Supplementary materials

FIGURE LEGENDS

Table S1 Sequences and accession numbers of computational identified snoRNA genes in *C. elegans*.
Structure elements of snoRNAs are boxed. The sequences complementary to the target RNAs are indicated by shadows. The sequences in red indicate the genes were also identified from Deng´s cDNA library [20]. The gene names used in Deng´s study are also shown behind the sequences.

Table S2. Oligonucleotides used in this study

Figure S1. Functional prediction of the novel box C/D snoRNAs. Methylation guide duplex between snoRNAs and rRNAs/snRNAs are predicted. The snoRNA sequences in a 5' to 3' orientation are shown in upper strands, while rRNA/snRNA sequences in a 3' to 5' orientation are shown in lower strands. Sequence motifs of box D′ /D are boxed. The positions of predicted methylation sites are indicated by numbers. Only one isoform is shown if the snoRNA has more isoforms. (A) Methylation sites predicted by novel snoRNAs. (B) Novel methylation sites predicted in this study for the snoRNAs identified previously [18].

Figure S2. Functional prediction of the novel box H/ACA snoRNAs. Pseudouridylation guide duplex between snoRNAs and rRNAs/snRNAs are predicted. The snoRNA sequences in a 5' to 3' orientation are shown in upper strands, while rRNA/snRNA sequences in a 3' to 5' orientation are shown in lower strands. The two sequence motifs are boxed and the upper parts of the hairpins are represented by solid lines. The positions of pseudouridines are indicated by numbers. Only one isoform is shown if the snoRNA has more isoforms. (A) Pseudouridylation sites predicted by novel snoRNAs. (B) Novel pseudouridylation sites predicted in this study for the snoRNAs identified previously [18].

Figure S3. Structural and functional evolution of H/ACA snoRNA. Schematic representation of the homologous sno/scaRNAs in yeast and metazoa are shown. The box elements are present and the homologous guide hairpins are boxed. The names of snoRNAs and the guiding functions of the hairpins are also shown. *S.c.*, *C.e.*, *D.m.* and *H.s.* represent *S. cerevisiae*, *C. elegans*, *D. melanogaster* and *H. sapiens*, respectively.

Table S1 Sequences and accession numbers of computational identified snoRNA genes in *C. elegans*

Novel SnoRNA	Accession No.	Sequence
Box C/D snoRNA		
MeCeSSU-A90	AM050148	caagcg**atgacga**ttgatatctgctctaatgagt**ctga**attaccatgttgagatcttgt**ctga**gctc (CeN65)
MeCeSSU-C400	AM050149	aggctg**gcgatga**ttgagattgttcccacacgcaatttctcctgatccacatgaaggctaaacttccttggacgt**ctga**gccg (CeN33)
MeCeSSU-A422	AM050287	aaaagt**gcgatga**atgacttggataagtttcgg**ctga**aacttggtgatgccaacttttttaaaactg**ctga**gcac (CeN108)
MeCeSSU-A601	AM050288	aagcaa**atgacga**aatcgcacctcggcccgactccaaccctgggggcgaaatgagctttttaact**caga**tgct (CeN24)
MeCeSSU-C980	AM050289	catcca**atgatga**gagtttgcgactagggcggt**ctta**cacaatcatggtgattctagtcatt**ctga**tggt (CeN27)
MeCeSSU-C1180	AM050290	acgatg**atgatga**agaatttttgatatggtgtcaggacct**ctga**gagttccgtgatgatgtttagagttc**ctga**atct (CeN106)
MeCeSSU-C1196	AM050291	cgggca**atgatgt**catactcaaatctgtcagtcct**caga**ccatccctggtgataacacggttttccggagtg**ctga**gcag
MeCeSSU-U1342	AM050838	ggattg**aagatga**aatcaacgacgctgaccggtagtgatgtgttgtcgaaacctctaagaagta**ctga**ccgc
MeCeLSU-G668	AM050839	aagtca**atgatgt**tttttcaagacggga**ccga**ctggtgaatgatgcataaatgaaatg**ctga**gact (CeN13)
MeCeLSU-A678	AM050840	ctggca**gtgatga**tcacaaatccgtgttt**ctga**caagcgattgacgatagaaaaccgg**ctga**gcca (CeN5)
MeCeLSU-G860	AM050841	atgtgc**gcgatga**actctttaccatctttcgggg**cata**acactcttgatgataacatacccatttg**ctga**gcgc (CeN70)
MeCeLSU-A1185	AM050842	gaatcg**gtgatgt**gatatccagttctgcta**ctga**gttattgtgaagattaactttccccgt**ctga**gatt (CeN118)
MeCeLSU-C1502	AM050843	tgtccg**ttgatga**caacatacatacaccattacgatct**ctga**agacttcgtgctgatcatgtatccatgcaacaccaa**ctga**ggac (CeN15)
MeCeLSU-U1566	AM050844	gggtcg**atgatta**gattattctactgttcacgta**ccga**ctgttcagtggtggataaaccttct**atga**gact
MeCeLSU-C2300	AM050845	gggttt**gtgatga**ctgcatacggatcactgggct**ctga**atctctatgaaccgataatatccgtt**ctga**tacc (CeN54)
MeCeLSU-A2317	AM050846	atgtca**atgatgt**ctaaaaaattactacgatttaat**tcga**attgctgtgagatcaatcttatacaatt**ctga**gaca (CeN119)
MeCeLSU-G2343	AM050847	acggct**gtgatga**tttcctattgccgtttacccgt**ctga**ggaaaaccgtgcttgatacaacttggaaaagg**ctga**gccg (CeN121)
MeCeLSU-A2384a	AM050848	cactaa**atgacga**atcctaataacccaatgggtttcattgcggatatgaggcatttgt**ctga**gcgg (CeN23-1)
MeCeLSU-A2384b	AM050849	cactaa**atgacga**atcctaatagcccaatgggtttcattgcggatatgaggcatttgt**ctga**gcgg (CeN23-2)
MeCeLSU-A2384c	AM050850	tcacca**atgatgc**aatggttaaatcagacgagtctattttggctatctattcgagttcttcgaagaaattgccgctaagcggggtgaaattgaggcatttgt**ctga**ggtg (CeN17)
MeCeLSU-C2407	AM051009	ccgcct**gtgatgc**ttttcatgcgtgtcag**ctga**gatcccctgattagaaccacccttgaggttgtattaa**ctga**ggct
MeCeLSU-U2417	AM051010	gaggag**ttgatga**gaactctaatccattct**ctga**gcgagaaggatggccgaagcgggttcgcatttgaggcattaaggtagacgacagagttcttctggaaactactgcctcgcg**ctga**cgtc (CeN117)

