Table S3. 122 *Oryza* miRNA loci.

family	gene	chromosome	begin	end	strand	miRNA sequence
156	156a	Contig13033	1493	1513	-	UGACAGAAGAGAGUGAGCACA
156	156b	Contig15626	3509	3529	-	UGACAGAAGAGAGUGAGCACA
156	156c	Contig15626	3900	3920	-	UGACAGAAGAGAGUGAGCACA
156	156d	Contig274	4663	4683	-	UGACAGAAGAGAGUGAGCACA
156	156e	Contig5039	6201	6221	-	UGACAGAAGAGAGUGAGCACA
156	156f	Contig439	18369	18389	+	UGACAGAAGAGAGUGAGCACA
156	156g	Contig1967	15322	15342	-	UGACAGAAGAGAGUGAGCACA
156	156h	Contig6735	1527	1547	+	UGACAGAAGAGAGUGAGCACA
156	156i	Contig62236	952	972	+	UGACAGAAGAGAGUGAGCACA
156	156j	Contig69405	187	207	-	UGACAGAAGAGAGUGAGCACA
156	156k	Contig5213	8914	8934	-	UGACAGAAGAGAGAGAGCACa
156	156l	Contig6603	2305	2325	-	CGACAGAAGAGAGUGAGCAUA
159/JAW	159a	Contig2393	7117	7137	-	UUUGGAUUGAAGGGAGCUCUG
159/JAW	159b	Contig3622	3590	3610	-	UUUGGAUUGAAGGGAGCUCUG
159/JAW	159c	Contig25407	2212	2232	-	AUUGGAUUGAAGGGAGCUCCA
159/JAW	159d	Contig12045	1208	1228	-	AUUGGAUUGAAGGGAGCUCCg
159/JAW	159e	Contig22151	1752	1772	+	AUUGGAUUGAAGGGAGCUCCU
159/JAW	159f	Contig11829	4932	4952	-	CUUGGAUUGAAGGGAGCUCUA
159/JAW	319a	Contig7071	3206	3225	+	UUGGACUGAAGGGUGCUCcc
159/JAW	319b	Contig21652	4082	4101	+	uuggacugaagggugcuccc
160	160a	Contig42	30830	30850	+	UGCCUGGCUCCCUGUAUGCCA
160	160b	Contig12962	3751	3771	+	UGCCUGGCUCCCUGUAUGCCA
160	160c	Contig5072	6548	6568	-	UGCCUGGCUCCCUGUAUGCCA
160	160d	Contig278	19015	19035	+	UGCCUGGCUCCCUGUAUGCCA
160	160e	Contig3516	2475	2495	+	UGCCUGGCUCCCUGUAUGCCG
160	160f	Contig7096	4672	4692	+	UGCCUGGCUCCCUGAAUGCCA
162	162a	Contig2445	8502	8522	-	UCGAUAAACCUCUGCAUCCAG
162	162b	Contig51	11868	11888	+	UCGAUAAGCCUCUGCAUCCAG
164	164a	Contig11673	2054	2074	+	UGGAGAAGCAGGGCACGUGCA
164	164b	Contig14283	4403	4423	+	UGGAGAAGCAGGGCACGUGCA
164	164c	Contig36005	1290	1310	+	UGGAGAAGCAGGGUACGUGCA
164	164d	Contig4937	9146	9166	-	UGGAGAAGCAGGGCACGUGCU
164	164e	Contig15421	324	344	+	UGGAGAAGCAGGGCACGUGAG
166	166a	Contig4452	2329	2349	-	UCGGACCAGGCUUCAUUCCCC
166	166b	Contig2319	1100	1120	-	UCGGACCAGGCUUCAUUCCCC
166	166c	Contig1481	3013	3033	-	UCGGACCAGGCUUCAUUCCCC
166	166d	Contig9514	1093	1113	+	UCGGACCAGGCUUCAUUCCCC
166	166e	Contig7441	4253	4273	+	UCGGACCAGGCUUCAUUCCCC
166	166f	Contig1210	18646	18666	-	UCGGACCAGGCUUCAUUCCCC
166	166g	Contig10486	2698	2718	+	UCGGACCAGGCUUCAUUCCUC
166	166h	Contig38876	1710	1730	+	UCGGACCAGGCUUCAUUCCUC
166	166i	Contig3185	377	397	+	ucggaucaggcuucauuccuc
166	166j	Contig75985	6	26	-	ucggaucaggcuucauuccuc
