

**Compilation of small RNA sequences**

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This is the second collection of small RNA sequences. Small RNAs are broadly defined as the RNAs not directly involved in protein synthesis. These are grouped under three categories: 1) Capped small RNAs; 2) Noncapped small RNAs; and 3) Viral small RNAs. The sequences reported here are based on actual sequence determinations and not on the basis of oligonucleotide catalogs. The references are restricted to the citation of the latest publication in those cases, where several papers deal with one sequence; for some early references, see 1985 compilation of small RNA sequences (Reference 1). The abbreviations used for base and sugar modifications are as follows: m<sub>3</sub>G, N2,2,7, trimethylguanosine; A3, 2'-O-methyladenosine; C3, 2'-O-methylcytosine; G3, 2'-O-methylguanosine; U3, 2'-O-methyluridine; F, pseudouridine; A6, N6-methyladenosine; G2, 2-methylguanosine; N, unidentified nucleotide. The nucleotides that are identical to those in the sequence shown in the upper line are indicated by dashes and only the differences from the sequence on the top line are shown. Dots in the middle of a sequence indicate absence of any nucleotide. The references are given in the figure legends and the letters in brackets next to the references indicate the method used in deducing the sequence; C, complementary DNA sequence; D, sequence inferred from the DNA sequence of a putative gene and R, direct RNA sequence. In the case of capped small RNAs, the cap nucleotide is not included in the numbering system. Though earlier publications included cap nucleotide in counting RNA length, to facilitate comparison with genes and pseudogenes, the first transcribed nucleotide is numbered 1. Partial sequences where available are included in references but not in Figures. Minor variants of some small RNAs are not included. The readers are requested to refer to the original publications in all such instances. Any information regarding missing material or erroneous presentation is welcome. The literature survey was completed during January, 1986.

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Human U2	m3GpppA3U3GCUUCUUGCG3CUCUUUG3G	20	40	60	80	100
Rat U2	-PF-----PF-----F-----F-----F-----					
Mouse U2	------U-----U-----U-----U-----U-----					
Chick U2	------U-----U-----U-----U-----U-----					
Frog U2	------U-----U-----U-----U-----U-----					
F.Fly U2A	1 CUAG3AUCAA3 CUAAGAUCAAGGUAGUAAU	20	40	60	80	100
F.Fly U2B	------U-----U-----U-----U-----U-----					
F.Fly U2C	------U-----U-----U-----U-----U-----					
Amoeba U2	NPPdGAUAAAGAAACAGAA UUAUUAUGCGCACAAAGUU	1 20	40	60	80	100
Human U2	1 UUUNGGAGGGAGAAGGA AUAGGAGCTUGCUCUCCUA	20	40	60	80	100
Rat U2	-A-U-----U-----U-----U-----U-----					
Mouse U2	-AGUA-----U-----U-----U-----U-----					
Chick U2	-CCG-----U-----U-----U-----U-----					
Frog U2	-A-----A-----A-----A-----A-----					
F.Fly U2A	1 UCAGACGAGGCHAGAGC UUGCUCCACUCUGUGGCC	20	40	60	80	100
F.Fly U2B	------G-----G-----G-----G-----					
F.Fly U2C	-Q-----Q-----Q-----Q-----Q-----					
Amoeba U2	1 UUAAAAGGUGAGUGGCUUA GAUCCUCUUCUUCANGAUC	20	40	60	80	100
U2 RNA	Human:placenta,17(D),18(D),19(D);HeLa cells(partial),20(R). Rat:Novikoff hepatoma,21(R);brain(partial),20(R). Mouse:liver,22(D).					
Bird:chicken,liver,23(D); pheasant liver(partial),20(R). F.Fly:D.melanogaster,24(D). Frog:X.laete,26(D). Amoeba:D.dimidium,27(R).						
Plants:wheat(partial),28(R);pea(partial),15(R);broad bean(partial),29(R). Human U2 RNA has other post-transcriptional modifications not yet localized. The homology between U2 RNA of amoeba and U2 RNA of larger eukaryotes is not established.						