MeCeLSU-A2429	AM051011	ctgggaatgatgaccttctgtgtaggaatctcaatgagtgactgtgacataaaaatgcagtaaattcactgacccc (CeN122)
MeCeLSU-C2440	AM051012	ctgtgtatgacgacaacgtgttagggacatctgcaccaaccgtgaagatttaacgaaagtagtactgacaca (CeN50-1)
MeCeLSU-U2762	AM051013	agagttgtgatgacttacttaggacaccttttggagggcctcggtctgctgaaaagaaaaatagtacgccttagtctgagctc
MeCeLSU-U2841	AM051014	actgcggtgacgatcaactcttacctactatgacaaaaacaatggttagacgttactcgtactgtctgagcag (CeN14)
MeCeLSU-G2903	AM051015	cgagcggtgatgaatgcacgtattgctctgacacctcttatgttagcggtaaatttccgtgccgcgatgagtccactaggatctctgagctc (CeN124)
MeCeLSU-A3023	AM051016	ttggctgtgacgattactattcccaacgcttggaatgaccaaagtgattattaaccaatccttttctgagcca (CeN30)
MeCeLSU-C3060	AM051017	gcttctgtgatcgacaacaaacccagctctctgaatgattgtgaggttaagaaaacagctgagaag
MeCeLSU-C3071	AM051018	agatctatgatgagactttcacgacggtcttccgatgtaatacatacctgtggatatctttacgtgaagctgagatc (CeN123)
MeCeLSU-A3159	AM051019	aagccgatgattaccaaaaaataaccaaatgtttgagtgattgtttgtgatcgaattttgtcactatcgctgaggct (CeN120)
MeCeU2-G12	AM051020	gacaggatgatgagtcactcgctgagtgacaataagccgagtgttagcggtttttatgtaatcgtgatgatcattccctcaaaagcgataattgtgaactgactgt (CeN69)
MeCeU2-C42	AM051021	tgcgcgatgaggattgataacacatacacacactctgaagttatgtgaagagataattgaagaacggatatctgagcgc (CeN89)
MeCeU2-U49	AM051022	cccaaaatgacgaaaccctatccccaatgattttacgacgtttcaagtcatttccatatcaatcggcgacctactacaccgtcttttgattgacctgaaacgcgtagagctagtgaatcttcatcataacgataagatctgattgg
MeCeU6-A48	AM051024	cctgcgatgatgagaatcatgctaatcatgaagagactcgtcagtctccaacctccatgttgtaaaacaatcgatagctgagcag
MeCeU6-C55	AM051023	agtgctgtgaagagaacgtgccactgtactttgcccatcggaagggcattgaaatggagatatacctggcacaggggccatctgagcac (CeN98)

Box H/ACA snoRNA

ΨCeSSU-499	AM051025	tgtgtcatttattcctcggatagtgatatttgttccttttcgagctcatactgacacaaaagcaatggcagatttgcaactagttgcgttctacttgatcgtggctatacttcttctgaccacattt
ΨCeSSU-513a	AM051026	gcagatgtccattacgcgaatgcctgtgcttttcgacgtttagtttagtctgcaagatagaattgatgtctcgaagacaggtacatcagcttttagagttcctgttcaagatcagcaaacaaac (CeN110)
ΨCeSSU-513b	AM051027	cactctacgcctttcctttcaatgggttggtatgatttaataaggatgcaagagaatagtacgggaagttcgaagacttgcggatttgcttccacggctccgcgagttcataactgtcaccacaatt (CeN41)
ΨCeSSU-970	AM051028	cgcttcatggtgcattacactttgtaacttgcattgtgtgcgcgactaatgaagcaatacaatatctgcagtatctgctcattcgccgtgttgtagttgatgagctcgtagtccgataacaatt (CeN64)
ΨCeSSU-971	AM051029	gcacacagaaagtcgagattgccggcttcaaaggctttatcgttgattacgtgtgctataacaacgccataggtatcattcttcgtcttcttaggcgcggaacatctaaatatggctacattt (CeN48)
ΨCeSSU-1146	AM051030	cacccacggcaacaaattccaattgtgtgtaacattcaatgcaattgagccgacgccgtgggaaatcactttcgtggaaccatttgatcccacgctcgttactgttaatgattgtggtttgcacgtggtttacaatt (CeN125)
ΨCeSSU-1152	AM051031	cacggccagtttgagttgattcgctctttcgcaatagagctttgagtcaaattactgtccggaaattatagagatgaagctcatttggagcaataacaattgttgaagctccgaaaatgatgctcttcccacaatt (CeN59)
ΨCeSSU-1266a	AM051032	tcctcgcttataagccgggggactagcctttgctgagttcactagtaaaacgcgaggaaaattatagagacatccagacaccgattatgttgtgatatattcggtcacttcgtatgtcccacagtt
ΨCeSSU-1266b	AM051033	acctctcttataagccgggggactagcattttgttaagttcactagtaaaatgagaggtaaagcatagagacaaccagacaccgagaatgttttgatgttttcggtcactttgtgtgtcccacaaat (CeN101)
ΨCeSSU-1523a	AM051034	gcatcgaaaaatggacggactttcccgatggatcgtttcgtataatttggtgcaaaatagtagagagacgcagtgttactcttccttacgttacctgtatctggagagtgtcaagcgtttcccacattt (CeN80)