166	166k	Contig38876	1881	1901	+	UCGGACCAGGCUUCAAUCCCU
166	166l	Contig7215	4788	4808	+	UCGGACCAGGCUUCAAUCCCU
167	167a	Contig1917	10902	10922	-	UGAAGCUGCCAGCAUGAUCUA
167	167b	Contig5097	1075	1095	-	UGAAGCUGCCAGCAUGAUCUA
167	167c	Contig981	7221	7241	+	UGAAGCUGCCAGCAUGAUCUA
167	167d	Contig1	16472	16492	-	UGAAGCUGCCAGCAUGAUCUG
167	167e	Contig10736	6185	6205	-	UGAAGCUGCCAGCAUGAUCUG
167	167f	Contig15194	2195	2215	-	UGAAGCUGCCAGCAUGAUCUG

167	167g	Contig15604	3032	3052	-	UGAAGCUGCCAGCAUGAUCUg
167	167h	Contig1917	7095	7115	-	UGAAGCUGCCAGCAUGAUCUG
167	167i	Contig3960	12432	12452	-	UGAAGCUGCCAGCAUGAUCUG
168	168a	Contig5181	6314	6334	+	UCGCUUGGUGCAGAUCGGGAC
168	168b	Contig725	2290	2310	+	aggcuuggugcagcucgggaa
169	169a	Contig267	1556	1576	+	CAGCCAAGGAUGACUUGCCGA
169	169b	Contig13498	6461	6481	+	CAGCCAAGGAUGACUUGCCGG
169	169c	Contig675	3784	3804	+	CAGCCAAGGAUGACUUGCCGG
169	169d	Contig45157	681	701	+	UAGCCAAGGAUGAAUUGCCGG
169	169e	Contig2398	8572	8592	+	uagccaaggaugacuugccgg
169	169f	Contig1393	18877	18897	-	UAGCCAAGGAUGACUUGCCUa
169	169g	Contig3158	5985	6005	-	UAGCCAAGGAUGACUUgccua
169	169h	Contig12780	1801	1821	+	UAGCCAAGGAUGACUUgccug
169	169i	Contig20759	2212	2232	-	UAGCCAAGGAUGACUUgccug
169	169j	Contig30457	1664	1684	-	UAGCCAAGGAUGACUUgccug
169	169k	Contig38505	1415	1435	+	UAGCCAAGGAUGACUUgccug
169	169l	Contig4845	5060	5080	-	uagccaaggaugacuugccug
169	169m	Contig4845	7663	7683	-	UAGCCAAGGAUGACUUgccug
169	169n	Contig10799	6187	6207	+	UAGCCAAGAAUGACUUgccua
169	169o	Contig29602	1007	1027	+	UAGCCAAGAAUGACUUgccua
169	169p	Contig21032	3041	3062	+	UAGCCAAGGACAaacuugccgg
169	169q	Contig99778	87	107	-	UAGCCAAGGAGACUGcccaug
171	171a	Contig15726	6391	6411	+	UGAUUGAGCCGCGCCAAUAUC
171	171b	Contig14063	4647	4667	+	UGAUUGAGCCGUGCCAAUAUC
171	171c	Contig1580	11153	11173	-	UGAUUGAGCCGUGCCAAUAUC
171	171d	Contig2941	11346	11366	+	UGAUUGAGCCGUGCCAAUAUC
171	171e	Contig3749	5908	5928	+	UGAUUGAGCCGUGCCAAUAUC
171	171f	Contig3911	11181	11201	+	UGAUUGAGCCGUGCCAAUAUC
171	171g	Contig13529	3267	3287	+	gagGUGAGCCGAGCCAAUAUC
172	172a	Contig7420	8032	8052	+	AGAAUCUUGAUGAUGCUGCAU
172	172b	Contig11136	6459	6479	-	GGAAUCUUGAUGAUGCUGCAU
172	172c	Contig3578	6657	6677	+	UGAAUCUUGAUGAUGCUGCAC
393	393	Contig4493	10056	10076	+	UCCAAAGGGAUCGCAUUGAUC
394	394	Contig15318	315	335	+	UUGGCAUUCUGUCCACCUCC
395	395a	Contig16544	4585	4605	+	GUGAAGUGCUUGGGGGAACUC
395	395b	Contig16544	4745	4765	+	GUGAAGUGUUUGGGGGAACUC
395	395c	Contig16544	4886	4906	+	GUGAAGUGUUUGGAGGAACUC
