

National Institute of Advanced Industrial Science and Technology

## National Metrology Institute of Japan



## Reference Material Certificate

NMIJ CRM 6204-b  
No. +++

## Ribonucleic Acid (RNA) Solutions for Quantitative Analysis

This certified reference material (CRM) is produced in accordance with the NMIJ's management system and is in compliance with ISO 17034 and ISO/IEC 17025. This CRM consists of five solutions of ribonucleic acid (RNA) having different lengths (533 or 1033 bases of single-strand RNA) and sequences. This CRM is primarily intended to be used in the evaluation, and control of the precision, of RNA analytical instruments and methods such as DNA microarrays (DNA chips), quantitative reverse-transcription PCR, and next-generation DNA sequencers. It can also be used to assign the value of an RNA sample for evaluation of DNA chips.

**Certified Values**

The certified values for the mass concentrations of total RNA (i.e., whole RNA materials in the sample solution regardless of sequence) of the five solutions (RNA500-A, -B, -C, RNA1000-A, and -B) at 25 °C are given in the table below. The uncertainty of the certified value is the expanded uncertainty multiplying the combined standard uncertainty by a coverage factor ( $k$ ) of 2, and it is the half-width of an interval estimated to have a level of confidence of approximately 95 %.

Sample name	Concentration as Total RNA	
	Certified value Mass concentration (ng/μL)	Expanded uncertainty Mass concentration (ng/μL)
RNA500-A	33.4	2.6
RNA500-B	32.3	2.6
RNA500-C	32.1	3.1
RNA1000-A	68.2	5.8
RNA1000-B	64.1	5.5

RNA500-A, -B, -C, RNA1000-A, and -B are the solutions of RNA designed and prepared as shown in Fig. 1 in this certificate.

**Analysis**

The certified values for each material were determined by converting the mass fractions of the total RNA obtained using the following analytical methods into the mass concentration using its density:

## (1) Isotope dilution mass spectrometry (IDMS)

Ribonucleotides, which were obtained from RNA using enzymatic digestion, were quantified by liquid chromatography mass spectrometry (LC/MS). The mass fraction of RNA was calculated from the sum of the ribonucleotides mass fractions considering dehydration at phosphodiester bonds in RNA molecule.

## (2) Inductively coupled plasma mass spectrometry (ICP-MS)

The phosphorus content in each solution was quantified by ICP-MS after acid digestion. The mass fraction of RNA was calculated from the determined mass fraction of the phosphorus content and number of phosphorus including target RNA.

**Metrological Traceability**

The certified values were determined by IDMS with the ribonucleotide standard solutions which were evaluated by <sup>1</sup>H NMR using internal standards calibrated with potassium hydrogen phthalate CRM (NMIJ CRM 3001-b), by ICP-MS with a phosphate ion standard solution from the Japan Calibration Service System (JCSS), and by density measurement with a density standard

liquid (water) from JCSS. The certified values, therefore, are traceable to the International System of Units (SI).

### Indicative Values

The indicative values of this CRM are the amount of substance concentrations for the RNA molecules shown in Fig. 1, in this certificate, at 25 °C are given in the table below. These values were obtained by using the certified values and the base compositions in the RNA molecules shown in Technical Information.

The quoted uncertainty is the expanded uncertainty multiplying the combined standard uncertainty by a coverage factor ( $k$ ) of 2, and it is the half-width of an interval estimated to have a level of confidence of approximately 95 %.

Sample name	Concentration as a targeted RNA molecule	
	Indicative value Amount of substance concentration (pmol/μL)	Expanded uncertainty Amount of substance concentration (pmol/μL)
RNA500-A	0.195	0.014
RNA500-B	0.188	0.014
RNA500-C	0.187	0.017
RNA1000-A	0.205	0.015
RNA1000-B	0.193	0.014

### Expiration of Certification

This certificate is valid for 6 months from the date of shipment, provided that this CRM is stored in accordance with the instructions given in this certificate.

### Description of the material

This CRM is in the form of a clear and colorless liquid containing 533 or 1033 bases synthesized single-stranded RNA dissolved in sterilized water at room temperature. This CRM of ca. 300 μL in net volume is kept in a semi-transparent plastic vial (for low-adsorption of nucleic acids), and a set of five solutions is sealed in a nylon bag hermetically by decompression. The bag is then sealed in an aluminum-laminated plastic bag.

### Instructions for Storage

This CRM should be stored in a freezer at a temperature between -30 °C and -20 °C.

### Instructions for Use

Prior to use, the frozen solution to be analyzed should be allowed to stand at room temperature (25 °C ± 5 °C) until it thaws (heating is strictly forbidden). After confirming that the cap of the vial is tightly closed, the vial is gently inverted several times to ensure complete mixing. The thawed solution should be used immediately and is for single use only. The use of low-adsorption and RNase-free pipette tips and vials is recommended. This CRM is intended for *in vitro* laboratory use only.

### Precautions for Handling

Refer to the safety data sheet (SDS) on this CRM before use.

### Preparation

This CRM was designed, synthesized, purified, and bottled by Biomedical Research Institute, National Institute of Advanced Industrial Science and Technology (AIST). The RNAs in the five solutions have different lengths (533 or 1033 bases of single-strand RNA) and sequences. These random sequences, which do not code for specific genes, were inserted into a plasmid, which was duplicated in *E. coli*. The plasmid was then extracted from *E. coli* and purified. Thereafter, the plasmid was fragmented with a restriction enzyme, and the target RNA sequence was synthesized enzymatically and purified. Then, the purified RNA solutions were diluted by sterilized water. Approximately 300 μL of each solution was bottled in a half transparent plastic vial, and a set of five solutions was kept in a decompression hermetically sealed nylon bag. This bag was then kept in an aluminum-laminated bag.