ΨCeSSU-1523b	AM051035	tgcattgaaaagggacgggctatctttatggattgtttcgtataatttgatgcg**agataa**tagagaggcgcactgttactcttcatcacttttctgtgaacggagagtgtcaagcgcttccc**aca**att (CeN79)
ΨCeLSU-1058	AM051036	ggcatgtatttaatgcgggtggatgattttagttcagccgtttagcttgcatgca**aaatta**ttttctattgaagattttattcgctgcactcggtcatattcgagtctatgatttttctgaaattggt**aca**ttt
ΨCeLSU-1176	AM051037	ccgcgcctagaaactcagcggtgttttgtctttatgatcatgagctggtagcaggtgca**atattg**attcggagtcttaccggcgtccaccttctgaagggtgtgcgttaaaagttcatccga**aca**att (CeN55)
ΨCeLSU-1573	AM051038	gcgtctctccaacccggacaactcgcaacttgttgcaatggtcagttcgactgttagagacgct**aaataa**gaaaattggcggattcgctaacgtcgcgtcccatgaccggagtttgaaagtaggccgctga**aca**att (CeN58)
ΨCeLSU-1996	AM051039	gcccggtctttgccccggatcaactttaatagttatgacggtttccgactcggg**aaaata**gattcttgcataacgacggaaatttcagagtttgttctgaatgtctccagtcgatgtgcaat**aca**att (CeN43)
ΨCeLSU-2294a	AM051040	tgcctcagtgtctgggcaaagcagactgtttttaatagtactgacttaatccactgatggct**atagaa**aaacattgtgtaatccccgaacaccttgtggtgtattggattcatgacattgtg**aca**att (CeN84)
ΨCeLSU-2294b	AM051041	tgccacaatgtctgggcaaggtgccctgcacccgcttgagcaacctaatccattgatggct**atataa**aattcattgagtaatccctgttgagttatatagaactttccagtttcatatcattgat**aca**att
ΨCeLSU-2361	AM051042	cgaagagtagtcatttgtatacgtgatacacttatacagagttacacttttcgt**atataa**gtgcttcttctcgttggagttgtttatttaatgagcaattacctctaaactgaaagca**aca**att (CeN81)
ΨCeLSU-2483	AM051043	tttccattaagtagtcccgtacgttgaaattttcacgtacggtttggacaatatggag**aaatga**atgttactctttttgcaatgacgctgaaacatattttcgctttgttagccataagtttaaa**aca**att (CeN93)
ΨCeLSU-2519a	AM051044	cgaggtccagagtcacctctgagaatatgtatctcttcggtgctaggattgctcga**aaatga**attaggtactttatgcaaacagttcttgcgttatgctcatctgtttcacctccgtttctg**aca**aaa