395	395d	Contig16544	5024	5044	+	GUGAAGUGUUUGGGGGAACUC
395	395e	Contig16544	5167	5187	+	GUGAAGUGUUUGGGGGAACUC
395	395f	Contig16544	5308	5328	+	GUGAAGUAUUUGGGGGAACUC
395	395g	Contig16544	5446	5466	+	GUGAAGUGUUUGGGGGAACUC
395	395h	Contig32015	1661	1681	+	GUGAAGUGUUUGGGGGAACUC
395	395i	Contig32015	1823	1843	+	GUGAAGUGUUUGGGGGAACUC
395	395j	Contig32015	1996	2016	+	GUGAAGUGUUUGGGGGAACUC
395	395k	Contig32015	2156	2176	+	GUGAAGUGUUUGGGGGAACUC
395	395l	Contig32015	2331	2351	+	gugaaguguuuggggggaacuc
395	395m	Contig32015	2506	2526	+	GUGAAGUGUUUGGGGGAACUC
395	395n	Contig5127	4895	4915	+	gugaaGUGUUUGGGGGAACUC
395	395o	Contig15161	2207	2227	+	GUGAAGUGUUUGGGGAAACUC
395	395p	Contig13097	4446	4466	-	gugaagcguuuggggggaaauc
395	395q	Contig13097	5098	5118	-	GUGAAGUGUUUGGGGAAACUC
395	395r	Contig48729	1123	1143	+	GUGAAGUGUUUGGGGAAACUC
395	395s	Contig15161	2859	2879	+	gugaagcguuuggggggaaauc

396	396a	Contig1552	6429	6449	+	UUCCACAGCUUUCUUGAACUG
396	396b	Contig6441	8392	8412	+	UUCCACAGCUUUCUUGAACUG
396	396c	Contig1552	13825	13845	-	UUCCACAGCUUUCUUGAACUU
397	397a	Contig146	12104	12124	-	UCAUUGAGUGCAGCGUUGAUG
397	397b	Contig414	5761	5781	+	UUAUUGAGUGCAGCGUUGAUG
398	398a	Contig11355	3212	3232	+	UGUGUUCUCAGGUCACCCCUU
398	398b	Contig14993	2506	2526	-	UGUGUUCUCAGGUCGCCCCUG
399	399a	Contig11471	1413	1433	+	UGCCAAAGGAGAAUUGCCCUG
399	399b	Contig15629	4245	4265	-	UGCCAAAGGAGAAUUGCCCUG
399	399c	Contig9607	6567	6587	-	UGCCAAAGGAGAAUUGCCCUG
399	399d	Contig9306	3519	3539	+	ugccaaaggagaguugcccug
399	399e	Contig11471	282	302	-	UGCCAAAGGAGAUUUGCCCAG
399	399f	Contig28748	2409	2429	-	UGCCAAAGGAGAUUUGCCCAG
399	399g	Contig3634	914	934	+	UGCCAAAGGAGAUUUGCCCGG
399	399h	Contig9607	8629	8649	+	UGCCAAAGGAGACUUGCCCAG
399	399i	Contig2354	6504	6524	-	UGCCAAAGGAGAGCUGCCCUG
399	399j	Contig772	12030	12050	-	ugccaaaGGAGAGUUGCCCUA
399	399k	Contig31542	1939	1959	-	UGCCAAAGGAAAUUUGCCCCG

All *Oryza* miRNAs identified by computational prediction and homology to validated miRNAs are list
as are the portions of each locus computationally predicted to have miRNA encoding potential. Nucle
region are in lowercase. For miRNA families that have not been cloned, the 5' end of the sequence is
mobility on a Northern blot. For miRNA loci that are related to a cloned miRNA, the 5' and 3' ends are
miRNAs have heterogeneity at either the 5' or 3' end, the ends of the sequences listed should be cor
sequence length containing the miRNA, miRNA*, and intervening sequence.