Rat	U6	AppGUGGUCCUCGGCAGCAGA UAUACUAUAAAURGGAAACAF ACAGGAGA3AGAUUA3G3CAUGC3 40	ACAGGAGA3AGAUUA3G3CAUGC3 CC3C3UCCGCAA3GG2AUGCAC3ACG CAAAUFGUGAAGGGUICCA 60
Mouse	U6	-----	-----
U6 RNA. Rat:Novikoff hepatoma, 46(R). Mouse: liver, 47(R), 48(D). Human:HeLa cells(partial), 49(R).	-----	-----	-----
Dinoflagellates:C.cohnii(partial), 50(R). X is an unidentified non-nucleotide.	X	-----	(106,107)
U7 RNA. Sea urchin, <i>P. milliaria</i> , 51(C).	-----	m3Gppp...UCUUUCAGUUUCUCUAGAA 20	GAGGGAGGUGGCCAA(U)-OH 80
Rat	U8	m3GpppA3U3CGUCAGGUUAUCCF 1 20	GEGUCUCGCCCCGUAGUUC 40
Mouse	U8	-----	A U C 60
RAT	U8	AUUCAGGACCGAACACACU 120	GGGGAGAUAAAUAUGUA 80
Mouse	U8	-----	UUGGAGCUUGAUAGUCIG 100
U8 RNA. Rat:Novikoff hepatoma, 52(R). Mouse:Friend leukemia, 53(R).	-----	-----	-----
Human 7SL-A	-----	1 20	-----
Human 7SL-B	-----	-----	-----
Rat 7SL	-----	Ac- 40	-----
Frog 7SL	-----	GCGGGCGUGGGCGUGUUG 20	GGGGAGGUAGGGAGGAAU 60
F.Fly 7SL-A	-----	GACUGGAAGGUUGGCAACU 130	CCUUCAGGAGGUAGGGAU 60
F.Fly 7SL-B	-----	PGGACUUGGCCGGCGUUG 130	AGGCUUCAGGAGGUAGGG 40
Bact. 7SL	-----	GCUCUCGCCGACACCCUGA 130	GUAGCUACAGCCGCGAC 150
Human 7SL-A	-----	CCCACAUAGUUCGCAUUA 130	CRRGGGACCCUCCGGAG 170
Human 7SL-B	-----	-----	UAGGAGGGUGAACCGGCC 190
Rat 7SL	-----	GGCACUAGUUCGCAUUAU 130	CAGGAGGGGAAACGGAGGG 210
Frog 7SL	-----	-----	UCCGUGGCAUUCAGGUUG 170
F.Fly 7SL-A	-----	CCGCACAUAGGUUCGCAUUA 130	UAGGAGGGGAUACGGGGC 190
F.Fly 7SL-B	-----	-----	CAGGGGGAAACCGAAC 210
Bact. 7SL	-----	AAGCCUCUCCGUCCGGAC 130	CAGGACCGAGUGGCAACCCC 170
			CGCAAGGGGUUCGGUGUG 170
			CAGGACCGAGUGGAGAC 190
			GGACCCUGGGCAUCCGGCC 190