ΨCeLSU-2519b	AM051045	acgaggtccagagtcacctttcagcaaaatgcagttagggtgctaggatttctcga**aaatga**aattggtcgtattatgcaagtagggcttgcttttgcgctccccgacttcacctcattcccaa**aca**ttt (CeN100)
ΨCeLSU-2836	AM051046	gcgtgagtactggctgtttgtgtctccgccattgccaatcagctgtcattaaccacgca**ataccg**tacagaggtgctaaattaccgtgaaaaccactttaaaaattggtcactcggaggaggcacctca**aca**ttt (CeN51)
ΨCeLSU-2966	AM051047	tgctcttcaaaagcactggttttaggatccactattatccaagccagccgtcaaaactgagct**ataaga**attatcttgttttgggtgaggtgtattcaattcagaatgcgtctcaataacacgatg**aca**att (CeN87)
ΨCeLSU-2992	AM051048	ggcggataggaaccaggtcatgtttgtacgtgatttgggccgacatcccgccg**aaatag**tcctgcgaagatctaaggtcgcttctggatggatttgccaccgggtgcacaccatcttcgtgc**aca**att (CeN39)
ΨCeLSU-3220	AM051049	aagcagtcgttcagtggcgcaagcgattttcccatgcgcttgtgcctaatctttgactgcg**agataa**aaatgtcagagtcaagcggtccgctatctgtgtagcattctgttcaag**ata**ttt (CeN126)
ΨCeU2-16	AM051050	gcggatattataaacagattcctataaatttacaaggtagtcagtagccgaaattcgca**atataa**attttccattaaaggcgagaaatcatccttttgtcggtgatatctagaaataaaactggag**aca**att
ΨCeU2-41	AM051051	ggtataagaaagcgagcaattattcgatggctgctcgttgataattcctcgaaaactaatgg**aaatta**agcgattatccgcctcatgaacagatccatcatttaacttgaatgataaggtctacttgacaggcgt**aca**tta
ΨCeU4-62	AM051052	cgcaccttgttttcacgcattctcggtcagttttctcgagattcaacgcgtccagaaagtgtgctat**atagaa**ataaagagtactttataagggccatacacggtttaatttaccccggctatggtagtttaacggtgct**aca**aat
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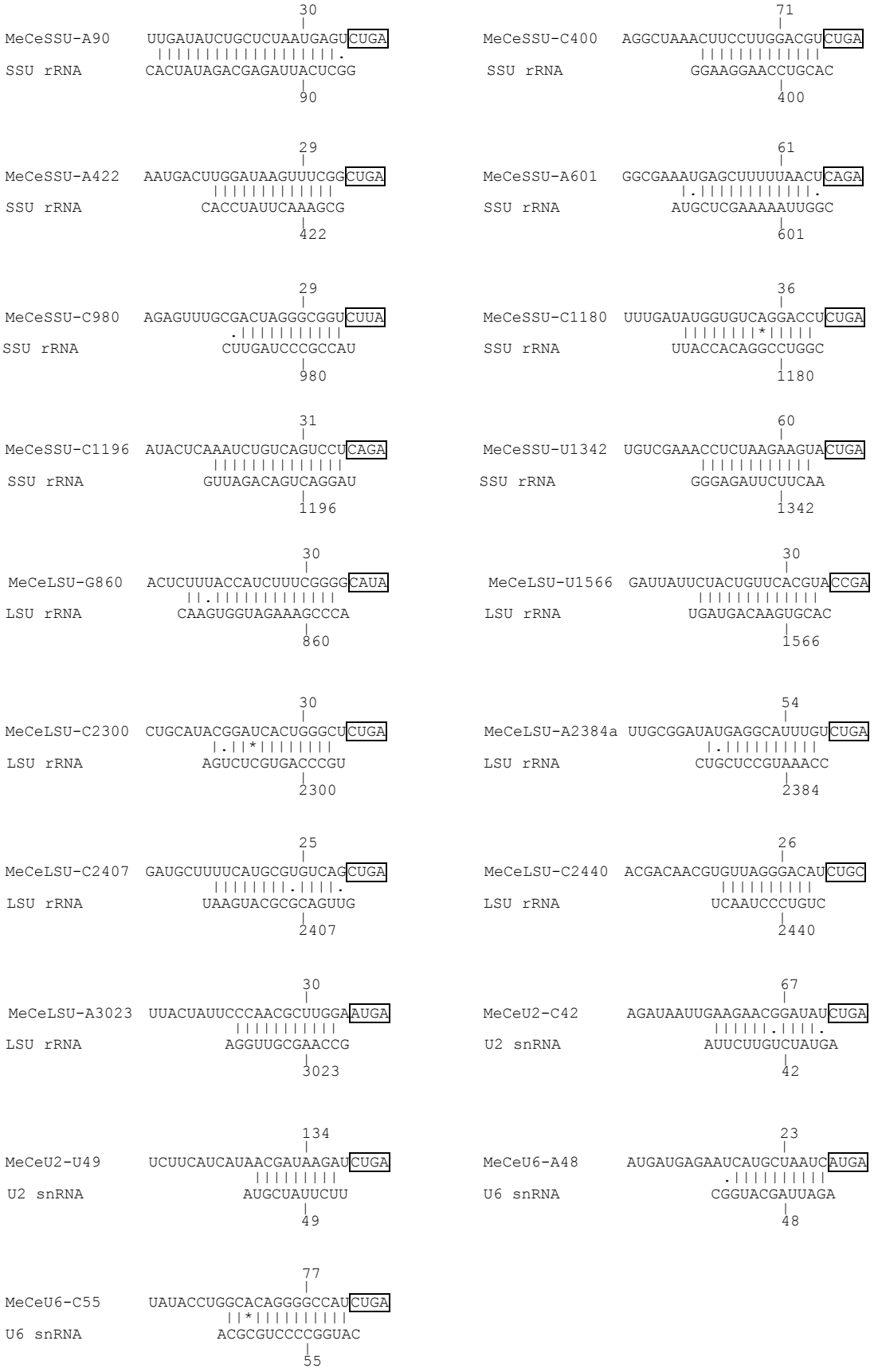
CeACAOrph2	AM051058	gggcacagagttgcagttgattgaaactaaatcatgtgcgctaagttgtcccg**aaagta**tctttggatttctacaaaacagtgttccttctttaggatgctgatccgtagagatctga**aca**aat (CeN45)