sequence predicted to encode miRNA	hairpin arm	hairpin length	hairpin sequence
GGUGACAGAAGAGAGUGAGCACACGUGGUUG	5'	89	ACUAGGAGGGu
GUCUGACAGAAGAGAGUGAGCACACACGGUGC	5'	86	UUUGGAGGUCu
GGCUGACAGAAGAGAGUGAGCACACAUGGUGA	5'	91	GAGGUGAGGCu
UUGACAGAAGAGAGUGAGCACACAGCGUG	5'	84	CUCAUGAGAUu
GGUGACAGAAGAGAGUGAGCACACGG	5'	86	UGGCGCGAGG
AGUUGACAGAAGAGAGUGAGCACAC	5'	183	UGGUGGCAGUu
GGCUGACAGAAGAGAGUGAGCACACAGCGGG	5'	112	CGCGGCUGGC
UUGUUGACAGAAGAGAGUGAGCACACGG	5'	86	GCGAGAUUGUu
GGUGACAGAAGAGAGUGAGCACACGG	5'	86	CGCUGGGCGG
UUGUUGACAGAAGAGAGUGAGCACACGG	5'	86	GCGAGAUUGUu
UGACAGAAGAGAGAGAGCAC	5'	86	UUGAGAGUGAu
GCCGACAGAAGAGAGUGAGCAUAUAU	5'	136	GCUAGGGAGCc
UCUUUGGAUUGAAGGGAGCUCUG	3'	252	GUUGUGGACGu
ACUCUUUGGAUUGAAGGGAGCUCUG	3'	168	GGUUAUGAAGU
UGAUUGGAUUGAAGGGAGCUCCAC	3'	177	GAGGAGGAAGA
UGAUUGGAUUGAAGGGAGCUCC	3'	169	UGAUGUGAGGA
UUGAUUGGAUUGAAGGGAGCUCCU	3'	171	GAUGAAGAAGA
UUAUGCUUGGAUUGAAGGGAGCUCUA	3'	168	GAAGAAGAAGA
UGGUUGGACUGAAGGGUGCUC	3'	171	UGUGUAAGAAC
	3'	177	GAUGGAUGGAA
GUGUGCCUGGCUCCCUGUAUGCCACACA	5'	82	GUGUAGUGUGu
AGCGUGCCUGGCUCCCUGUAUGCCACUC	5'	82	CUUGAGAGCGu
AUGUGCCUGGCUCCCUGUAUGCCACUC	5'	87	AUUGGGAAUGu
GAUAUGCCUGGCUCCCUGUAUGCCACUCG	5'	130	AAAGGGGAUAu
GGAUAUGCCUGGCUCCCUGUAUGCCGC	5'	94	GUAGGGGAUAu
CUGCCUGGCUCCCUGAAUGCCAUC	5'	80	GGAUUAACGCu
GGAAUCGAUCGAUAAACCUCUGCAUCCAGU	3'	156	GUGGUGAUGCc
GAAUCGAUCGAUAAGCCUCUGCAUCCAGA	3'	114	UGGGUGAUGCc
ACGGUGGAGAAGCAGGGCACGUGCA	5'	93	CCGUGCACGGu
CCGCGUUGGAGAAGCAGGGCACGUGCAUGC	5'	122	AGGACCGCGUu
UUGUUGGAGAAGCAGGGUACGUGCAA	5'	99	AGGUUCUUGUu
