCAGCAGTGGGAATATTGGACAATGGCGAAAGCCTGATCC
 AGCCATGCCCGTGTGAAGAAGGTCTCGGATTGTAAAGCACTTTAAGTT
 GGGAGGAAGGGTTGTAGATTAATACTCTGCAATTGACGTTACCGACAGAA
 TAAGCACCGCTAACTCTGTGCCAGCAGCCGCGTAATACAGAGGGT GCAAG
 CGTTAACCGAATTACTGGCGTAAAGCGCGTAGGTGGTTGTTAAGTTG
 GATGTGAAAGCCCCGGGCTCAACCTGGGA ACTGCATTCAAAACTGACAAGCT
 AGAGTATGGTAGAGGGTGGAAATTTCCTGTGAGCGGTGAAATGCGTAG
 ATATAGGAAGGAACACCAGTGGCGAAAGCGTGGGGAGCACACAGC GATTAGATA
 GACACTGAGGT GCGAAAGCGTGGGGAGCACACAGC GATTAGATA
 GGAGTCCCACGCCGCCAACGACGTCAACTAGCCGTCTGGAAGCCTCGAGCTA
 CTA CTGGCGCACCTAACGCA

Strain NRCJ-17

TGGATTGAGAGCTTGCTCTCAAGAAGTTAGCGGCGGACGGGTGAGTAACAC
 GTGGGTAACCTGCCATAAGACTGGATAACTCCGGAAACCGGGCTAATA
 CCGGATAAACATTGAACTGCATGGTCGAAATTGAAAGGCGGCTCGGCTG
 TCACTTATGGATGGACCCCGTGCATTAGCTAGTTGGT GAGGTAACGGCTC
 ACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGT GATCGGCCACACTGGG
 ACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTCCGC
 AATGGACGAAAGTCTGACGGAGCAACGCCCGTGAGTGATGAAGGCTTCGG
 GTCGTAAA ACTCTGTTAGGGAAGAACAGTGCTAGTTGAATAAGCTGGC
 ACCTTGACGGTACCTAACAGAAAGCCACGGCTAACTACGTGCCAGCG
 CGGTAAACGTAGGTGGCAAGCGTTATCGGAATTATTGGCGTAAAGCGCG
 CGCAGGTGGTTCTTAAGTCTGATGTGAAAGCCCACGGCTCAACCGTGGAGG
 GTCATTGGAAACTGGGAGACTTGAGTGAGAAGAGGAAAGT GGAATTCCATG
 TGTAGCGGTGAAATGCGTAGAGATATGGAGGAACACCAGTGGCGAAGGGCA
 CTTCTGGTCTGTAACTGACACTGAGGC GCGAAAGCGTGGGGAGCAAACAGG
 ATTAGATAACCTGGTAGTCCACGCCGAAACGATGAGTGCTAAGTGTAGAG

GGTTCCGCCCTTACTGCTGAAGTTAACGCATTAAGCACTCCGCCTGGGGAG
 TACGGCCGCAAGGCTGAAACTCAAAGGAATTGACGGGGCCCGACAAGCGG
 TGGAGCATGTGGTTAATTGAAGCAACCGAAGAACCTTACCAAGTCGTGA
 CATCCTCTGAAAACCCTAGAGATAGGGCTTCCTCCGGAGCAGAGTGACA
 GGAGGTGCATGGTGTCTCAGCTCTGTC

Strain NRCJ-18

TGACTGAGAGGCCTAACACATGCAAGTCGGCGCTGAAGAGAGCCTGGTCTCT
 GATTCCCGGGAGGGTGAGTGTGCCTCTAAATCTGCCTGGTGTGGGACA
 ACATCTCGAAAGGGACGCTCATATACCATACTGCCTATAGGAGAAAGCGCGG
 GACCTCTGGCCTGTGCTATCTCATGAGCCTAGGTGGATTATAGTTGGT
 GAGGTAATGGCTACCAAGGCACAATCCGTAAGTGGTCTGAGAGGATGATC
 TCTCACACTGGAACGTGAGACACGGTCCCCACTCCTACGGGAGGCACCAGTGG
 GGAATATTGTACACTGGCGAAAGCCTGATCCAGCCCTGCCCGTGTGAA
 AAAAGTCTCTGATTGAAAGCACTTAAGTTGGAGGAAAGGGTGTATT
 AATACTCTCAGTTTGACGTTACCCACACAATAAGCACCCGCTAACTCTGTG
 CCAGCACCCCCGTAATACACAGGGTGACCGTTAATCTCAATTACTCGGC
 GTATAGCGCGCGTATGTGTTAAGTTGGATGTGAAAACCTCGGGCTCA
 AACTGTGAACTGTGCTCTCAACTGTGTTAGCTCTAGTGTGTTAGAGTGTGGT
 GATTTCTGTGTCGCTGAGATGCGTATATATGAGGGAGCACCCCTGTC
 GAAAGCGCCCCCATGTACTGATGACTCTCACTCTGGTGCAGCGTGG
 GCAACACACGAGTATGATATACTCTTAGTACTCCGACGCCACGATGATGA
 ACAACTACGCTGGTGAGCCCTTGCTCTTATTAGCGGGCGCTCTCACAGTT
 ATTATTGTCGAGCGGGCCGCGCACACCGCATGAGCATGATGGTGTATT
 CTAGCACAGCGCGAAGCACTTACCGACGCTGTCCACTAGCATTCTCAA
 AGAGTAGTGGTCGCTTGCAGCAGTTGACCAGGCAGCATGAGCGATTGCTAC
 ACTAGGCTGCATTGCGTATGTGCCTACACGGCACTTCC

Strain NRCJ-19

TCGAGCGGCAGCACGGTACTTGTACCTGGTGGCGAGCGGGACGGGTGAG
 TAATGCCTAGGAATCTGCCTGGTAGTGGGGATAACGCTCGAACGGACGC
 TAATACCGCATACTGCCTACGGGAGAAAGCAGGGACCTCGGGCCTGCGC
 TATCAGATGAGCCTAGGTCGGATTAGCTAGTTGGTGGTAATGGCTCACCA

AGGCAGCGATCCGTAACGGTCTGAGAGGGATGATCAGTCACACTGGAACTGA
 GACACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATATTGGACAATG
 GGCAGAAAGCCTGATCCAGCCATGCCGCGTGTGAAGAAGGTCTCGGATTG
 TAAAGCACTTTAACGTTGGAGGAAGGGCATTACCTAACGTAAGTGTGTTT
 GACGTTCCGACAGAATAAGCACCGGCTAACCTGTGCCAACAGCCGCGTA
 AAGAAAAGGGAGGTACGTAAAATGGAAAAAAAGGGGGAAAAGCCCCGGA
 AGGTGGTCCCCTAAATTGAGATGTTAAAATCCGTGGAAACAAAG

Strain NRCJ-20

CCGTGGCAGGCCCTCCGAAGGTTAACGCTACCTGCTTCTGGTGCAACAAAC
 TCCCATGGTGTGACGGGCGGTGTGTACAAGGCCGGAACGTATTACCGCA
 GCAATGCTGATCTGCGATTACTAGCGATTCCGACTTCATGGAGTCGAGTGCA
 GACTCCAATCCGGACTGAGATAGGGTTCTGGATTGGCTTACCGTCGCCGG
 CTTGCAGCCCTCTGTCCTACCATTGTAGTACGTGTAGCCCTGGCCGTAAG
 GCCATGATGACTTGACGTACCCCCACCTTCCTCCGGTTGTACCGGCCGGT
 CTCCCTAGAGTCCCACCATTACGTGGCAACTAAGGACAAGGGTTGCGCT
 CGTTCGGGACTTAACCAAACATCTCACGACACGAGCTGACGACAGCCATGC
 AGCACCTGTGTTCGAGTTCCGAAGGCACCAATCCATCTGGAAAGTTCTCG
 ACATGTCAAGGCCAGGTAAAGGTTCTCGCGTTGCATCGAATTAAACCACATA
 CTCCACCGCTTGTGCGGGCCCCGTCAATTCTTGAGTTTAGTCTGCGAC
 CGTACTCCCCAGGCAGCGAACCTAACGCGTTAGCTCGATACTGCGTGCCAA
 ATTGCACCAAACATCCAGTTGCATCGTTAGGGCGTGGACTACCAGGGTATC
 TAATCCTGTTGCTCCCCACGCTTCGTGCCTCAGTGTCAAGTGTGGTCCAGGT
 AGCTGCCTCGCCATGGATGTTCCCTGATCTCACGCATTCACTGCTACA
 CCAGGAATTCCGCTACCCCTTACCCACACTCTAGTCGTCCAGTTCCACTGCAG
 TTCCCAGGGTTGAGCCCAGGGCTTCACAACAGACTAAACGACCACTACGC
 ACGCTTACGCCAGTAATTCCGAGTAACGCTTGCACCCCTCGTATTACCGCG
 GCTGCTGGCACGAAGTTAGCCGGTGTATTCTTGGTACCGTCATCCCAA
 CCGG

Strain NRCJ-21

TACACCGTGGTAACCGTCCTCCGAAGGTTAGACTAGCTACTTCTGGTGCAAC
 CCACTCCCATGGTGTGACGGGCGGTGTACAAGGCCGGAACGTATTAC
 CGCGACATTCTGATTGCGATTACTAGCGATTCCGACTTCACGCAGTCGAGTT

GCAGACTGCGATCCGGACTACGATCGGTTTGAGATTAGCTCCACCTCGCG
 GCTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGT
 AAGGGCCATGATGACTTGACGTACCCCCACCTCCTCCGGTTGTCACCGGC
 AGTCTCCTTAGAGTGCCACCATAACGTGCTGGTAACTAAGGACAAGGGTTG
 CGCTCGTTACGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCC
 ATGCAGCACCTGTGTCAGAGTCCGAAGGCACCAATCCATCTCTGGAAAGT
 TCTCTGCATGTCAAGGCCTGGTAAGGTTCTCGCGTTGCTCGAATTAAACCA
 CATGCTCCACCGCTTGTGCGGGCCCCGTCAATTCAATTGAGTTAACCTTG
 CGGCCGTACTCCCCAGGCGGTCAACTTAATGCCTAGCTGCGCCACTAAAAT
 CTCAAGGATTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGG
 TATCTAACCTGTTGCTCCCCACGCTTCGCACCTCAGTGTCAAGTATCAGTCC
 AGGTGGTCGCCTCGCCACTGGTGTTCCTCCTATATCTACGCATTCACCGCT
 ACACAGGAAATTCCACCAACCCCTCTACCGTACTCTAGCTGCCAGTTGGGA
 TGCAAGTCCAGGTTGAGCCCGGGGTTCACATCCAACCTAACAAACCACCT
 ACGCGCGCTTACGCCAGTAATTCCGATTAACGCTTGCACCTCTGTATTACC
 CGGGCTGCTGGCACAGAGTTAGCC