Table S2. Oligonucleotides used in this study

Oligonucleotide name	Sequence
Oligonucleotides used for northern blotting	
PMeCeSSU-C1180	5'-CTCTAAACATCATCACGGAACTCT-3'
PMeCeSSU-U1342	5'-GAGGTTTCGACAACACATCACTAC-3'
PMeCeLSU-C2300	5'-CATAGAGATTCAGAGCCCAGTG-3'
PMeCeLSU-A2317	5'-TTGTATAAGATTGATCTCACAGCA-3'
PMeCeLSU-A2384c	5'-GCTTAGCGGCAATTTCTTCG-3'
PMeCeLSU-C2440	5'-TCGTTAAATCTTCACGGTTGGT-3'
PMeCeLSU-U2841	5'-AGTACGAGTAACGTCTAACCATTG-3'
PMeCeLSU-C3071	5'-TATTACATCGGAAGACCGTCGT-3'
PMeCeU2-C42	5'-TCCGTTCTTCAATTATCTCTTCAC-3'
PMeCeU6-C55	5'-TGCCAGGTATATCTCCATTTCA-3'
PMeCeSSU-C1196	5'-CTGAGGACTGACAGATTTGAGTAT-3'
PMeCeLSU-C2407	5'-TCAAGGGTGGTTCTAATCAGG-3'
PMeCeLSU-C3060	5'-TAACCTCACAATCATTCAGAGAG-3'
PMeCeU2-U49	5'-CGTTATGATGAAGATTCACTAGC-3'
PΨCeSSU-499	5'-GTCAGAAGAAAGTATAGCCACGAT-3'
PΨCeSSU-970	5'-AACACGGCGAATGAGCAGATA-3'
PΨCeSSU-1146	5'-GCAAACCACAATCATTAACAGTA-3'
PΨCeLSU-1573	5'-TTAGCGAATCCGCCAATTTTCTT-3'
PΨCeLSU-1996	5'-TTCCGTCGTTATGCAAGAATCTA-3'
PΨCeLSU-2294a	5'-GCCATCAGTGGATTAAGTCAGTAC-3'
PΨCeLSU-2361	5'-AAACAACTCCAACGAGAAGAAGC-3'
PΨCeLSU-2992	5'-AAATCCATCCAGAAGCGACCT-3'
PΨCeU5-48	5'-CTATTTCAGAAACTTCCAGCATTA-3'
PCeACAOrph1a	5'-GAGATGCAAGACAAACACCGAA-3'
PΨCeSSU-1266a	5'-CCGAATATATCACAACATAATCG-3'
PΨCeLSU-2294b	5'-CTGGAAAGTTCTATATAACTCAACA-3'
PΨCeLSU-2519a	5'-TCATTTTCGAGCAATCCTAGC-3'
PΨCeU2-16	5'-ATTTCGGCTACTGACTACCTTGT-3'
PΨCeU2-41	5'-CGCCTGTCAAGTAGACCTTATC-3'
PΨCeU4-62	5'-AGCACCGTTAAACTACCATAGC-3'
PΨCeU6-26	5'-GATGAGCACCGCTAATGACA-3'
Oligonucleotides used for cDNA libraries construction and screening	
*Hin*dIII(T)$_{16}$	5'-CCCCAAAAGCTTTTTTTTTTTTTTTTTTTTTTT-3'
*Bam*HI(C)$_{16}$	5'-GGAATTCGGATCCCCCCCCCCCCCCCC-3'
P47	5'-CGCCAGGGTTTTCCCAGTCACGAC-3'
P48	5'-AGCGGATAACAATTTCACACAGGA-3'

Figure S1

A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LSU rRNA
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2440
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LSU rRNA
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30
3023
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U2 snRNA
AUUCUUGUCUAUGA
67
42
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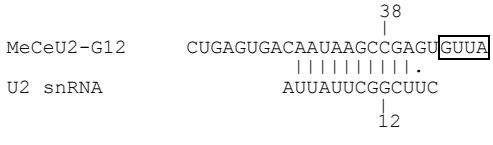
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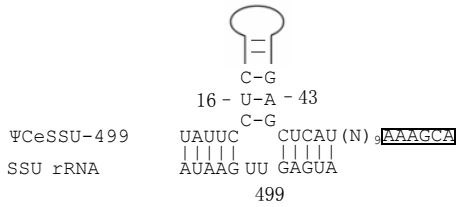
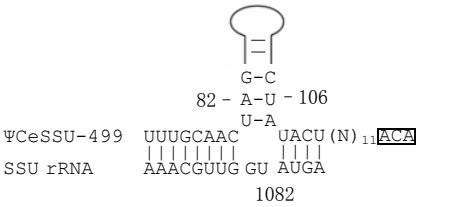
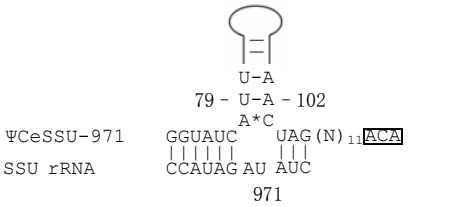
U2 snRNA
AUUAUUCGGCUUC
38
12