CCGUGCUGGAGAAGCAGGGCACGUGCUC	5'	74	CAAACCGUGCu
AGGGUGGAGAAGCAGGGCACGUGAGC	5'	112	UUGUGCAGGGu
UCUCGGACCAGGCUUCAUUCCCCUCAGA	3'	111	UUGCUUCUGAC
UCUCGGACCAGGCUUCAUUCCCCCC	3'	170	UUCAUUUUGAC
UUCGGACCAGGCUUCAUUCCCCCC	3'	102	UGGCAGUUGAC
UCUCGGACCAGGCUUCAUUCCCCUCAAGU	3'	93	UUCACUUUGAC
UCUCGGACCAGGCUUCAUUCCCCUCAGA	3'	119	UUGUUUUUGGC
UCUCGGACCAGGCUUCAUUCCCCUCAACA	3'	77	AUCUUGUUGAC
UCUCGGACCAGGCUUCAUUCCUCACA	3'	125	AGCAUGGUGUC
UCGGACCAGGCUUCAUUCCUCGCAA	3'	99	GGUGGCUUGU
	3'	125	AGAUAGGUGUU
	3'	120	AGAUAGGUGUU
GGAGCCUCGGACCAGGCUUCAAUCCCUU	3'	107	AUUAGGUUAAC
CUCGGACCAGGCUUCAAUCCCUU	3'	97	GUUAGGUUAAC
GAGUGAAGCUGCCAGCAUGAUCUAGCUCUG	5'	115	UGUGAAUGAGu
CGUGAAGCUGCCAGCAUGAUCUAACUU	5'	127	AGAGAAAGCGu
GAGUGAAGCUGCCAGCAUGAUCUAGCUC	5'	127	AGGGAACGAGu
AGCUGAAGCUGCCAGCAUGAUCUGAUGA	5'	90	CAUUAGGAGCu
AUGAAGCUGCCAGCAUGAUCUGGU	5'	253	UGUGAGAGAAu
UGGAUGAAGCUGCCAGCAUGAUCUGAUCA	5'	93	CACAAGUGGAu

GGUGAAGCUGCCAGCAUGAUCU	5'	62	CAUAAGCAGGu
UUGGUGAAGCUGCCAGCAUGAUCUGAUGA	5'	100	CACAAGUUGGu
GGCUGAAGCUGCCAGCAUGAUCUGGU	5'	181	UGUGAGAGGCu
CUCGGGCUCGCUUGGUGCAGAUCGGGACCC	5'	67	CGCCUCGGGCu
	5'	86	UGGUCUUGUGa
AUGGUGCAGCCAAGGAUGACUUGCCGAUC	5'	95	GGCCAUGGUGc
AAUGCAGCCAAGGAUGACUUGCCGGU	5'	108	GAACGGAAUGc
GGAUGCAGCCAAGGAUGACUUGCCGGCUC	5'	92	GAACGGGAUGc
GUGUAGCCAAGGAUGAAUUGCCGGC	5'	83	AUUUAUCGUGu
	5'	112	GCUGGUUGUG
UUCGGUAGCCAAGGAUGACUUGCCU	5'	149	GCUGAUUCGGu
CUCUGGUAGCCAAGGAUGACUU	5'	106	CUGCCUCUGGu
CUCUGGUAGCCAAGGAUGACUU	5'	102	UUAGCUCUGGu
CUCUGGUAGCCAAGGAUGACUU	5'	88	GUAGCUCUGGu
AUCUGGUAGCCAAGGAUGACUU	5'	105	UCGCAUCUGGu