Strain NRCJ-22

GTAGGCTCTACACATGCAAGTCGAACGGCAGCACAGGTAAAGAACCTGCTCT
 TATGGGTGGCGAGTGGCGATGGGTGATGAATACTACATAACTCTACGGTTA
 ACTTGGGGAAACGTAATGGAAACTTACTTAAAATTCAAGATAACTAAAAA
 TCGATGGCATTATGGTGGGAAATGTTGATTGAATGAGCCGATGTCGG
 ATTAGCTAGTTGGCGGGTAAAGGCCACCAAGGCGACGATCCGTAGCTGGT
 CTGAGAGGATGATCAGCCACACTGGAACCTGAGACACGGTCCAGACTCCTACG
 GGAGGCAGCAGTGGGAATATTGGACAATGGCGCAAGCCTGATCCAGCCA
 TACCGCGTGGGGAGAAAAGGCTGGCCGTTGTAAACACTTTTTGGAG
 GAAAAGACACGATGCAACACCTGGTGTAGAGTTGCAAAAGAACCAATCA
 ACCTGGGGAGTTCTACATGTCTAGGGGGTAAAATTGTTGCGCTTC
 AACTAAAAAAACAGATACTCAAGCGGGTGCACCCCCGAAACTTACTT
 TGAGAACACTCTGTGACCGTCCTCCCCGGCGGGAAACTGAAAGCGTTA
 CTGTGCTACAGCGTGCAAAATCCCCCGCATCTAACACATCTGTAGGG
 GGGTGGATACAAGGTTATCAATTGTTGCTCCCCACACTTGTGCCACA
 GAGTGAGAGTTGTGAGGAAACAGCCCTCGAGACGGTGGTTGTCACCACT
 CGAGAGTTAACTGCGCGGGGGATATTGCGCTCTCATTCAACACTAAA

CCTTCATTCGCTGCACGCGAGTGGCTGCCGGCCTCCACACACAAAT
ATAA

Strain NRCJ-23

TCGAGCGGATGAAGAGAGCTGCTCTGATTCAAGCGCGGACGGGTGAGTA
ATGCCTAGGAATCTGCCTGGTAGTGGGGACAACGTTGAAAGGAACGCTA
ATACCGCATACGTCCTACGGGAGAAAGCAGGGACCTCGGGCCTGCGCTA
TCAGATGAGCCTAGGTGGATTAGCTAGTTGGTGAGGTAATGGCTACCAAG
GCGACGATCCGTAACTGGTCTGAGAGGATGATCAGTCACACTGGAAC TGAGA
CACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATATTGGACAATGGC
GAAAGCCTGATCCAGCCATGCCCGTGTGAAGAAGGTCTCGGATTGTAA
AGCACTTAAGTGGGAGGAAGGGTTGTAGATTAATACTCTGCAATTGAC
GTTACCGACAGAATAAGCACCGGCTAACCTGTGCCAGCAGCCCGTAATA
CAGAGGGTGCAAGCGTTAACCGAATTACTGGCGTAAAGCGCGCTAGGTG
GTTGTTAAGTGGATGTGAAAGCCCCGGCTAACCTGGGAAC TGCAATTCA
CAACTGAGAAGCTCTAGTATGGTAGAGGGTGCTGGAAAAATCTCAAGGTTCC
CGGGGGCTAGCTAACATCTATTACGGACTGCATTCCCCGGTGCCACAATC
CGGTTGCACACCACGCTTGGGCCAAAGGGGGGGGACACTACAGGG
ATATCATACCGGGGAAGGGGAGAACCTTACAACAATGTCAAATATTGCCT
GGGACCCATTAAAGTTCTACCCC GGCTCTCCAAGCGCTTAATTGACGT
ACAAGAGAACAAAGTGCCCGGTTGGGTCAAGTGGCTATAAAATAGACAGG
ACCCCCGACACACCCCCGCCGCACCTGTTAGATATAATGTGGGACAT
CTGCGGAGAGAACACACCAAGGGAGGAGAACAAACGAAGCACCCCTCC
CCGAAATATGGGTGGTGGTTTATTTCGGAGAACATACACACACCACG
CCTGCCAGCCATGGCATGA

Strain NRCJ-24

TGCAAGCTCGAGCGGATGAAGAAGAAGCTCTGCTCTGATTCAAGCGCGGA
CGGGTAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGACAACGTCGAA
AGGGACGCTAACCGCATACGTCCTACGGGAGAAAGCAGGGACCTCGG
GCCTGCGCTATCAGATGAGCCTAGGTGGATTAGCTAGTTGGTGAGGTAAT
GGCTCACCAAGGCAGCAGTGGCTGAGAGGATGATCAGTCACAC
TGGAACTGAGAACCGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATATT
GGACAATGGCGAAAGCCTGATCCAGCCATGCCCGTGTGAAGAAGGTCT

TCGGATTGTAAAGCACTTAAGTGGGAGGAAGGGTTAGATTAATACTCT
 GCAATTGACGTTACCGACAGAATAAGCACCGGCTAACTCTGTGCCAGCAG
 CCGCGGTAAATACAGAGGGTGCAAGCGTTAACCGAATTACTGGCGTAAAGC
 GCGCGTAGGTGGTTGTTAACGGATGTGAAAGCCCCGGCTAACCTGGG
 AACTGCATTCAAAACTGACAAGCTAGAGTATGGTAGAGGGTGGTGAATTTC
 CTGTGTAGCGGTGAAATCGTAGATATAGGAAGGAACACCAGTGGCGAAGG
 CGACCACCTGGACTGATACTGACACTGAGGTGCGAAAGCGTGGGAGCAA
 CAGGATTAGATACCCCTGGTAGTCCACGCCGTAAACGATGTCAACTAGCCGTT
 GGGAGCCTTGAGCTCTAGTGGCGCAGCTAACGCATTAAGTGANCGCCTGG
 GGAGTACGGCCGCAAGGTTAAAACCAAATGAATTGACGGGCCCCACAA
 GCGGTGGAGCATGTGGTTAACCGAAGCAACGGCGAAGAACCTTACCAAGGC
 CCTTGACATCCAATGAACCTTCCAGAGATGGATTGGTGCCTCG

Strain NRCJ-25

ACATGCAAGTCGAGCGGCAGCACGGTACTTGTACCTGGTGGCGAGCGGCGG
 ACGGGTAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGATAACGCTCGGA
 AACGGACGCTAACCGCATACTGCCTACGGGAGAAAGCAGGGGACCTTCG
 GCCCTGCGCTATCAGATGAGCCTAGGTCGGATTAGCTAGTTGGTGAGGTAA
 TGGCTACCAAGGCACGATCCGTAACTGGTCTGAGAGGATGATCAGTCACA
 CTGGAACGTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATAT
 TGGACAATGGCGAAAGCCTGATCCAGCCATGCCGCGTGTGAAGAAGGTC
 TTCGGATTGTAAAGCACTTAAGTGGGAGGAAGGGATTACCTAACGTT
 AAGTGTGTTGACGTTACCGACAGAATAAGCACCGGCTAACTCTGTGCCAGCA
 GCCCGGTAATACAGAGGGTGCAAGCGTTAACCGAATTACTGGCGTAAAG
 CGCGCGTAGGTGGTCTAAATTGGATGTTGAAATCCCCGGCTAACCGG
 GGCACGGATTCAAAACGTTAACGTTAACGTTGGAAAAGGGGGGGAAATT
 TCCGGCGCCACGGCAATGCGACAACATT

Strain NRCJ-26

GCAGGCCTAACACATGCAAGTCGAGCGGCAGCACGGTACTTGTACCTGGT
 GCGAGCGGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGG
 ATAACGCTCGAACGGACGCTAACCGCATACTGCCTACGGGAGAAAGC
 AGGGGACCTTCGGGCCTGCGCTATCAGATGAGCCTAGGTCGGATTAGCTAG
 TTGGTGAGGTAATGGCTACCAAGGCACGATCCGTAACTGGTCTGAGAGGA

TGATCAGTCACACTGGAACTGAGACACGGTCCAGACTCCTACGGGAGGCAGC
 AGTGGGAATATTGACAATGGCGAAAGCCTGATCCAGCCATGCCGCGTGT
 GTGAAGAAGGTCTCGGATTGTAAGCACTTAAGTTGGGAGGAAGGGCAGT
 TACCTAATACGTATCTGTTGACGTTACCCCCAGAGTACGCACCCGGCCAAC
 TCTGTGCCCGCAGCCCCGGTAAGACAGAGGGTGCACGCGTTAGATCGGAATT
 ACTGGGGCGTAAAGCGCGTAGGTGGTCTTAAGTTGGATGTGAAATCCC
 TGGGATCAACCTGGCAACTGCCTCAAAACTGACAAACTAGACTTGGTAGA
 GGGTGGTGGAGTTCACTGCTGCGGTGCAATGTGTATCTATAAGAGGAGC
 ACCACTGGGGAAGGAATC

Strain NRCJ-27

AGGCCTAACACATGCAAGTCGAGCGGATGACGGGAGCTTGCTCCTGATTCA
 CGGGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGACAAC
 GTTCGAAAGGAACGCTAATACCGCATACTGCCTACGGGAGAAAGCAGGG
 ACCTTCGGGCCTTGCCTATCAGATGAGCCTAGGTCGGATTAGCTAGTTGGT
 GGGTAATGGCTACCAAGGCGACGATCCGTAACGGTCTGAGAGGATGATCA
 GTCACACTGGAACTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTGG
 GGAATATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCGCGTGTGAA
 GAAGGTCTCGGATTGTAAGCACTTAAGTTGGGAGGAAGGGCAGTAAGCG
 AATACCTTGCCTGTTGACGTTACCGACAGAATAAGCACCGGCTAACTCTGTG
 CCAGCAGCCGCGGTAAATACAGAGGGTGAAGCGTTATCGGAATTACTGGC
 GTAAAGCGCGTAGGTGGTCTTAAGTTGGATGTGAAAGCCCCGGCTCA
 ACCTGGGAACTGCCTCCAAAAGTGGCCGCTAGAGTACGGGAGAGGGGGGC
 TAAAATTCTCAAGGATTCAACGACTCGTTGACATCGTTACGGCACGCCAC
 CCAGATGCCTAATCCCTGGTTGCTCCAGCCTTTCGACCCCAAAGCTAGT
 AAACAACCAGGTGTTGACTTCTGGAAGGCGCGCTCCAATATCTAACATT
 ACCGCTACCCCGTGAATTCAACCGCCTCCAGACATATTACTGCCGCTG
 GGGCGAATGCCCGGGGTTACAGGCATTGAATTGCCAGGTGACCAGACG
 ACCTGCTGCACCATGTGGCTAATTGCTAGCCACTCCAATATCCCTCCCTAGG
 GCTCACAAAGCCCCAGCACTCACGTAGTGGAGTGGATTGCTTGGT