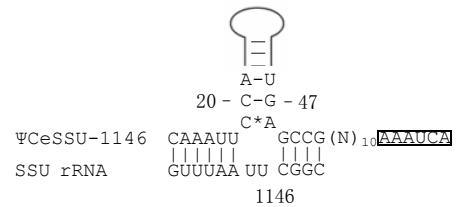
Figure S2

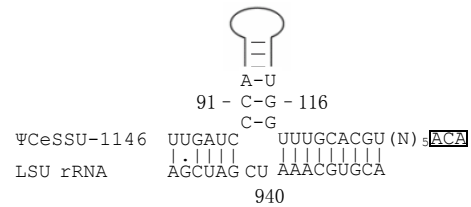
A

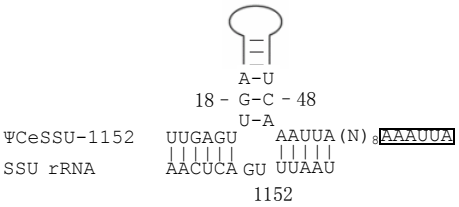

C-G
16 - U-A - 43
C-G
ΨCeSSU-499 UAUUC CUCAU(N)9AAAGCA
SSU rRNA AUAAG UU GAGUA
499

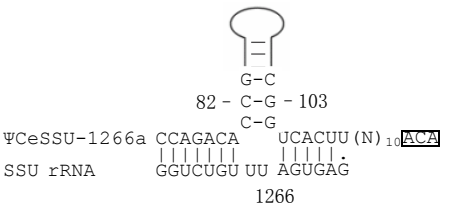

G-C
82 - A-U - 106
U-A
ΨCeSSU-499 UUUGCAAC UACU(N)11ACA
SSU rRNA AAACGUUG GU AUGA
1082

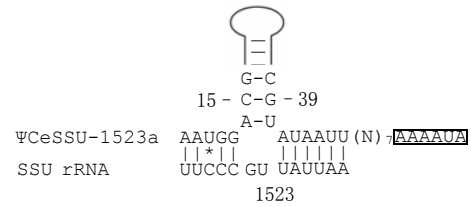

U-A
79 - U-A - 102
A*C
ΨCeSSU-971 GGUAUC UAG(N)11ACA
SSU rRNA CCAUAG AU AUC
971

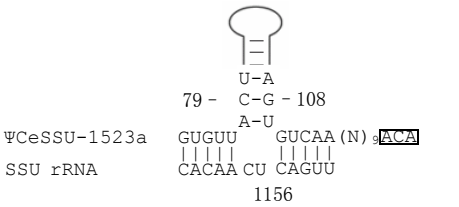

A-U
20 - C-G - 47
C*A
ΨCeSSU-1146 CAAAUU GCCG(N)10AAAUCA
SSU rRNA GUUUAA UU CGGC
1146

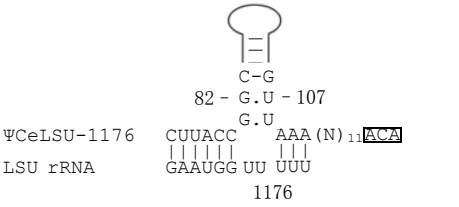

A-U
91 - C-G - 116
C-G
ΨCeSSU-1146 UUGAUC UUUGCACGU(N)5ACA
LSU rRNA AGCUAG CU AAACGUGCA
940

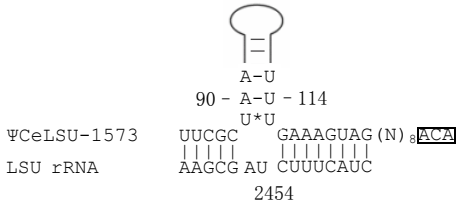

A-U
18 - G-C - 48
U-A
ΨCeSSU-1152 UUGAGU AAUUA(N)8AAAUUA
SSU rRNA AACUCA GU UUAAU
1152

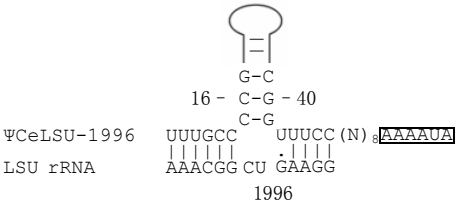

G-C
82 - C-G - 103
C-G
ΨCeSSU-1266a CCAGACA UCACUU(N)10ACA
SSU rRNA GGUCUGU UU AGUGAG
1266

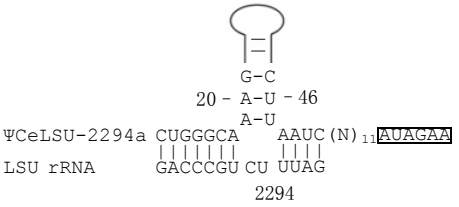

G-C
15 - C-G - 39
A-U
ΨCeSSU-1523a AAUGG AUAAUU(N)7AAAAUA
SSU rRNA UUCCC GU UAUUAA
1523

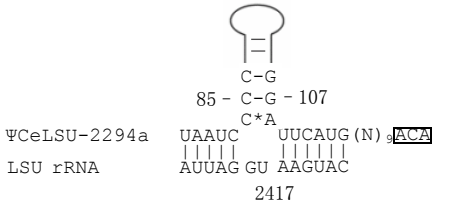

U-A
79 - C-G - 108
A-U
ΨCeSSU-1523a GUGUU GUCAA(N)9ACA
SSU rRNA CACAA CU CAGUU
1156

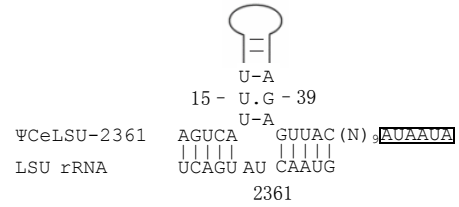