UAGAUAGCCAAGGAUGACUU	5'	136	UCUGUCUAGAu
	5'	88	UUAUCUCUGAu
CCUGGUAGCCAAGGAUGACUU	5'	118	UGAGUCCUGGu
UUUGGUAGCCAAGAAUGACUU	5'	157	CUCCCUUUGGu
UUUGGUAGCCAAGAAUGACUU	5'	146	CUCCCUUUGGu
AGCAAGGUGUAGCCAAGGACA	5'	106	GAGCAAGGUGu
UCAGGCUAGCCAAGGAGACUG	5'	88	CCACUCAGGCu
GUAUCUGAUUGAGCCGCGCCAAUAUCUC	3'	78	UGGAAAGAGCC
UCUUUUGAUUGAGCCGUGCCAAUAUCACGUC	3'	80	GCGACGACGGC
CUCUUUGAUUGAGCCGUGCCAAUAUCACGUC	3'	79	GUGGGAACGGC
UUCUGAUUGAGCCGUGCCAAUAUCUCAGC	3'	115	UUGUAGCUAUC
UUUCUGAUUGAGCCGUGCCAAUAUCUUAG	3'	99	UGGUAGCUAUC
GUCUGAUUGAGCCGUGCCAAUAUCAC	3'	107	GGGAGAGUGCC
GUGAGCCGAGCCAAUAUCAC	3'	69	GACAUGGCAUC
GGCUGAGAAUCUUGAUGAUGCUGCAUCCGCA	3'	89	GUGUUUGCGG
GGGAAUCUUGAUGAUGCUGCAUCGGAA	3'	218	GUGAUUUCUGA
UGCGUGAAUCUUGAUGAUGCUGCACCAGCAA	3'	91	CUUGUUGCGGC
GGGGAAGCAUCCAAAGGGAUCGCAUUGAUCCUUC	5'	95	UGGGGAAGCAu
GAGAGUUCUUUGGCAUUCUGUCCACCUCCUUG	5'	90	UACUGAGAGUu
AGUGAAGUGCUUGGGGGAACUCCAG	3'	122	UUGUCCACUGC
CGUGAAGUGUUUGGGGGAACUCUUA	3'	66	GAGUCCCUAGC
GUGAAGUGUUUGGAGGAACUCUCGG	3'	69	GUAUUAUCAAG
UGUGAAGUGUUUGGGGGAACUCUCGGU	3'	68	GUAUUGUCGUC
UGUGAAGUGUUUGGGGGAACUCUCGA	3'	68	GUAUUAUCGAC
GUGAAGUAUUUGGGGGAACUCUCGA	3'	68	GUAUUAUCGCC
UGUGAAGUGUUUGGGGGAACUCUCG	3'	66	GUAUCACCGUC
GUAGUGAAGUGUUUGGGGGAACUCUAGGUGGCA	3'	92	UUGUUACCUGC
UGUGAAGUGUUUGGGGGAACUCUUGGU	3'	79	GUUUUACCGGC
CUGUGAAGUGUUUGGGGGAACUCUAGGUGGCA	3'	92	GUGUUCCCAAC
UGUGAAGUGUUUGGGGGAACUCUUGA	3'	79	GUUUUAUCGGC
	3'	112	GUGGCCCCAGC
UUGUGAAGUGUUUGGGGGAACUCUUG	3'	114	AUGUCCCUAAC
GUGUUUGGGGGAACUCUCGA	3'	66	GUAUCACCGUC
AGUGAAGUGUUUGGGGAAACUCCGG	3'	123	UUAUCCACUGC