Strain NRCJ-28

TGCGCAGGCCTAACACATGCAAGTCGAGCGGATGAAGGGAGCATTGCTCCTG
 AATTCAAGCGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGG

GACAACGTTCGAAAGGAACGCTAATACCGCATACGTCTACGGGAGAAAGC
 AGGGGACCTTCGGGCCTGCGCTATCAGATGAGCCTAGGTCGGATTAGCTAG
 TTGGTGAGGTAATGGCTACCAAGGCAGATCCGTAACTGGTCTGAGAGGA
 TGATCAGTCACACTGGAACGTGAGACACGGTCCAGACTCCTACGGGAGGCAGC
 AGTGGGAAATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCCGCTGT
 GTGAAGAAGGTCTCGGATTGTAAGCACTTAAGTTGGGAGGAAGGGTTGT
 AGATTAATACTCTGCAATTGACGTTACCGACAGAATAAGCACCGGCTCACT
 CTGTGCCAGCACCCCCGGTGATACAGAGAGGGAAAGCGCTAAAGTAGTTTC
 TGGCGTAAAACCGCGCATACGTGTCTAAAAGTGAGATGAAAAATCCGA
 GTTCACACCTTGGAACTGCATCCAAAAAGTGGCACCTAAAATTCTGGAAC
 GGGCGGTTGAAATCTCCAGGGCTCCGATGGCTTACATTCTTA

Strain NRCJ-29

GCAGGCCTAACACATGCAAGTCGAGCGGCAGCACGGTACTTGTACCTGGT
 GCGAGCGCGGACGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGG
 ATAACGCTCGAACGGACGCTAACACCGCATACGTCTACGGGAGAAAGC
 AGGGGACCTTCGGGCCTGCGCTATCAGATGAGCCTAGGTCGGATTAGCTAG
 TTGGTGAGGTAATGGCTACCAAGGCAGATCCGTAACTGGTCTGAGAGGA
 TGATCAGTCACACTGGAACGTGAGACACGGTCCAGACTCCTACGGGAGGCAGC
 AGTGGGAAATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCCGCTGT
 GTGAAGAAGGTCTCGGATTGTAAGCACTTAAGTTGGGAGGAAGGGAGT
 TACCTAACGTAATTGTTTGACGTTACCGACAGAATAAGCACCGCTAACT
 CTGTGCCAGCCGCGTAATACAGAGGGTGCAAGCGTTAACGGAAATTAC
 TGGCGTAAAAGCGCGCTAGGTGGTCTGTTAAGTTGGATGTGAAATCCCCGG
 GCTAACCTGGAACTCGCTTACAGTGAGAGCTAGAGTATGGTGGAGGGT
 GGTGG

Strain NRCJ-30

TCGAACGGCAGCACAGAGGAGCTTGCTCCTGGGTGGCGAGTGGCGGACGGG
 TGAGGAATGCATCGGAATCTGCCTATTGTGGGGATAACGTAGGGAAACTT
 ACGCTAACACCGCATACGTCTACGGGAGAAAGTGGGGACCGCAAGGCCTC
 ACGCAGATAGATGAGCCGATGCCGGATTAGCTAGTTGGCGGGTAAAGGCC
 ACCAAGGCGACGATCCGTAGCTGGTCTGAGAGGATGATCAGCCACACTGGAA
 CTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATATTGGACA

ATGGGCGCAAGCCTGATCCAGCCATGCCCGTGTGAAGAAGGCCTCGGG
 TTGTAAAGCACTTTGTCCGGAAAGAAAAGCGCTCGATTAATACTCGGGTGT
 ATGACGGTACCGGAAGAATAAGCACCGGCTAACCTCGTGCAGCAGCCCG
 TAATACGAAGGGTGCAAGCGTTACTCGGAATTACTGGCGTAAAGCGTGC
 AGGTGGTTGTTAAGTCTGATGTGAAAGCCCTGGGCTAACCTGGGAAC
 ATTGGAAACTGGCTTACTAGAGTCGGTAGAGGGGTGTTGAAATTCCC
 AGCAGTGAAATGCGTAGATATCGGGAGGAACATCTGTGGCGAAGGC
 CCTGGGCCAGCACTGACACTGAGGCACGAAAGCGTGGGAGCAA
 TTAGATAACCCTGGTAGTCCACGCCCTAAACGATGCGAACTGGATGTTGG
 GGCAACTTGGCCCTCAGTCTAAAGCTAACGCGTTAACGCTAACG
 AAGTCCGTCGACGACTGAAACACAAAAGGAAATTGAAGGGAGCGCG
 CAAAGACGGTCGGGTGGTGGTTAATTAGAGGACGGCAGAGAACAA
 TACCTTCTACTTGCCTGTCACGAACTGCCACTATTCCGAGTAGTTGG
 TTTGGTATAGGGGACGCACGATAGAAGACTGGC

Strain NRCJ-31

TGCCTAATACATGCAAGTCGAGCGAATGGATATAGAGAGCTTGTCTTAAGA
 AGTTAGCGGCGGACGGGTGAGTAACACGTGGTAACCTGCCATAAGACTGG
 GATAACTCCGGAAACCGGGGCTAACCGGATAATATTGAACTGCATGG
 TTCGAAATTGAAAGGCGCTCGGCTGTCACTTATGGATGGCCCCCGTCCC
 ATTCCAGTTGTGAAGGACCGGCTCACCAAGGCAACGATCCTACCCAAATG
 AGAAGGGAATCGGCAAACGGACTGAGACACGGCAAACCTACGGAGG
 CACAGTAGGGAATCTCCGCAATGGAAAAAGTCTGACGGACAACGCCCG
 AGTGTGAAGGCTTCGGCGTAAACTCTGTTGGAAAACAAGTGCTA
 TTGAATAAGCTGGACCTTGACGTACATAACCAGAAAGCCCTGCTAACTACGT
 GCCACACCGCGTGATACTAGTTGCAAGATATCCGAAATTGGCGTAAGCCG
 CGCAGGTGGTTTAATTAGATGTACACCCCGGTTCTCCGTGTTGGTCATTG
 AACTGGGACACTTACTGCAAAGAAGAAAGTGGATTCCATGTGTACGGTGAA
 ATGCTAGTGTATGGCGGAACACCTGCCTATGCGAACATCAGGTTGAAACTG
 CCACGGGCTCGAAACCTGGGCTTGGGCCATTGGAATAGAAGCGCGCAGTCCCC
 CCCAAACAAGATTGCAAATGAATAACGGTTCCGCCATAGTGCCTTACAA
 ATTAAAAAAACCCACCGGGCTTACCGCAACATGCTAAACTAATAGGAATTAC
 ATGCGCCCCCGCCGCGACATGTGGTTAATTCAAGCACGTCAAAGACCTA
 CCAAGTCTGAATTCTTGACAACCTTAGATGAGATTCACTTGACCGAGT
 GCCCGGGAAATCATGTTTCTCAACTATCT

Strain NRCJ-32

ATGCGCATGCGCGTGCCTAATAACACTGCAAGCTGAGCGAACTGGATCAGAG
 AGCTTGCTCTTAAGAACGCTAGCGCGGACGGGTGAGTAACACGTGGTAACC
 TGCCCATAAGACTGGATAACTCCGGAAACCGGGGCTAATACCGGATAATA
 TTTGAACTGCATGGTCGAAATTGAAAGGCGGCTCGCTGTCACTTATGGA
 TGGACCCCGTGCATTAGCTAGTTGGTGAGGTAACGGCTACCAAGGCAAC
 GATGCGTAGCCGACCTGAGAGGGTATCGGCCACACTGGGACTGAGACACG
 GCCCAGACTCCTACGGGAGGCAGCAGTAGGAAATCTCCGCAATGGACGAA
 AGTCTGACGGAGCAACGCCCGTGAGTGATGAAGGCTTCGGTCGTAAAAC
 TCTGTTAGGAAAGAACAAAGTGTAGTTGAATAAGCTGGCACCTGACGG
 TACCTAACCAAGAAAGCCACGGCTAACTACGTGCCAGCAGCCCGGTAATACG
 TAGGTGGCAAGCGTTATCGGAATTATTGGCGTAAAGCGCGCAGGTGGT
 TTCTTAAGTCTGATGTGAAAGCCCACGGCTCAACCCTGGAGGGTCATTGGAA
 ACTGGGAGACTTGAGTGCAGAAGAGGAAAGTGGATTCCATGTGTAGCGGTG
 AAATGCGTAGAGATATGGAGGAACACCAAGTGGCGAAGGCGACTTCTGGTCT
 GTAACGTGACACTGAGGCACGAAAGCGTGGGAGCAAACAGGATTAGATACC
 CTGGTAGTCCACGCCGTAAACGATGAGTGCTAAGTGTAGAGGGTTCCGCC
 CTTTAGTGCTGAAGTTAACGCATTAAGCACTCCGCTGGGAGTACGGCCGC
 AAGGCTGAAACTCAAAGGAATTGACGGGCCGCACAAGCGTGGAGCATG
 TGGTTAACCGAAGCAACCGAAGAACCTTACCGGTCTGACATCCTCTGA
 AAACCCCTAGAGATAGGGCTTCTCCTCGGGAGCAGAGTGACAGTGGTGCAT
 GGTTGTCGTAGCTCGTGTGAGATGTTGGTTAGTCCGCACGAGCGCA
 CCCTTGATCCTAGGTGGCAATCATAGTGGCACCTCTAGTGACTGCTGTGACAA
 ACCGGAGAGGTGGAAATGAACCGTCA

Strain NRCJ-33

TCGAGCGAATGGATAAGAGAGACTTGCTCTCAAGAACGTTAGCGCGGACGGGT
 GAGTAACACGTGGTAACCTGCCATAAGACTGGATAACTCCGGAAACCG
 GGGCTAATACCGATAACATTTGAACGCTGCATGGTCGAAATTGAAAGCGG
 CTTCGGCTGTCACCTATGGATGGACCCCGCGTCGCATTAGCTAGTTGGTGGAGGT
 AACGGCTACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTATCGGCC
 AACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTAGGGA
 ATCTTCCGCAATGGACGAAAGTCTGACGGAGCAACGCCCGTGAGTGATGAA
 GGCTTCGGTCGAAACTCTGTTAGGAAAGAACAGTGTAGTTGAA