C-G
82 - G.U - 107
G.U
ΨCeLSU-1176 CUUACC AAA(N)11ACA
LSU rRNA GAAUGG UU UUU
1176

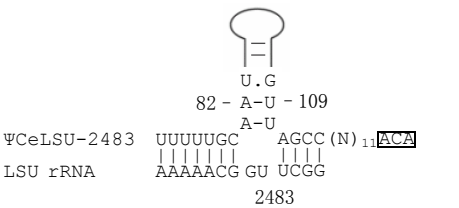

A-U
90 - A-U - 114
U*U
ΨCeLSU-1573 UUCGC GAAAGUAG(N)8ACA
LSU rRNA AAGCG AU CUUUCAUC
2454

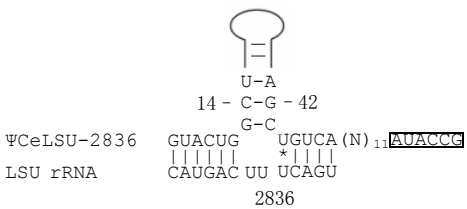

G-C
16 - C-G - 40
C-G
ΨCeLSU-1996 UUUGCC UUUCC(N)8AAAAUA
LSU rRNA AAACGG CU GAAGG
1996

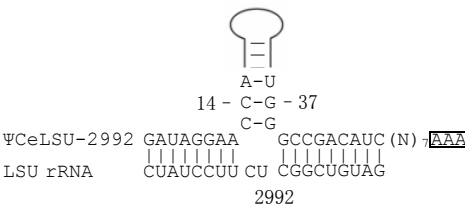

G-C
20 - A-U - 46
A-U
ΨCeLSU-2294a CUGGGCA AAUC(N)11AUAGAA
LSU rRNA GACCCGU CU UUAG
2294

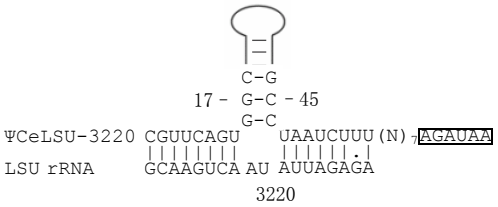

C-G
85 - C-G - 107
C*A
ΨCeLSU-2294a UAAUC UUCAUG(N)9ACA
LSU rRNA AUUAG GU AAGUAC
2417

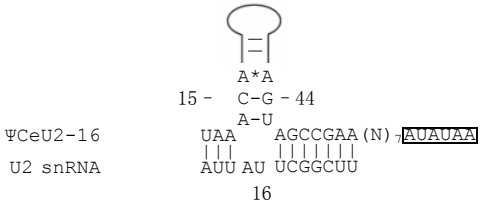

U-A
15 - U.G - 39
U-A
ΨCeLSU-2361 AGUCA GUUAC(N)9AUAAUA
LSU rRNA UCAGU AU CAAUG
2361

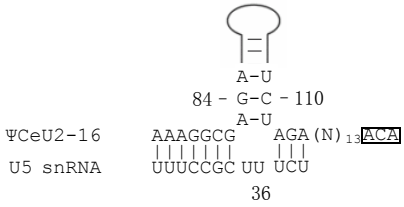

U.G
82 - A-U - 109
A-U
ΨCeLSU-2483 UUUUUGC AGCC(N)11ACA
LSU rRNA AAAAACG GU UCGG
2483

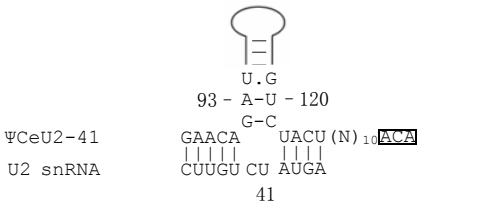

U-A
14 - C-G - 42
G-C
ΨCeLSU-2836 GUACUG UGUCA(N)11AUACCG
LSU rRNA CAUGAC UU UCAGU
2836

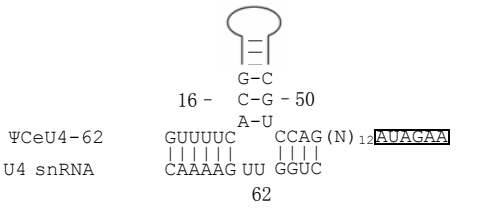

A-U
14 - C-G - 37
C-G
ΨCeLSU-2992 GAUAGGAA GCCGACAUC(N)7AAAUAG
LSU rRNA CUAUCCUU CU CGGCUGUAG
2992

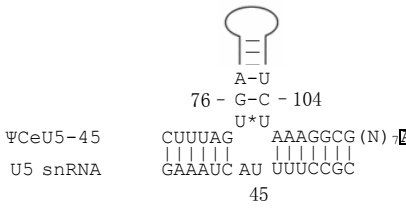

C-G
17 - G-C - 45
G-C
ΨCeLSU-3220 CGUUCAGU UAAUCUUU(N)7AGAUAA
LSU rRNA GCAAGUCA AU AUUAGAGA
3220