	3'	66	ACACUGCCAGC
AGUGAAGUGUUUGGGGAAACUCCGG	3'	123	UUAUCCACUGC
AGUGAAGUGUUUGGGGAAACUCCGG	3'	123	UUAUCCACUGC
	3'	66	ACACUGCCAGC

AUCUUCCACAGCUUUCUUGAACUGC	5'	134	CUUUGUGAUCu
GUCUUCCACAGCUUUCUUGAACUGC	5'	96	CUUUGUGGUCu
CAUGCCUUUCCACAGCUUUCUUGAACUUCU	5'	121	UGCCAUGCCUu
GCAUCAUUGAGUGCAGCGUUGAUGAA	5'	94	AUCAAAUGCAu
	5'	98	AGGGAAGGCAu
ACUGUGUUCUCAGGUCACCCCUUUGGG	3'	95	GCUGAACCCAC
CGUGUGUUCUCAGGUCGCCCCUGCCG	3'	68	GGAGUUCCUAC
GUGCCAAAGGAGAAUUGCCCUGC	3'	129	CUGUGAAUUAC
CGUGCCAAAGGAGAAUUGCCCUGCC	3'	77	GUGAGAAUCAC
CGUGCCAAAGGAGAAUUGCCCUGC	3'	90	CGGCGAAUUAC
	3'	266	AAGACAGUAGU
ACCACUGCCAAAGGAGAUUUGCCCAG	3'	98	ACAUGCAUUAC
UGUUCUCUCUGCCAAAGGAGAUUUGCCCAG	3'	97	UGGUGGAUUAC
UCUGCCAAAGGAGAUUUGCCCGGCGAU	3'	98	AUGUGCAUUGC
CCACUGCCAAAGGAGACUUGCCCAGCAA	3'	80	CCAUGCAUUAC
CCCUGCCAAAGGAGAGCUGCCCUGCCA	3'	96	GUGAGAAUCAC
GGAGAGUUGCCCUAAAACUGGA	3'	86	AGUCCAGUUUC
ACUGCCAAAGGAAAUUUGCCCCGGAAUUCA	3'	86	AGCUGCAUUGC

ed . The sequences of the mature miRNAs are shown,

eotides in the mature miRNAs outside of the computationally predicted

s determined by PCR of the miRNA and the length is inferred from

e inferred from the ends of the cloned homolog. Because many plant

nsidered to be approximations. Hairpin length is defined as the minimal

ugacagaagagagugagcacaCGUGGUUGUUUCCUUGCAUAAAUGAUGCCUAUGCUUGGAGCUACGCGUG
ugacagaagagagugagcacaCACGGUGCUUUCUUAGCAUGCAAGAGCCAUGCUGGGAGCUGUGCGUGCU
ugacagaagagagugagcacaCAUGGUGACUUUCUUGCAUGCUGAAUGGACUCAUGCUUGAAGCUAUGUG
ugacagaagagagugagcacaCAGCGUGAUGGCCGGCAUAAAAUCUAUCCCGUCCUCGCCGCGUGCUCACU