TAAGCTGGCACCTGACGGTACCTAACAGAAAGCCACGGCTAATACGTGC
 CAGCAGCCCGGTAATACGTAGGTGGCAAGCGTTATCCGAATTATTGGCG
 TAAAGCGCGCAGGTGGTTCTTAAGTCTGATGTGAAAGCCCACGGCTCAA
 CCGTGGAGGGTCATTGGAAAAGTGGAGACTTGAGTCACAAAAGGGAAAA
 TGGGATTCCCCGTGAACGGTGTAAATGTGTAGGAAATAATGAGAACACCCC
 TGGGGCAGGGGCCATTCTGGCTTTGCCCTCCCATTGCGGCGAAACCC
 GGGGAACAAAAGAGGAAAGGATGTCCTGGTTGCCCCCCCCCCCCAAAAC
 GAACAAGTGCCTAGTTAGGAGGGTTCCCCCCCCCTATTACTGCTAAATC
 TTAACCGAATTAAAAACCCCCCGGGGAAGAACGGGCCCCAACGCGGGTGCA
 GACATTGTGGTTAAAATCGAACGCCAACACAAAAAAACCTTACCGGAGG
 TTCTGGACAATCCTCTAGAAAAACCCAAATATAAGGGTCTTCCCCCT
 TTAGAGAACAGAAATGACACAGGGCGTACCAAAGGGTATTGTCGCTACG
 ACACACCTGCGTCGAGGGGAAGATGGTTGGGGGATAAAGTCTCCTCACACA
 AAACTACCGCGCGAACCGCTTGTGGAATCCC

Strain NRCJ-34

GTGCCTAATACATGCAAGTCGAGCGAATGGATTGAGAGCTGCTCTCAAGAA
 GTTAGCGCGGACGGGTGAGTAACACGTGGTAACCTGCCATAAGACTGGG
 ATAACCTCCGGAAACCGGGCTAATACCGATAACATTGAACTGCATGGT
 TCGAAATTGAAAGGCGGCTCGGCTGTCACTTATGGATGGACCCGCGTCGCA
 TTAGCTAGTTGGTGAGGTAAACGGCTACCAAGGCAACGATGCGTAGCCGACC
 TGAGAGGGTATCGGCCACACTGGACTGAGACACGGCCAGACTCCTACGG
 GAGGCAGCAGTAGGAAATCTCCGCAATGGACGAAAGTCTGACGGAGCAAC
 GCCCGTGAGTGTGAAGGCTTCGGCTGTAAAACCTGTTAGGGAAG
 AACAAAGTCTAGTTGAATAAGCTGGCACCTGACGGTACCTAACCAAGAAC
 CACGGCTAACTACGTGCCAGCAGCCCGGTAAACGTAGGTGGCAAGCGTTA
 TCCGGAATTATTGGCGTAAAGCGCGCAGGTGGTTCTTAAGTCTGATGTG
 AAAGCCCACGGCTAACCGTGGAGGGTATTGAAACTGGAGACTTGAGTG
 CAGAAGAGGAAAGTGGATTCCATGTGTAGCGGTGAAATGCGTAGAGATAT
 GGAGGAACACCAGTGGCGAAGCGACTTCTGGCTGTAACTGACACTGAGG
 CGCGAAAGCGTGGGAGCAAACAGGATTAGATAACCTGGTAGTCCACGCCGT
 AAACGATGAGTGTAAAGTGTAGAGGGTTCCGCCCTTAGTGCTGAAGTTA
 ACGCATTAAGCACTCCGCCTGGGAGTACGGCCGCAAGGCTGAAACTCAAAG
 GAATTGACGGGGCCCGACAAGCGGTGGAGCATGTGGTTAATTGAAAGCA

ACCGGAAGAACCATACCAGGTCTGACATCCTCTGAAAATCCTAACACGATA
GGGCTTCTCATT CGC

Strain NRCJ-35

CGGAGCCTGCGCGTGCCTAAACATGCAAGTCAGCGAAAGATAAGGAGCTTGC
TCCTTCGATGTTAGCGGCGGAGGGGTGAGTAACACGTGGCAACCTACCTTA
TAGTTTGGGATAACTCCGGAAACCGGGGCTAATACCGAATAATCTGTTCA
CCTCATGGTGAAACACTGAAAGACGGTTTCGGCTGCGCTAGGATGGGCC
CGCGCGCATTAACTAGTTGGTGAGGTAACGGCTACCAAGGCGACGATGCG
TAGCCCACCTGAGAGGGTGTAGCGGCCACACTGGGACTGAAACACGGCCATA
CTCCTACGGGAGGCAGCAGTGGGAATCTTCCACAATGGCAAAAGCCTGAT
GGAGCAACGCCCGTGAGTGAAGAAGGATTCGGTTCGTAAAACCTCTGTTGT
GAGGGAAAGAACAAAGTACAGTAATAACTGGCTGTACCTTGACGGTACCTTATT
AAAAAGCCACGGCTAACTACGTGCCAGCAGCCGGTAATACGTAGGTGGC
AAGCGTTGCCCGAATTATTGGCGTAAAGCGCGCAGGTGGTTCTTAAG
TCTGATGTGAAAGCCCACGGCTAACCGTGGAGGGTCATTGAAAACGGAG
ACTTGAGTGCAGAACAGAGGATAGTGGATTCCAAGTGTAGCGGTGAAATGCGT
AGAGATTGGAGGAACACCAAGTGGCGAAGGCGACTATCTGGTCTGTAACG
CACTGAGGCCGAAAGCGTGGGAGCAAACAGGATTAGATAACCTGGTAGT
CCACGCCGTAAACGATGAGTGTCTAAGTGTAGGGGGTTCCGCCATTAGT
GCTGCAGCTAACGCATTAAGCACTCCGCCTGGGAGTACGGTCGCAAGGCTG
AAACTCAGAGGAATTGACGGGGCCCCGCACAAGCGTGGAGCATGTGGTT
TATTAGCAGCAACCGAAGAACATTACCAAGGTCTTGACATCCTGTTGACCAC
TGAAAAGAAAGTTGTTCTCCCCTCGGGGACAACGGTACTGGTGGCTGCA
TGGTTGTCGTAGCTCGTGTGAGATGGTGTATTAGTCCAGCACCGAGC
GCCAACTCGATGATCCTAGCTGCCATTTAGTGGTACTCTGAGCTGA
CCTGGCTGTTGAACAACCGGAGAGTGGATGGACGATTCCATCATGCACTGCC
CTCTTATCGAGCGCGTGCACCTAC

Strain NRCJ-36

TGCAGCTTGCCTGCGCTAATACATGCAAGTCAGCGAATGGATTGAGAGCT
TGCTCTCAAGAACAGTTAGCGGCGGACGGGTGAGTAACACGTGGTAACCTGCC
CATAAAGACTGGATAACTCCGGAAACCGGGGCTAATACCGATAACATT
GAACTGCATGGTCGAAATTGAAAGGCGGCTCGGCTGTCACTTATGGATGG

ACCCGCGTCGCATTAGCTAGTTGGTGAGGTAACGGCTACCAAGGCACGAT
 GCGTAGCCGACCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACGGCC
 CAGACTCCTACGGGAGGCAGCAGTAGGAAATCTTCCGCAATGGACGAAAGTC
 TGACGGAGCAACGCCGCGTGAGTGATGAAGGCTTCGGGCGTAAAACCTCTG
 TTGTTAGGAAAGAACAAAGTGCTAGTTGAATAAGCTGGCACCTGACGGTACC
 TAACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGGTAATACGTAGG
 TGGCAAGCGTTATCCGGAATTATTGGCGTAAAGCGCGCGCAGGTGGTTCT
 TAAGTCTGATGTGAAAGCCCACGGCTAACCGTGGAGGGTCATTGGAAACTG
 GGAGACTTGAGTGCAGAACAGGAAAGTGGATTCCATGTGTAGCGGTGAAA
 TCGTAGAGATATGGAGGAACACCAGTGGCGAAGGCACCTTCTGGTCTGTA
 ACTGACACTGAGGCGCAAAGCGTGGGAGCAAACAGGATTAGATAACCTG
 GTAGTCCACGCCGTAACGATGAGTGCTAACGTGTTAGAGGGTTCCGCCCTT
 AGTGTGAAGTTAACGCATTAAGCACTCCGCTGGGAGTACGGCGCAAGG
 CTGAAACTCAAAGGAATTGACGGGGCCCGACAAGCGGTGGAGCATGTGG
 TTTAATTGAAAGCAACCGAAGAACCTTACCAAGGTCTTGACATCCTCTGAAA
 ACCCTAGAGATAGGCTTCCTCGGAGCAGAGTGACAGGTGGTCATGGT
 TGTGTCAGCTCGTGTGAGATGTTGGGTTAGTCCGCACCGAGCGCAAC
 CCTTGGATCTTAGTTGCCATCATTAGTGGTCACCTCTAACGTGACTGCCGTGA
 CAACCGAGGAATGTGCGATGACGTCCGAATCAATCCATTGAACACAT

Strain NRCJ-37

TGCAGCTTGCCTGCGCTAACATGCAAGCCAGCGAATGGATTGAGAGCT
 TGCTCTCAAGAACGTTAGCGCGGACGGGTGAGTAACACCGTGGTAACCTGCC
 CATAAGACTGGATAACTCCGGAAACCGGGCTAACCGGATAACATT
 GAACTGCATGGTCGAAATTGAAAGGCGGCTCGGCTGTCACTTATGGATGG
 ACCCGCGTCGCATTAGCTAGTTGGTGAGGTAACGGCTACCAAGGCACGAT
 GCGTAGCCGACCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACGGCC
 CAGACTCCTACGGGAGGCAGCAGTAGGAAATCTTCCGCAATGGACGAAAGTC
 TGACGGAGCAACGCCGCGTGAGTGATGAAGGCTTCGGGCGTAAAACCTCTG
 TTGTTAGGAAAGAACAAAGTGCTAGTTGAATAAGCTGGCACCTGACGGTACC
 TAACCAGAAAGCCACGGCTAACGTGCCAGCAGCCGGTAATACGTAGG
 TGGCAAGCGTTATCCGGAATTATTGGCGTAAAGCGCGCGCAGGTGGTTCT
 TAAGTCTGATGTGAAAGCCCACGGCTAACCGTGGAGGGTCATTGGAAACTG
 GGAGACTTGAGTGCAGAACAGGAAAGTGGATTCCATGTGTAGCGGTGAAA
 TCGTAGAGATATGGAGGAACACCAGTGGCGAAGGCACCTTCTGGTCTGTA

ACTGACACTGAGGCGCAAAGCGTGGGAGCAAACAGGATTAGATAACCCTG
 GTAGTCCACGCCGTAAACGATGAGTGCTAAGTGTAGAGGGTTCCGCCCTT
 AGTGCTGAAGTTAACGCATTAAGCACTCCGCCTGGGAGTACGGCCGCAAGG
 CTGAAACTCAAAGGAATTGACGGGGCCCGCACAAGCGGTGGAGCATGTGG
 TTTAATTGAGCAACCGAAGAACCTTACCAAGGTCTGACATCCTCTGAAA
 ACCCTAGAGATAGGGCTTCTCCCTCGGGAGCAGAGTGACAGGTGGTGCATG
 GGTTGTCGTAGCTCGTGTGAGATGTTGGTTAGTCCGCACGAGCGCA
 CCCTGATCTTAGTTGCCAATCATAGTGGCACTCTAGTGACTGGCGTGACGAC
 GGAGGAAGGTGGATGACGTTCCAATCATTCACTGGCAC