Human	7SL-A	CGUGCUGAUCAGUAUGGGA	240	260	280		
Human	7SL-B	-----	-----	-----	-----		
Rat	7SL	GUCCCAUCAGUAUGGGAU	-----	-----	-----	(A)-OH	(299, 300)
Frog	7SL	-----	-----	-----	-----		
F.Fly	7SL-A	CGUGGUAUCAGUAUGGGA	240	260	280		
F.Fly	7SL-B	-----	-----	-----	-----		
Bacteria	7SL	ACCCGGGGCGGCGGGU	-----	-----	-----		
TS RNA. (7S RNA; L RNA; SRP RNA).	Human: HeLa cells, 54(C), 55(D). Rat: Novikoff hepatoma, 56(R).	CGGGACUACGCCAACGCU	CGGUUAUCACCCAAACGCU	GGGGACUACGCCAACGCU	GGGGACUACGCCAACGCU	CGGGGGCUACCCGACU	CGAC-OH (304)
58(C). Insect: <i>D. melanogaster</i> , 58(C), 59(C). Bacteria: <i>Halobacterium halobium</i> , 60(R).		-----	-----	-----	-----	-----	
Human	7SK	pppGAUUGCAGGGGAUCUGC	20	40	60	80	100
Rat	7SK	-----	-----	-----	-----	-----	110
Human	7SK	UUGGUCCCUCCGAGAUGC	-----	-----	-----	-G-	
Rat	7SK	-----	-----	-----	-----	-----	
Human	7SK	UUGGUCCCUCCGAGAUGC	130	150	170	190	210
Rat	7SK	-----	-----	-----	-----	-----	220
Human	7SK	CUCAAAAAGCUCUCAAGG	-----	-----	-----	-A-	
Rat	7SK	UUCUCCCCUCCUAGCCU	240	260	280	300	320
TSK RNA. (7-3 RNA; K RNA). Human: HeLa cells, 61(C). Rat: Novikoff hepatoma, 62(R).		-----	-----	-----	-----	-----	
Chicken CEH-RNA	CUAACAGGCCAACCAAACCA	ACCCUCGCCCUCCUAGGAC	ACCUACCCUGAAGGCC	CCCAACCCACUGAUCCAGC	CCUAUAUCUAUAUCGAAACA	100	
	CCACCUACUCAUAGGCCCA	UUAAGGCCUAGGUAGGCCU	AACAGGAACCCICACAGCUC	GUACACCACAUACUACAUU	AUCUCUAGGCCACAUUCGC	200	
	CCUUCUACUCAUAAACCU	CAAUUCCCCCCUACGCCA	CUCAUCCAUUCCCACUAC	CACUUCUAGGCCUACGCC	CACUUCUAGGCCUACGCC	300	
CEH-RNA. (Chicken Embryonic Heart RNA), 63(C).	UUCUCCCCUCCUAGCCU	CUACUCAAGAAAUAUUTU	AAAAAAUAUAUAUAUAUA	AAAUAUAUAUAUAUAUA	AAAUAUAUAUAUAUAUA	393	
Chicken tc-102	UCGGUGAGAGAGAGUGUU	UCGGUGAGAGAGAGUGUU	UCGGUGAGAGAGUGUU	GAGAGUGUUAUAUAUAGU	UAGGGUGUGGGGGGGGGGC	G	
Chicken tc-89	AAGAGAGAGAGAGAGUUG	UAGAGAGAGAGAGAGUUG	UAGAGAGAGAGAGAGUUG	UAGGGUGAGAGAGAGUGAA	UAGGGUGAGAGAGAGUGAA	AUCAGAGACA	
Chicken tc-90	UUGAAUCGUAGAGGGUGA	UAGGGGGGGGGGGGGGGGG	UAGGGGGGGGGGGGGGGGG	AUCAGAGAGAGAGAGUGAA	UAGGGUGAGAGAGAGUGAA	CUCGCCAGCA	
tc-RNAs. (translational-control RNAs). Chicken embryonic skeletal muscle, 64, 65(R).							

Hamster 6S	pppGGGUGGAGAGAUGCUCAGA GGGCGUGUGAGUGUCUCAG	20	40	60	80	100
Mouse 6S	GUUAACAGAGCACUGACUGCU UGGUAAAAGAACCCACUG	130	150	170		
Hamster 6S	UGCCCUUUCUUUCUGUGAG ACUCCCUUCUUCUGUGUC					(183)
House 6S	AUAUAUCAUAGGAAGCAGAA UAUACAUAGGAAGCAGAA					
6S RNA. (Type 2 all RNA; B2 RNA). Hamster:Chinese hamster ovary, 66(D). Mouse:Murine fibroblasts, 67(C). The sequence shown for mouse 6S RNA is a consensus cDNA sequence; for sequences of individual cDNA clones see Reference 67.						
Hamster 4.5S	1 Mouse 4.5S	20	40	60	80	100
4.5S RNA. Mouse:Friend leukemia, 68(R). Hamster:Kidney, 68(R); ovary, 69(R).						
4.5S1 RNA. Rat:Novikoff hepatoma, 70(R).		1 20	40	60	80	100
Human Y1	pppGGGUGGUGGUCAAGGG GUUAUCUCAAUUGAUUU	20	40	60	80	100
Human Y3	GUUAUCUCAAUUGAUUU UGGUUCAACUAAUGAUCA					
Human Y5	pppAGUUGUCCGAGUGUG GUUAUCUCAAUUGAUUU					
X RNAs. (Ro RNAs). Human:Y1,Y3 RNAs:placental DNA, 71(D); Y5 RNA, HeLa cells, 72(R).						
E.coli 4273	1 GAACCGUACGAGACUGC	20	40	60	80	100
E.coli ts709						
E.coli 4273	GCCGGGGGAAACCCAG	120	140	160	180	200
E.coli ts709						
E.coli 4273	GCCCCGGGUAAACGUCCG	220	240	260	280	300
E.coli ts709						
E.coli 4273	UGACCCGGGUAGACUGUC	320	340	360		
E.coli ts709						
Rhase P RNA E.coli 4273,73(D); E.coli ts709,74(D).						
						(375)
						(375)



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