ugacagaagagagugagcacaCGGCCGGGCGUGACGGCACCGGCGGGCGUGCCGUCGCGGCCGCGUGCU
ugacagaagagagugagcacaCAGCGGCCAGACUGCAUCGAUCUAUCAAUCUUCCCUUCGACAGGAUAGCU
ugacagaagagagugagcacaCAGCGGGCAGACUGCAUCUGAAAUAAACUGGUGACGACGAAGAAGACGAC
ugacagaagagagugagcacaCGGCGCGGCGGCUAGCCAUCGGCGGGAUGCCUGCCCCCGCCGCGUGCUC
ugacagaagagagugagcacaCGGCCGGGCGGAACGGCACCGGCGGAUGUGCCGUCGCGGCCGCGUGCU
ugacagaagagagugagcacaCGGCGCGGCGGCUAGCCAUCGGCGGGAUGCCUGCCCCCGCCGCGUGCUC
ugacagaagagagagagcacaACCCGACAGCAGCGACGACGGCGGUCGCUUCUGCCAGGGCCGUGUGCUC
cgacagaagagagugagcauaUAUAGUUCUUUCCUUGCAUAUGUGGUCAUAUGUGUGUUGACUGAAGAGAU
UUGAGCUCCUUUCGGUCCAAAAAGGGGUGUUGCUGUGGGUCGAUUGAGCUGCUGGGUCAUGGAUC
UGGAGCUCCUUUCGUUCCAAUGAAAGGUUUAUCUGAAGGGUGAUACAGCUGCUUGUUCAUGGUUCC
AGGAGCUCCUUUCGAUCCAAUUCAGGAGAGGAAGUGGUAGGAUGCAGCUGCCGAUUCAUGGAUACC
AGGAGCUCCUUUCGAUCCAAUUCAGGAGAGGAAGUGGUGGGAUGCAGCUGCCGGUUCAUGGAUACC
AAGAGCUCCCUUUCGAUCCAAUUCAGGAGAGGAAGUGGUAGGAUGCAGCUGCCGGUUCAUGGAUAC
ACGAGCUCCCUUCGAUCCAAUCCAGGAGAGGAAGUGGUAGGAUGCAGCUGCCGGUUCAUGGAUACC
GAGAGCUCUCUUCAGUCCACUCUCAGAUGGCUGUAGGGUUUUAUUAGCUGCCGAAUCAUCCAUUCA
AGAGAGCGUCCUUCAGUCCACUCAUGGGCGGUGCUAGGGUCGAAUUAGCUGCCGACUCAUUCACCC
ugccuggcucccuguaugccaCACAUGUAGACCAACCCAUGGUGUCUGGUUGCCUACUGGGUGGCGUGCAA
ugccuggcucccuguaugccaCUCAUGUAGCCCAAUCCAUGGUGUGUUUGGAUGCUGUGGGUGGCGUGCAA
ugccuggcucccuguaugccaCUCAUCUAGAGCAACAAACUUCUGCGAGAGGUUGCCUAUGAUGGAUGGCGU
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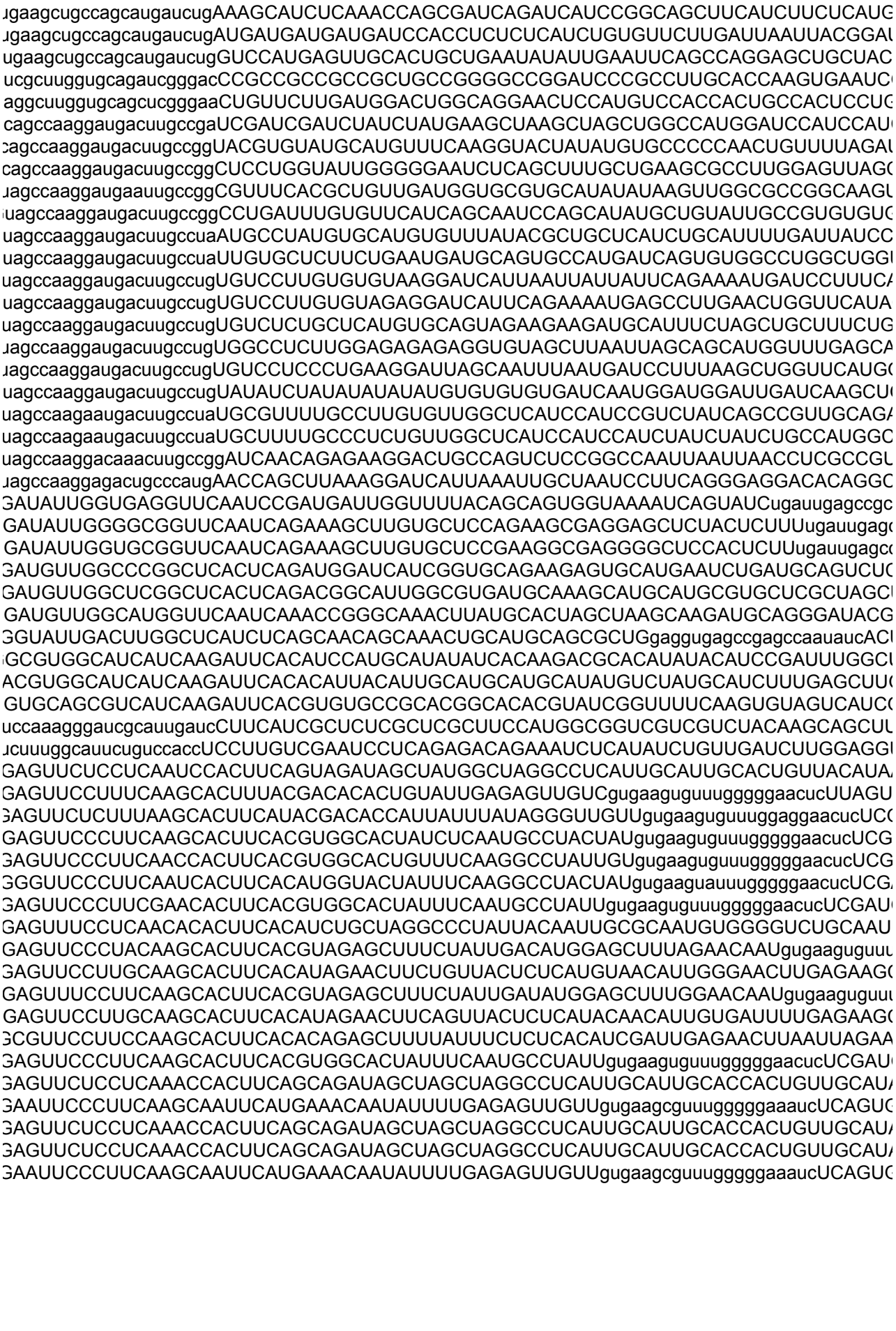

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