Strain NRCJ-38

ATGCCCTGCGCGCTAATACATGCAAGCTCGAGCGAATTGGATTGAGAG
 CTTGCTCTCAAGAAGTTAGCGGCGGACGGGTGAGTAACACGTGGTAACCTG
 CCCATAAGACTGGGATAACTCCGGAAACCGGGCTAATACCGGATAACATT
 TTGAAC TGCA TGGT CGAA ATT GAA AGG CGG CT CGG CT GT CA CT TAT GG ATG
 GACCCCGCGTCG CATT AGCT AGTTGGT GAGG TAACGGCTACCAAGGCAACGA
 TCGTAGCCGACCTGAGAGGGTATCGGCCACACTGGACTGAGACACGGCC
 CAGACTCCTACGGGAGGCAGCAGTAGGAAATCTCCGCAATGGACGAAAGTC
 TGACGGAGCAACGCCGCGTGAGTGATGAAGGCTTCCGGTGTAAAACCTG
 TTGTTAGGAAAGAACAAAGTGCTAGTGAAATAAGCTGGCACCTGACGGTACC
 TAACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGGTAATACGTAGG
 TGGCAAGCGTTATCCGAATTATTGGCGTAAAGCGCGCGAGGTGGTTCT
 TAAGTCTGATGTGAAAGCCCACGGCTAACCGTGGAGGGTATTGGAAACTG
 GGAGACTTGAGTGAGAACAGGAAAGTGGATTCCATGTGTAGCGGTGAAA
 TCGTAGAGATATGGAGGAACACCAGTGGCGAAGGCGACTTCTGGTCTGTA
 ACTGACACTGAGGCGCAAAGCGTGGGAGCAAACAGGATTAGATAACCCTG
 GTAGTCCACGCCGTAAACGATGAGTGCTAAGTGTAGAGGGTTCCGCCCTT
 AGTGCTGAAGTTAACGCATTAAGCACTCCGCCTGGGAGTACGGCCGCAAGG
 CTGAAACTCAAAGGAATTGACAGGGCCCGCACAAGCGGTGGAGCATGTGG
 TTTAATTGAGCAACCGAAGAACCTTACCAAGGTCTGACATCCTCTGAAA
 ACCCTAGAGATAGGGCTTCTCCCTCGGGAGCAGAGTGACAGGTGTGCATG
 GTTGTGTCAGCTCGTGTGGAGATGTTGGTTAGTCCGCACGAGCGCAC
 CCTGATCTAGTTGCCATT CATAGTGGCACTCTAGGTGACTGCCGTTGACAACC
 GGAGCAGTGGATGGACCGCGCTCGAATTCCATCA

Strain NRCJ-39

TGAACTTGCGCAGGCCTAACACATGCAAGCTCGAGCGGATGAAGAGAGCTTG
 CTCTCTGATT CAGCGCGGACGGGTGAGTAATGCC TAGGAATTCGCCTGGTA
 GTGGGGACAACGTT CGAAAGGAACGCTAATACCGCATACGTCCTACGGGA
 GAAAGCAGGGACCTCAGGCTT GCGCTATCAGATGAGCCTAGGT CGGATT
 AGCTAGTTGGTGAGGTAATGGCTCACCAAGGCAGCAGATCCGTAACGGTCTG
 AGAGGATGATCAGTCACACTGGA ACTGAGACACGGTCCAGACTCCTACGGGA
 GGCAGCAGTGGGAATATTGGACAATGGCGAAAGCCTGATCCAGCCATGCC
 GCGTGTGAAGAAGGTCTCGGATTGTAAAGCACTTAAGTTGGAGGAAG
 GGTGTAGATTAATACTCTGCAATTGACGTTACCGACAGAATAAGCACCG
 GCTAACTCTGTGCCAGCAGCCCGTAATACAGAGGGTGCAAGCGTTAATCG
 GAATTACTGGCGTAAAGCGCGGTAGGTGGTTGTTAAGTTGGATGTGAAA
 GCCCCGGGCTCAACCTGGGA ACTGCAATTCAAAGACTGACAAGCTAGAGTATGG
 TAGAGGGTGGTGGAATT CCTGTAGCGGTGAAATGCGTAGATATAGGAAG
 AACACCACTGGCGAAGGC GACCACCTGGACTGATACTGACACTGAGGTGC
 GAAAGCGTGGGAGCAAACAGGATTAGATA CCCTGGTAGTCCACGCCGTAA
 ACGATGTCAACTAGCC GTGGAGCCTGAGCTCTTAGTGGCGCAGCTAACG
 CATTAAGTTGACCGCCTGGGAGTACGCCGCAAGGTTAAA ACTCAAATGAA
 ATTGACGGGGGCCGCACAAGCGGTGGAGCATGTGGTTAATTGAAAGCAAC
 GCGAAAGAACCTTACCAAGGCCTGACATCCAATGAACTTCCAGAGATGGAT
 TGGTGCCTTCGGAAACATTGAGACAGGTGCTGCATGGCTGTCAGCTC
 GTGTCCGTGGAGATGTTGGGTTAAGTCCGTAACGAGCGCAACGCTTGTCC
 AGT TACCAGCACTTAATGGTGGACTCTAAGGAAACTGCCGGATGACAAACC
 GGAAGCAGGTTGGGATGACCGGTACAGTTACATACATAGAGCACTCACATA
 TATACGAGGCACCACT

Strain NRCJ-40

GCTCAGAAGATGGTGAGAGTGGCGCACGGTGCGTAACCGTGAGCAACCT
 ACCTCTATCAGGGGATAGCCTCTCGAAAGAGAGATTAACACCGCATAACAT
 CAACAGTCGATGTTGGTGTAAATATTAGGATAGAGATGGCTCG
 CGTGACATTAGCTAGTTGGTAGGGTAACGGCTTACCAAGGCAGCAGATGTCTA
 GGGGCTCTGAGAGGAGAATCCCCCACACTGGTACTGAGACACGGACCAGACT
 CCTACGGGAGGCAGCAGTAAGGAATTGGTCAATGGCGGAAGCCTGAAC
 CAGCCATGCCCGTGCAGGATGACTGCCATGGTTGAAACTGCTTTGTC

CAGGAATAAACCCAGATACGTATCTGGCTGAATGTACTGGAAGAATAAGG
 ATCGGCTAACTCCGTGCCAGCCGCCGGTATACGGGGAAACCCAGCGTTA
 TCCCGATTATTGGGTTAAAGGGGTGCTTAGCCGGAAATTAAATTCAAGGGTG
 AAATACGGTGGCTCAACCATCGCAGTGCCTTGATCCTGATTGGCTGAATCC
 ATTGAAATTGGCCGGAAAAAAACAAGTA

Strain NRCJ-41

TGCGCTGCGCAGGCCTAACACATGCAAGTCGGCGGATGAAGAGAGCTGCTC
 TCTGATTCAAGCGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTG
 GGGGACAACGTTCGAAAGGAACGCTAACATACCGCATACGTCTACGGGAGAA
 AGCAGGGGACCTCGGGCCTGCGCTATCAGATGAGCCTAGGTGGATTAGC
 TAGTTGGTAGAGTAATGGCTACCAAGGCACGATCCGTAAGTGGCTGAGA
 GGATGATCAGTCACACTGGAACTGAGACACGGTCCAGACTCCTACGGGAGGC
 AGCAGTGGGAATATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCGCG
 TGTGTGAAGAAGGTCTCGGATTGTAAAGCACTTAAGTGGAGGAAGGGT
 TGTAGAT

Strain NRCJ-42

CTGCACCTGCGCATGCCTAACATGCAAGTCGAGCGGCAGCACGGCTTCTG
 CCTGTTGGCGAGTGGCGAACGGGTGAGTAATACATCTGAACGTGCCGTGTG
 TGGGGATAACAAGTCGAAAGATTAGCTGATACCGCATACGAACGTGATGGTG
 AAAGCGGGGACCGCATGGCTATGGCTACACGAGCGGCCGATGTCTGATTA
 GCTACTTGGTGGGTAAAAGCCTACCAACGCCACAATCAGCGCTGGTCCGA
 AAGGACGATCAACCACACTGGACTGAAACACGGACCAGACTCCTACCGGA
 GGCACCACTGGGAATTGGACAATGGGGCAACCGTACGCAACAAATGCC
 GCGTGTGAAGAAGGCCTCGGGTTGTAAGCACTTGTCCGGAAAGAAA
 TGGCCTGGGTGAATACCCGGTCGCTGCCGGTACCGGAACAATAAGCACCG
 GCTAACTACGTGCCAACTACCGCGTAATACGTAGGGTGCAGCGTTAATCA
 GAATTACTGGCGTAAAGCGTGCAGCGGTTTGTAAGACAGGGCGTAAA
 TCCCCGAGCTTAACGGATGGCGCTTGTGACTGATTGGCTACAGTATGTC
 TTGGGGCGTAGAATTCCACGTGAGCAGTGAAATGCGTATAGATGTCGAGGA
 ATACCGATGGCGCACGCATGCCCTGGCACGTCACTGACGCTCATGCGACAC
 AGCGTGGGAGCACACAGGATTAGATACCGTGGTAGTCCACGCCCTAACGA
 TGTCAACTACGATGGGATTATTCTCAGTTACCGTACCGTACGCGCT

GAAGTTGACCGGCCTGGGGAGTACGGTCGCAGGGATTAAAACCTCAAAGGTA
 AATGACTCGGGACCACGTACCAGCGCGCATGAATGTGGGTTAATTCTTGTG
 CATCGACGTAAAACCATACCCTACCCATTGATATGCCGGTCTATCGACATCAT
 ATAAGCATTCTGATGCCCTAAAGGGTAAACTTGTACAGCAGGATAGCTGC
 ATCGGCTTGTGTTCATCGTCTTGTCACTGAGATGTTGGGTTAAGATTCCCTGCG
 ATCAAGCTCCAACCACCTGTCTTAGCTGACTCTAAGATAACCCTACCAA
 ACAGAACTTGCCCCTGACAAAGCCGCAATGCAAGGTGGAGGAATGCGACG
 ATGCAGGTCGTGCATAGTACTTACAAGTTACGGGCTTCGATATTAATCGATG
 CTGTAGCGAGTACATGCGGAGTGCATTTGGCCAAGGTCGGACAGAAAAG