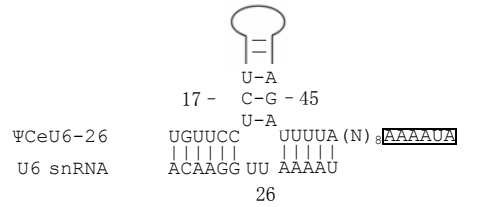

A*A
15 - C-G - 44
A-U
ΨCeU2-16 UAA AGCCGAA(N)7AUAUAA
U2 snRNA AUU AU UCGGCUU
16


A-U
84 - G-C - 110
A-U
ΨCeU2-16 AAAGGCG AGA(N)13ACA
U5 snRNA UUUCCGC UU UCU
36

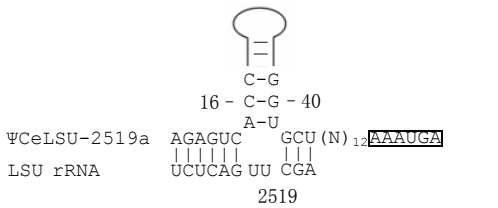

U.G
93 - A-U - 120
G-C
ΨCeU2-41 GAACA UACU(N)10ACA
U2 snRNA CUUGU CU AUGA
41

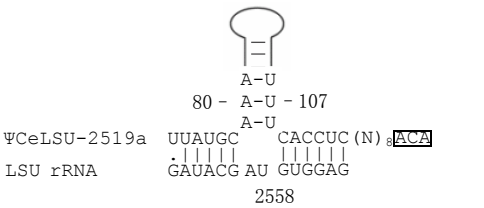

G-C
16 - C-G - 50
A-U
ΨCeU4-62 GUUUUC CCAG(N)12AUAGAA
U4 snRNA CAAAAG UU GGUC
62

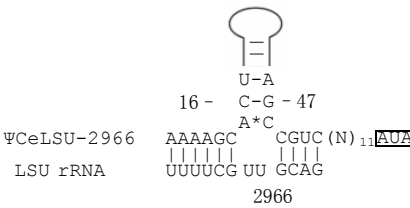

A-U
76 - G-C - 104
U*U
ΨCeU5-45 CUUUAG AAAGGCG(N)7ACA
U5 snRNA GAAAUC AU UUUCCGC
45


U-A
17 - C-G - 45
U-A
ΨCeU6-26 UGUUCC UUUUA(N)8AAAAUA
U6 snRNA ACAAGG UU AAAAU
26

B


C-G
16 - C-G - 40
A-U
ΨCeLSU-2519a AGAGUC GCU(N)12AAAUGA
LSU rRNA UCUCAG UU CGA
2519


A-U
80 - A-U - 107
A-U
ΨCeLSU-2519a UUAUGC CACCUC(N)8ACA
LSU rRNA GAUACG AU GUGGAG
2558


U-A
16 - C-G - 47
A*C
ΨCeLSU-2966 AAAAGC CGUC(N)11AUAAGA
LSU rRNA UUUUCG UU GCAG
2966

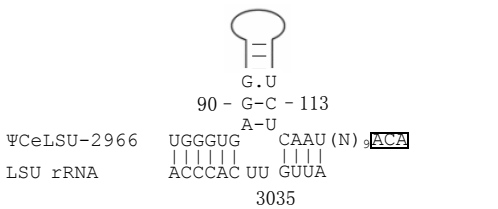
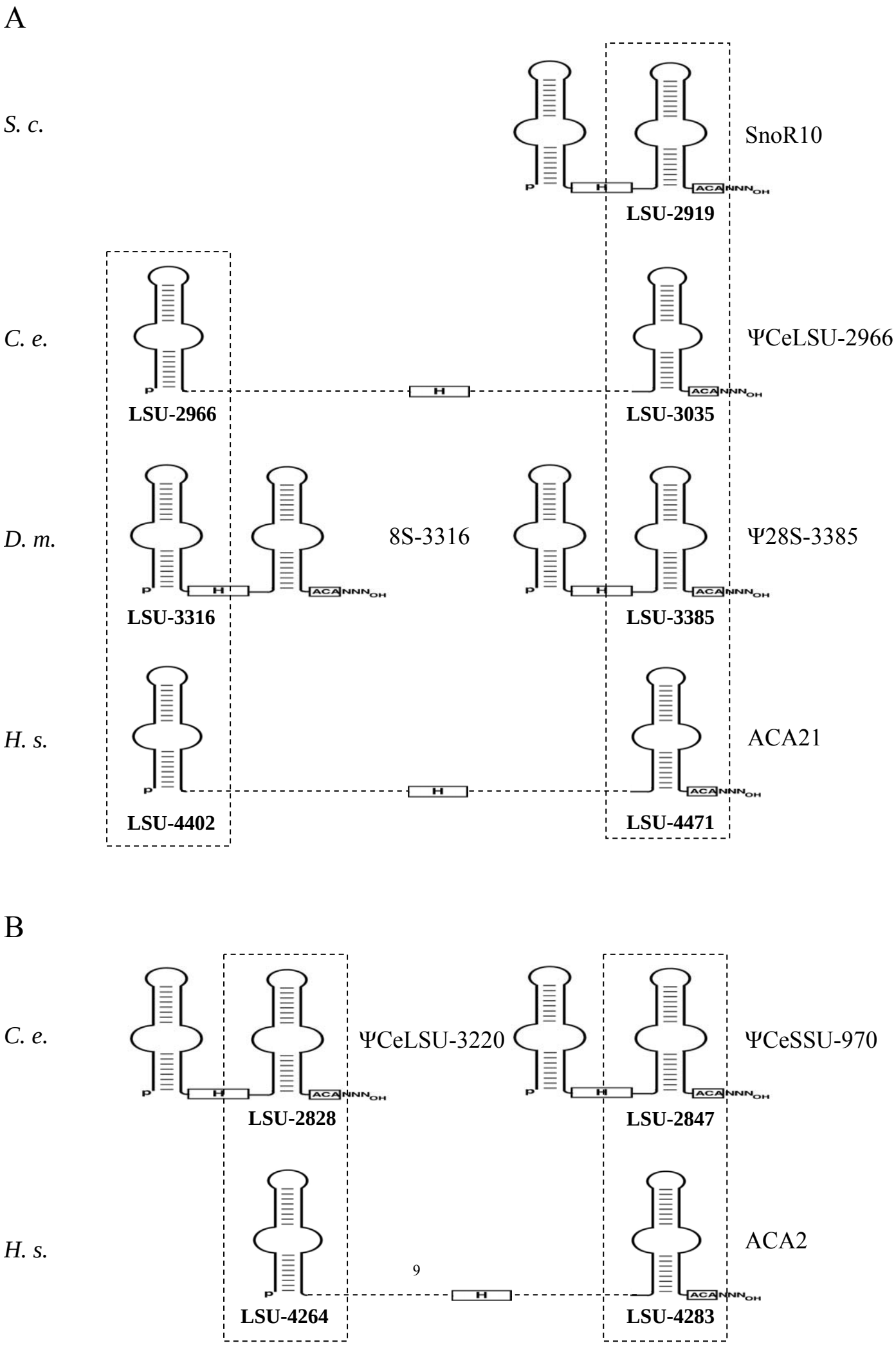

G.U
90 - G-C - 113
A-U
ΨCeLSU-2966 UGGGUG CAAU(N)8ACA
LSU rRNA ACCCAC UU GUUA
3035

Figure S3


A
S. c.
SnoR10
LSU-2919
C. e.
ΨCeLSU-2966
LSU-2966
LSU-3035
D. m.
8S-3316
Ψ28S-3385
LSU-3316
LSU-3385
H. s.
ACA21
LSU-4402
LSU-4471
B
C. e.
ΨCeLSU-3220
ΨCeSSU-970
LSU-2828
LSU-2847
H. s.
ACA2
LSU-4264
LSU-4283