Strain NRCJ-43

CACCGTGGTAACCGTCCTCCGAAGGTTAGACTAGCTACTTCTGGTGCAACCC
 ACTCCCATGGTGTGACGGGCGGTGTACAAGGCCGGAACGTATTCACCG
 CGACATTCTGATTGCGATTACTAGCGATTCCGACTTCACGCAGTCGAGTTGC
 AGACTGCGATCCGGACTACGATGGTTTATGGGATTAGCTCCACCTCGCGC
 TTGGCAACCCTTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGTAAG
 GCCATGATGACTTGACGTACCCCCACCTCCTCCGGTTGTACCGGAGT
 CTCCTTAGAGTGCCCACCATTACGTGCTGGTAACTAAGGACAAGGGTTGCGC
 TCGTTACGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCCATG
 CAGCACCTGTCTCAATGTTCCGAAGGCACCAATCCATCTCTGGAAAGTTCAT
 TGGATGTCAAGGCCTGGTAAGGTTCTCGCGTTGCTTCGAATTAAACCACATG
 CTCCACCGCTTGTGGGGCCCCCGTCAATTGAGTTAACCTTGC
 CGTACTCCCCAGGCAGGTCAACTTAATGCGTTAGCTGCCACTAAGAGCTCA
 AGGCTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGGTATC
 TAATCCTGTTGCTCCCCACGCTTCGCACCTCAGTGTCACTAGTCCAGG
 TGGTCGCCCTCGCCACTGGTGTCCCTATATCTACGCATTACCGCTACA
 CAGGAAATTCCACCAACCCTTACCCATACTCTAGCTGCCAGTTGGATGCAG
 TTCCCAGGTTGAGCCGGGATTTCACATCCAACCTAACGAACCACCTACGC
 GCGCTTACGCCAGTAATTCCGATTAACGCTTGCACCCTGTATTACCGCG
 GCTGCTGGCACAGAGTTAGCCCAGGTG

Strain NRCJ-44

AGGCCTAACATGCAGTCGAGCGGATGAAGGGAGCTTGCTCCTGAATTCA
 GCGGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGACAACGT

TTCGAAAGGAACGCTAATACCGCATACGCCCTACGGGAGAAAGCAGGGGAC
 CTTCGGGCCTTGCCTATCAGATGAGCCTAGGTGGATTAGCTAGTTGGTGAG
 GTAATGGCTCACCCAGGGCACCATCCCTAACTGGTCCGAAAGGATTATCAAT
 TCACTGGAAATTAACACGGGTCAAAAACCTCCCAGGAGGGCACCAGGTGCGAA
 AAATTGGACAAAAGGGTTATCTGCCCGATGCCCATGTGTGAAATGCTCT
 TCTGACTGAACAGACTTAAATTGGAAAGAAAGGTTACGATAAAACTGTGCT
 ATCTTCCCCTACCCAACCGAAAGAACCCCTGCTTATTCGTAAGTACTGCCGC
 GGTAAAACAGAGGGTGCATGCCTAATCTGAATTACTGGCGTAAAGCGCGC
 GTCGGTGGTCGTTAAGTTGGTTAAAATCCCCGTTCAAACCCCTGGGAACCT
 GCATCCAAAAGCTGGCAAGCTTAAGTCTTACAGCGTCACTAGA

Strain NRCJ-45

TCGAGCGAACTGGATTAGAGAGCTGCTCTCAAGAAGTTAGCGGCGGACGGG
 TGAGTAACACGTGGTAACCTGCCATAAGACTGGATAACTCCGGAAACC
 GGGGCTAATACCGATAATATTGAACTGCATGGTCGAAATTGAAAGGCG
 GCTTCGGCTGTCACTTATGGATGGACCCGCGTCGCATTAGCTAGTTGGTGAGG
 TAACGGCTACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGTAGCGGC
 ACACGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTAGGGA
 ATCTTCCGCAATGGACGAAAGTCTGACGGAGCAACGCCCGTGAGTGAA
 GGCTTCGGTCGTTAAACTCTGTTAGGAAAGAACAAAGTGTAGTTGAA
 TAAGCTGGCACCTGACGGTACCTAACAGAAAGCCACGGCTAACTACGTGC
 CAGCAGCCCGGTAATACGTAGGTGGCAAGCGTTATCCGAATTATTGGCG
 TAAAGCGCGCGAGGTGGTTCTTAAGTCTGATGTGAAAGCCCACGGCTAA
 CGTGGAGGGTATTGAAACTGGAGACTTGAGTGAGAACAGAGGGAAAGTG
 GAATTCCATGTGTAGCGGTAAATGCGTAGAGATATGGAGGAACACCAGTGG
 CGAAGGCGACTTCTGGTCTGTAACGACACTGAGGCGCGAAAGCGTGGGA
 GCAAACAGGATTAGATACCCCTGGTAGTCCACGCCGTAAACGATGAGTGCTAA
 GTGTTAGAGGGTTCCGCCCTTAGTGCTGAAGTTAACGCATTAAGCACTCCG
 CCTGGGAGTACGCCGCAAGGCTGAACCAAAGGAATTGACGGGGCCCG
 CACAAGCGGTGGAGCATGTGGTTAACCGAAGCAGCGAAGAACCTTACCA
 GGTCTGACATCCTCTGAAACCCAAGAGATAGAGCTTCGTCCTCCT

Strain NRCJ-46

GTCCCCTCGCCCTCCCCCGGAAACCGGTTGGGCCATGAGCTCGGGTGT
 CCAACTTCGTGACTTGACGGCGGTGTACAAGGCCGGAACGTATTCA
 CCGCAGCGTTGCTGATCTCGGATTACTAGCGACTCCGACTTCATGGGTCGAG
 TTGCAGACCCCAATCCGAAC TGAGACCGGCTTTGGGATTGCTCCACCTTA
 CGGTATCGCAGCCCTCTGTACC GGCCATTGTAGCATGCGTAAGCCCAAGAC
 ATAAGGGGCATGATGATTGACGTACCCACCTCCTCCGAGTTGACCCCG
 GCAGTCTCCATGAGTCCCCGGCATAACCGCTGGCAACATAGGACGAGGGT
 TCGCTCGTTGCGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAA
 CCATGCACCACCTGTGCACCGACCTAACGGGACCCATCTGAGTTGCG
 GGTGCATGTCAAGCCTGGTAAGGTTCTCGCGTTGCATCGAATTAAATCCGCA
 TGCTCCGCCGTTGTGCAGGGCCCCGTCAATTCTTGAGTTAGCCTTGCG
 GCCGTACTCCCCAGGCGGGGCACTTAATGCGTTAGCTGCGGCACGGAACTCG
 TGGAATGAGCCCCACACCTAGTGCCAACGTTACGGCATGGACTACCAGGG
 TATCTAATCCTGTTGCTCCCATGCTTCGCTCCTCAGCGTCAGTTGCGGCC
 AGAGACCTGCCATCGGTGTTCCCTGATATCTGCGCATTCCACCGC
 TACACCAGGAATTCCAGTCTCCCTACCGCACTTAGTCTGCCC
 GCAAGCCGAGGTTGAGCCTCGGGATTTCACACCAAGACGCGACAGACCGCCT
 ACGAGCTTTACGCCAATAATTCCGGACAAGCGCTGCGCCCTACGTATT
 ACCGCGGCTGCTGGCACGTAGTTAGCC

Strain NRCJ-47

GTGGCGCGTGCCTAATACATGCAAGTCGAGCGAATGGATATAGAGAGCTTGC
 TCTCAAGAAGTTAGCGGCGGACGGGTGAGTAACACGTGGTAACCTGCCAT
 AAGACTGGGATAACTCCGGAAACCGGGCTAATACCGGATAACATTGAA
 CCGCATGGTTGAAATTGAAAGGCGGCTCGGCTGTCACTTATGGATGGACC
 CGCGTCGCATTAGCTAGTTGGT GAGGTAACGGCTACCAAGGCAACGATGCG
 TAGCCGACCTGAGAGGGTGATCGGCCACACTGGACTGAGACACGGCCAG
 ACTCCTACGGGAGGCAGCAGTAGGGAATCTTCTCAATGGACGAAAGTCTGA
 CGGAGCAACGCCCGT GAGTGATGAAGGCTTGGGTGTAAGCTGTTG
 TTAGGGAAGAACAGTGCCTGTTGAATAAGCTGGCCCTGGCCGGTCCCAA
 ACCCAAAGCCCACGCTAACCTCCTGCCCTCCGCCGGTGTCCCTTA

Strain NRCJ-48

AAGAGCTTGCCTTCGATTCAGCGCGGACGGGTGAGTAATGCCTAGGAATC
 TGCCTGGTAGTGGGGACAACGTTCGAAAGGAACGCTAATACCGCATACGT
 CCTACGGGAGAAAGCAGGGACCTTCGGGCCTGCGCTATCAGATGAGCCTA
 GGTGGATTAGCTAGTGGTGGGTAAATGGCTACCAAGGCACGATCCGTA
 ACTGGTCTGAGAGGATGATCAGTCACACTGGAACGTGAGACACGGTCCAGACT
 CCTACGGGAGGCAGCAGTGGGAATATTGGACAATGGCGAAAGCCTGATC
 CAGCCATGCCCGTGTGAAGAAGGTCTCGGATTGTAAAGCACTTAAGT
 TGGGAGGAAGGGCAGTAAGCGAACCTTGCTGTTGACGTTACCGACAGA
 ATAAGCACCGCTAACTCTGTGCCAGCAGCCCGTAATACAGAGGGTCAA
 GCGTTAACGGAATTACTGGCGTAAAGCGCGCTAGGTGGTTGTTAAGTT
 GGATGTGAAAGCCCCGGCTAACCTGGGAACACTGCATCCAAAAGCAGC
 TAGAGTACGGTAGAGGGTGGTGAATTCCCTGTGTAGCGGTGAAATGCGTAG
 ATATAGGAAGGAACACCAGTGGCGAACGACCTGGACTGATACTGAC
 ACTGAGGTGCGAAAAGCGTGGGAGCAAACAGGATTAGATAACCTGGTAGT
 CCCACGCCGTAAACGATGTCAACTAACCGTTGGAATCCTGAGATTTATGTG
 GCGCAGCTAACGCTTAAGTTGACCGCCCCGGGGTACGGGCCGCGTTA
 AAAACACAAAATGAATTGAAAGGGGGCCGCACAAACAGATGGAGCAGGT
 GTTTTAGTTCTAATTACAACAAAAACCTAACCGCCCTTGACTTCTAGGG
 GCTTGCAGAGATAGAATGGGGCTTTGGCATTGTGAACACGTGAAAGAG
 CAGCATGTAGAGCTATCATCACACCGATACGCTGGAAATAGTGCAGTGAACA
 TCGACACACCAACGGTCACGCACTATACACAAGTGTCCGGCAGACGTATGTG
 TGTGTGGCAGCATTGATCAGTGACA

Strain NRCJ-49

CACCGTGGTAACCGTCCTCCGAAGGTTAGACTAGCTACTTCTGGTGCAACCC
 ACTCCCATGGTGTGACGGCGGTGTACAAGGCCGGAACGTATTACCG
 CGACATTCTGATTGCGATTACTAGCGATTCCGACTTCACGCAGTCAGTTGC
 AGACTGCGATCCGGACTACGATCGGTTGTGAGATTAGCTCCACCTCGCGGC
 TTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGTAA
 GGGCCATGACTTGACGTCATCCCCACCTCCTCCGGTTGTACCGGCAG
 TCTCCTTAGAGTGCCACCATAACGTGCTGGTAACTAAGGACAAGGGTTGCG
 CTCGTTACGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCCAT
 GCAGCACCTGTGTAGAGTTCCGAAGGCACCAATCCATCTGGAAAGTTC

TCTGCATGTCAAGGCCTGGTAAGGTTCTCGCGTGCTCGAATTAAACCACA
TGCTCCACCGCTTGTGCAGGGCCCCGTCAATTGAGTTAACCTGCG
GCCGTACTCCCCAGGCGGTCAACTTAATCGTTAGCTGCGCCACTAAAATCTC
AAGGATTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGGTAT
CTAACCTGTTGCTCCCACGCTTCGACCTCAGTGTCACTACAGTCCAG
GTGGTCGCCCTCGCCACTGGTGTTCCTCCTATATCTACGCATTCAACCGCTAC
ACAGGAAATTCCACCACCCCTACCGTACTCTAGCTCGCCAGTTGGATGCA
GTTCCCAGGTTGAGCCCCGGGCTTCACATCCAACCTAACGAACACACCTAC
GCGCGCTTACGCCAGTAATTCCGATTAACGCTGCACCCCTGTATTACC
GCGGCTGCTGGCACAGAGTTAGCCC

Strain NRCJ-50

CACCCGTGGTAACCGTCCTCCGAAGGTTAGACTAGCTACTTCTGGTGCAACC
CACTCCCATGGTGTGACGGCGGTGTACAAGGCCGGAACGTATTACC
GCGACATTCTGATTGCGATTACTAGCGATTCCGACTCACGCAGTCGAGTTG
CAGACTGCGATCCGGACTACGATCGGTTTGTGAGATTAGCTCCACCTCGCG
CTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTAGCCAGGCCGTA
AGGGCCATGATGACTTGACGTCATCCCCACCTCCTCCGGTTGTCACCGGCA
GTCTCCTTAGAGTGCCCACCATAACGTGCTGGTAACTAAGGACAAGGTTGC
GCTCGTTACGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCCA
TGCAGCACCTGTGTCAGAGTTCCGAAGGCACCAATCCATCTCTGGAAAGTT
CTCTGCATGTCAAGGCCTGGTAAGGTTCTCGCGTGCTCGAATTAAACCAC
ATGCTCCACCGCTTGTGCAGGGCCCCGTCAATTGAGTTAACCTGCG
GGCCGTACTCCCCAGGCGGTCAACTTAATCGTTACGGCGTGGACTACCAGGGT
TCAAGGATTCCAACGGCTAGTTGACATCGTTACGGCGTGGACTACCAGGGT
ATCTAACCTGTTGCTCCCCACGCTTCGACCTCAGTGTCACTACAGTCC
AGGTGGTCGCCACTGGTGTCCCTATATCTACGCATTCAACCGCT
ACACAGGAAATTCCACCAACCCCTACCGTACTCTAGCTCGCCAGTTGGATG
CAGTTCCCAGGTTGAGCCCCGGGCTTCACATCCAACCTAACGAACACACCTA
CGCGCGCTTACGCCAGTAATTCCGATTAACGCTGCACCCCTGTATTACC
GCGGCTGCTGGCACAGAGTTAGCCC

Strain NRCJ-51

AGGAAGCTTGCACCGTGCCTAATACATGCAAGACGAGCGAATTGGATTAGAGA
 GCTTGCTCTCAAGAACGTTAGCGGGGACGGGTGAGTAACACGTGGTAACCT
 GCCCATAAGACTGGATAACTCCGGAAACCGGGCTAATACCGGATAATAT
 TTTGAACACTGCATGGTCGAAATTGAAAGGGCTCGGCTGTCACTTATGGAT
 GGACCCCGCGTCGATTAGCTAGTTGGTGGAGGTAACGGCTCACCAAGGCAACG
 ATGCGTAGCCGACCTGAGAGGGTGGATCGGCCACACTGGGACTGAGACACGG
 CCCAGACTCCTACGGGAGGCAGCAGTAGGAAATCTTCCGCAATGGACGAAA
 GTCTGACGGAGCAACGCCCGTGAGTGATGAAGGCTTCGGTGTGAAACT
 CTGTTGTTAGGAAAGAACAAAGTGTAGTTGAATAAGCTGGCACCTGACGGT
 ACCTAACCAAGAACGCCACGGCTAACTACGTGCCAGCAGCCGCGTAATACGT
 AGGTGGCAAGCGTTATCCGAATTATTGGCGTAAAGCGCGCAGGTGGTT
 TCTTAAGTCTGATGTGAAAGCCCACGGCTCAACCGTGGAGGGTCATTGGAAA
 CTGGGAGACTTGAGTCAGAAGAGGAAAGTGGATTCCATGTGTAGCGGTGA
 AATGCGTAGAGATATGGAGGAACACCAGTGGCGAAGGCGACTTCTGGTCTG
 TAACTGACACTGAGGCGCGAACAGCTGGGGAGCAAACAGGATTAGATAACCC
 TGGTAGTCCACGCCGAAACGATGAGTGCTAAGTGTAGAGGGTTCCGCC
 TTTAGTGCTGAAGTTAACGCATTAAGCACTCCGCCTGGGAGTACGGCCGCA
 AGGCTGAAACTCAAAGGAATTGACGGGGCCCGACAAGCGGTGGAGCATGT
 GGTAAATTGCAAGCACCGAAGACCCCTTACCAAGGTCTGACATCCTCTGAA
 AACCCCTAGAGATAGGGCTCTCCTCGGGAGCAGAGTGACAGTGGTGCATGT
 TGTCGTCAGCTCGTGTGAGATGTTGGTTAGTCCGCACGAGCGCACCC
 GATCTTAGTTGCATCATAGTAGTCACTCTAACGTGACTGCCGTGACAACGAGA
 AGGTTGGATTGACGTCAAATCCATCATGTGACCT

Strain NRCJ-52

CCTACCGCACACCGTGGCAAGCGCCCTCCGAAGGTTAACGCTACCTGCTTCT
 GGTGCAACAAACTCCCATGGTGTGACGGCGGTGTACAAGGCCGGAAAC
 GTATTCACCGCAGCAATGCTGATCTGCGATTACTAGCGATTCCGACTTCATGG
 AGTCGAGTTGCAGACTCCAATCCGGACTGAGATAGGGTTCTGGGATTGGCT
 TACCGTCGCCGGCTGCAGCCCTCTGTCCTACCATTGTAGTACGTGTGAGC
 CCTGGCCGTAAGGGCCATGATGACTGACGTCACTCCCCACCTCCTCCGGTT
 GTCACCGGGCGGTCTCCTAGAGTCCCACCATTACGTGCTGGCAACTAAGGA
 CAAGGGTTGCGCTCGTGCAGGACTTAACCCAACATCTCACGACACGAGCTG

ACGACAGCCATGCAGCACCTGTGTCGAGTTCCGAAGGCACCAATCCATCT
 CTGGAAAGTTCTCGACATGTCAAGGCCAGGTAAGGTTCTCGCGTTGCATCG
 AATTAAACCACATACTCCACCGCTTGTGGGGCCCCGTCAATTCTTGAGT
 TTCAGTCTTGCACCGTACTCCCCAGGCCAGGCGAACCTAACCGCTTAGCTCGA
 TACTGCGTGCCTAACATTGCACCCAACATCCAGTTGCATCGTTAGGGCGTGGA
 CTACCAAGGGTATCTAATCCTGTTGCTCCCCACGCTTGTGCCTCAGTGTCA
 GTGTTGGTCCAGGTAGCTGCCTTCGCCATGGATGTTCCCTGATCTCTACGC
 ATTCACTGCTACACCAGGAATTCCGCTACCCCTACCAACACTCTAGTCGTC
 AGTTTCCACTGCAGTTCCCAGGTTGAGCCCAGGGCTTCACAACAGACTAAA
 CGACCACCTACGCACGCTTACGCCAGTAATTCCGAGTAACGCTGCACCC
 TCGTATTACCGCGGCTGCTGGCACGAAGTTAGCCCAGGCTTATTCTTGGGT
 ACCGTCATCACCAACCCGGGTATTAACCCAGCTGCATTCTTCCCACAAAAA

Strain NRCJ-53

ATACACCGTGGTAACCGTCCTCCGAAGTTAAACTAGCTACTTCTGGTGCAAC
 CCACTCCCATGGTGTGACGGCGGTGTGTACAAGGCCGGAACGTATTAC
 CGCGACATTCTGATTGCGATTACTAGCGATTCCGACTTCACGCAGTCCAGTT
 GCAGACTGCGATCCGGACTACGATCGGTTGTGGATTAGCTCCACCTCGCG
 GCTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTAGCCCAGGCCGT
 AAGGGCCATGATGACTTGACGTACCCACCTAACGTGCTGGTAACTAAGGACAAGGGTTG
 CGCTCGTTACGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCC
 ATGCAGCACCTGTCAGTGTCCGAAGGCACCAAACCATCTCTGGTAAGTT
 CACTGGATGTCAAGGCCTGGTAAGGTTCTCGCGTTGCTTCAATTAAACCAC
 ATGCTCCACCGCTTGTGGGGCCCCGTCAATTGAGTTAACCTTGC
 GGCGTACTCCCCAGGCCAGTCAACTAACGCTTAGCTGCGCCACTAAATC
 TCAAGGATTCCAACGGCTAGTTGACATCGTTACAGCGTGGACTACCAGGGT
 ATCTAACCTGTTGCTCCCCACGCTTGCACCTCAGTGTCACTGAGGCC
 AGGTGGTCGCCCTTC

Strain NRCJ-54

GCAGCAGGCCTAACACATGCAAGTCGAGCGGATGAAGGGAGCTTGCTCCTGA
 ATTCAAGCGGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGG
 ACAACGTTCGAAAGGAACGCTAACACCGCATACGTCCACGGGAGAAAGCA

GGGGACCTCGGGCCTGCGCTATCAGATGAGCCTAGGTGGATTAGCTAGT
 TGGTGAGGTAATGGCTACCAAGGCGACGATCCGTAACTGGTCTGAGAGGAT
 GATCAGTCACACTGGAACGTGAGACACGGTCCAGACTCCTACGGGAGGCAGCA
 GTGGGAAATATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCCGTGTG
 TGAAGAAGGTCTCGGATTGTAAAGCACTTAAGTTGGGAGGAAGGGTTGTA
 GATTAATACTCTGCAATTGACGTTACCGACAGAATAAGCACCGGCTAACCTC
 TGTGCCAGCAGCCCGGTAATACAGAGGGTGCAAGCGTTAACCGGAAATTACT
 GGGCGTAAAGCGCGTAGGTGGTCGTTAAGTTGGATGTGAAATCCCCGGG
 CTCAACCTGGAACTGCATCCAAAACGGCGAGCTAGAGTATGGTAGAGGGT
 GGTGGAATTCCCTGTGTAGCGGTGAAATCGTAGATATAGGAAGGAACACCA
 GTGGCGAAGGCGACCACCTGGACTGATACTGACACTGAGGTGCGAAAGCGTG
 GGGAGCAAACAGGATTAGATAACCGCTGGTAGTCCACGCCGAAACGATGTCAA
 CTAGCCGTTGGGAGCCTTGAGCTCTAGGGCGCAGCTAACGCATTAAGTTG
 ACCGCCTGGGAGTACGGCCGCAAGGTTAAACTCAAATGAATTGACGGGG
 GCCCGCACAAGCGGTGGAGCATGTGGTTAATTGAAAGCAACCGGAAGAAC
 TTACCAAGGCCTGACATCCAATGAACCTTCAGAGATGGATTGGTGCCTTCG
 GGAACATTGAGACAGGTGCTGCATGGCTGTCAGCTCGTGTGAGATG
 TTGGGTTAAGTCC

Strain NRCJ-55

CGGCACCTGCGCGTGTAAACATGCAAGTCGAACGATGATCAGGAGCTTG
 CTCCTGTGATTAGTGGCGAACGGGTGAGTAACACGTGAGTAACCTGCCCTG
 ACTCTGGGATAAGCGTTGGAAACGACGTCTAATACTGGATATGATCACTGGC
 CGCATGGTCTGGTGGGGAAAGATTTGGTTGGGATGGACTCGCGGCCT
 ATCAGCTTGGTGGTAGGTAATGGCTACCAAGGCGACGACGGTAGCCGGC
 CTGAGAGGGTGACCGGCCACACTGGACTGACACACCGGCCACACTCCTACG
 GGAGGCAGCAGTGGGAATATTGACACAATGGCGAACGCGTGTGAGCAA
 CGCCGCGTGAGGGATGACGGCCTCGGGTTGAAACCTTTAGTAGGGAA
 GAAGCGAAAGTGACGGTACCTGCAGAAAAAGCACCGGCTAACTACGTGCCA
 GCAGCCGCGTAATACGTAGGGTGCAAGCGTTGTCGGAAATTATTGGCGTA
 AAGAGCTCGTAGGCGGTTGTCGCGTCTGCTGTGAAATCCGAGGCTCAACC
 TCGGGCTTGCAGTGGGTACGGCAGACTAGAGTGCAGGTAGGGAGATTGGA
 ATTCCCTGGTGTAGCGGTGGAATGCGCAGATATCAGGAGGAACACCGATGGCG
 AAGGCAGATCTCTGGCCGTAACGTGACGCTGAGGAGCGAAAGCGTGGGAG
 CGAACAGGATTAGATAACCGCTGGTAGTCCACGCCGAAACGTTGGCGCTAGA

TGTAGGGACTTCCACGGTTCTGTGTCGTAGCTAACGCATTAAGCGCCCCG
 CCTGGGGAGTACGGCCGCGAGGCTAAAACCTCAAAGGAATTGACGGGGGCC
 GCACAAGCGGCGGAGCATCGGATTAATCGATGCAACCGAAGAACCTTAC
 CAAGGCTTGACATACACCGAAACGGCCAGAGATGGTCGCCCCCTGTGGTC
 GTGTACAGGTGGTGCATGGTTGTCGTAGCTCGTGTGAGATGTTGGT
 AAGTCCCAGCAACGAGCGAACCCCTCGTATCTATGTTAGCCAGCTGCGTTAT
 GGCTGGCACTCATAGGAGACTGCCGGTCCAGCTGGAGCAGTGGGGATG
 ACGGTCAAATTCCATGGCCCTTAGTCCTGGCTCACACGCCATGGC
 TTAACACTGGTGGCCG

Strain NRCJ-56

TTGAAATGCGCAGGCCTAACACACTGCAAGCTCGAGCGGATGACGGGAGCTT
 TGCTCCTCTGATTCAAGCGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTG
 GTAGTGGGGACAACGTTCGAAAGGAACGCTAACCGCATACTGCCTACG
 GGAGAAAGCAGGGGACCTTCGGGCCTGCGCTATCAGATGAGCCTAGGTCGG
 ATTAGCTAGTTGGTGGAGGTAATGGCTACCAAGGCGACGATCCGTAAGTGGT
 CTGAGAGGATGATCAGTCACACTGGAACGTGAGACACGGTCCAGACTCCTACG
 GGAGGCAGCAGTGGGAATTGGACAATGGCGAAAGCCTGATCCAGCCA
 TGCCCGTGTGAAGAAGGTCTCGGATTGTAAGCACTTAAGTTGGAG
 GAAGGGCAGTAAGCTAACCTTGCTGTTGACGTTACGACAGAATAAGC
 ACCGGCTAACTCTGTGCCAGCAGCCCGGTAAACAGAGGGTGAAGCGTTA
 ATCGGAATTACTGGCGTAAAGCGCGTAGGTGGTTGTTAAGTTGGATGT
 GAAAGCCCCGGGCTAACCTGGAAACTGCATCCAAAACGGCGAGCTAGAGT
 ACGGTAGAGGGTGGGAAATTCTGTAGCGGTGAAATGCGTAGATATAG
 GAAGGAACACCAGTGGCGAAGGCACCTGGACTGATACTGACACTGAG
 GTGCGAAAGCGTGGGAGCAAACAGGATTAGATACCGTGTAGTCCACGCCG
 TAAACGATGTCAACTAGCCGTTGGAATCCTGAGATTAGTGGCGCAGCTA
 ACGCATTAAGTTGACCGCCTGGGAGTACGGCCGCAAGGTTAAAACCTAA
 GAATTGACGGGGGCCGCACAAGCGGTGGAGCATGTGGTTAATTGAAAGCA
 ACGCGAAGAACCTTACCAAGGCCTGACATGCAGAGAACCTTCCAGAGATGGA
 TTGGTGCCTTCGGAACTCTGACACAGGTGCTGCATGGCTGTCAGCTCG
 TGTGAGATGTTGGTTAGGCTAACGAGCGCAACCCTGTCCCTAGT
 TAACCAAGCAGTTATGGTGGGACTCTAGAGACTGCAGTGACAACCCGGAG
 GAGTTGGGATGACGTTAGTCAGTCATGGACCTTACCGACCTGGCCTACAC
 ACACGTGGACTC

Strain NRCJ-57

TCGAACGGTAACAAGGAAGAAGCTTGCTGCTTGCTGACGAAGTGGCGGACG
 GGTGAGTAATGTCTGGAAACTGCCTGATGGAGGGGATAACTACTGGAAAC
 GGTAGCTAATACCGCATAACGTCGCAAGACCAAAGAGGGGGACCTCGGGC
 CTCTGCCATCGGATGTGCCAGATGGGATTAGCTTAGGTGGGTAACGG
 CTCACCTAGGCGACGATCCCTAGCTGGTCTGAGAGGATGACCAGCCACACTG
 GAACTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGTGGGAATATTGC
 ACAATGGCGCAAGCCTGATGCAGCCATGCCCGTGTATGAAGAAGGCCTC
 GGGTTGAAAGTACTTCAGCGGGAGGAAGGGAGTAAAGTTAACCTTG
 CTCATTGACGTTACCCGCAAGAAGCACCAGCTACTCCGTGCCAGCAGCC
 CGCGTAATACGGAGGGTGCAAGCGTTAACCGAATTACTGGCGTAAAGCGC
 ACGCAGGCGTTGTTAACGTAGATGTGAAATCCCCGGCTAACCTGGAA
 CTGCATCTGATACTGGCAAGCCTGAGTCTCGTAGAGGGGGTAGAATTCCAG
 GTGTAGCGGTGAAATGCGTAGAGATCTGGAGGAATACCGGTGGCGAAGGCG
 GCCCCCTGGACGAAGACTGACGCTCAGGTGCGAAAGCGTGGGAGCAAACA
 GGATTAGATAACCTGGTAGTCCACGCCGTAACCGATGTCGACTGGAGGTTG
 TGCCCTGAGGCGTGGCTTCCGGAGCTAACCGTTAACGCTGACCGCCTGGGG
 AGTACGGCGCAAGGTTAAACTCAAATGAATTGACGGGGCCGCACAAG
 CGGTGGAGCATGTGGTTAACCGATGCAACCGAAGAACCTTACCTGGTCTT
 GACATCCACGGAAGTTTCAGAGATGAGAATGTGCCTCGGGAAACC

Strain NRCJ-58

AGGCCTAACACATGCAAGTCGAGCGGCAGCACGGTACTTGTACCTGGTGGC
 GAGCGGCGGACGGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGATA
 ACGCTGGAAACGGACGCTAACCGCATACTGGCTACGGGAGAAAGCAGG
 GGACCTTCGGGCTTGCCTACAGATGAGCCTAGGTGGATTAGCTAGTTG
 GTGAGGTAATGGCTACCAAGGCGACGATCCGTAACGGTCTGAGAGGATGA
 TCAGTCACACTGGAACGTGAGACACGGTCCAGACTCCTACGGGAGGCAGCAGT
 GGGGAATATTGGACAATGGCGAAAGCCTGATCCAGCCATGCCCGTGTGTG
 AAGAAGGTCTCGGATTGTAAGCACTTAAGTTGGAGGAAGGGATTAC
 CTAATACGTAAGTGTGTTGACGTTACCGACAGAATAAGCACCAGCTAACCT
 GTGCCAGCAGCCCGGTAAATACAGAGGGTGCAAGCGTTAACCGAATTACTG
 GCGTAAAGCGCGCGTAGGTGGTCTGTTAACGTTGGATGTGAAATCCCCGGC
 TCAACCTGGAACTGCATTAAAACGTGAGCTAGAGTATGGTAGAGGGTG

GTGGAATTCTGTAGCGGTGAAATGCGTAGATATAGGAAGGAACACCAAG
TGGCGAAGGCAGCACCTGGACTGATACTGACACTGAGGTGCGAAAGCGTGG
GGAGCAAACAGGGTTATATACCCTGGTAGTCCACGCCGTAACAAATGTCTAC
TAGCTCTCCGGAGCCAAGAAATTACTCGCGCAGCCTACCACATTCAATTGA
CCGATTGGTGGGGTGCAGGCCCCGGGGAGTTGTGCTTAAATTACAGCACGG
GGCGAGCCCAACCCCAGGCAGGCGGGACTTGTGCTTAAATTACAGCACGG
GCCGGACCCCTGCCTGGACCTCCGACAACCAATGAACGATAACAGACACCGGA
TTTTGTCTGTCAGGAAAATAAGAAAAAACATTACATCACAGACGAACACCC
CCTCTGCTGCCCTGGATATTGTTGGGTTAACCTCCCCCCAACACGTCGCC
GAGCGC

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