**Technical Information**

## (1) Sequence analysis and molecular weight

The sequences, molecular, and weights of RNAs for this CRM are shown in Fig. 1. The database accession numbers (DDBJ/GenBank/EMBL) of the template DNA used in the enzymatic synthesis of this CRM are also shown in Fig.1. The sequences of the complementary DNA (cDNA) synthesized using the prepared materials were analyzed using an automated DNA sequencer, and the sequences were confirmed to be the same as the designed one.

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BASE COUNT      165 A    127 C    121 G    120 U
ORIGIN
   1 GGGCUCGACU AGUUAUACG GUACAGGAUA ACCGAUCGGC UUGCAACAUA ACGGCGUUA
  61 GAAUGCGGGA GUGCAGUUUC CGAUUCUCAC AUCAAUCGCC AAUAAGGCCU UGUCGCAUA
 121 UAGACUACAC GGUUCUAGUA GCUGAUCGGU AUUACGUGAC GCAACCGAUU AGACAUGCAC
 181 AAUCCUUGG  UCGCUAUACU ACGGAAAUCG UCAGGUACUA UAACCCGUCG CAGGCCUAAU
 241 ACGUGUCGUC ACAUCGCCAA CCUAUCGUCA GUCGGAAAGA CGUUGCUGUC UACCAUCGAA
 301 ACUAUUUACC GCUCCGAGAU UCACGAGUAC GAACUCACGA GGAAGUUGCC CUAUGUAAGG
 361 UAUCACUCCA GGUACUGCGC CGAUAGUACC AGGUGAUCAA ACGGUUGCAA GAAGGCCACG
 421 ACGUAUCGGG CUCUUUAGAC GUACGCUCGA GAUJAAACGC GCACUGAUUC ACUUUAGCCC
 481 GGAUUGUCUC GGUUGCGAUGU AGAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAA

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Molecular weight: 171 603.8
Accession number: AB610939 (registered as 6204-a-500-1)

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Fig. 1(a). Sequence of RNA molecule in RNA500-A

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BASE COUNT      163 A    130 C    130 G    110 U
ORIGIN
   1 GGGAGACUAA AUCUCGGCGU CGGUUCAUAC GCGCGAUCGU UUGCUGUCAG GGCAUACUCG
  61 AAUCCGGACU CCGACAAUUA UAGGCCAUCC UGAAUAGCCG AUCAUGCGAG UCACGAUAAG
 121 GCAGGCUCUG CGAUAUCCCG AUUACUGGA GAAGCUGAAU CCCACCUAGA GCGAACUGUC
 181 AGAGGAUCGA CCUCAGGCUC GCUAUCAUCA UAACGGCGGA CGACCUGUGU CACAUUCCGA
 241 ACGCUACGUG ACGAUUUUUA CUGUCGAAAG GCAUAGAACG CCGGUCAAUA UCCUGCGGCA
 301 UUCUCUUUUA CACCGCUAU AACUACUAGG UUCCGAGAU AUAGACUGCG CACGGAACAU
 361 GUAGAUAGAU CGAGUAGGGU AGCGAUUUAA CGACUCGACU UACAGACAGA GACGUAGAAC
 421 GUCAGACGAG UGGUAUGCCC ACCAGAGGCG AUACAGGCGU UACCUGCGUA GCACUAGAGU
 481 CGUGCGUCAU GCGGACCCUA UCUAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAA

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Molecular weight: 171 906.1
Accession number: AB610940 (registered as CRM6204-a-500-2)

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Fig. 1(b). Sequence of RNA molecule in RNA500-B

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BASE COUNT      171 A    126 C    116 G    120 U
ORIGIN
   1 GGGCUAAAC GCACUGAAUA CCGUACUACA ACAGACGAAG UUGUAAUAG CCGUGUAAU
  61 UAUGAACGAA UAUGGCCAUG UGUCCGUAA UCCGCGGUAC UAGCCAGUUA GCAACUCGAC

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121 CAAUCGCUCA CGUCAGUGGU UCUAUGCAAU AUGCUCCAGU ACCCUGUAAG UUCGCAAUCA
181 AUAGACGCGC CUUACUCCUC UCAAGAAGGG UAUCUGCAUG AGCCGACACA UCAAGACCCA
241 AUGGACGUUU GAGCGAGUGG CUUGGAGAGU AUUAACGCAC UAACUCUUCG AAGGCUUACU
301 UCGGCAAUUC CGCGAGCUCC ACUAUUAACA UGCCAAUACG ACAGGAUCAA UUCUGCGACU
361 GCACGACCGA AUUAUGCACC UACUUGUGA GGCACGAGAU UCGUCUUGCA GCUAUUUAAA
421 GGGUCCAGC UUAUGGAUAG GCGACUCUUC AGUGCGUAU AAAGCAACGC CCAAUCGGCA
481 UGUUACCGGA UAGUACGGGC GAUAAAAAAA AAAAAAAAAA AAAAAAAAAA AAA

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Molecular weight: 171 547.8

Accession number: AB610942 (registered as 6204-a-500-4)

Fig. 1(c). Sequence of RNA molecule in RNA500-C

BASE COUNT	283 A	258 C	257 G	235 U
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ORIGIN

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1 GGGCGAUUCG AAGAGGUACG AGUGGACGCG UAAGCGAAUG ACCUAGACCU CGGCGUUAU
61 UAGGACCCUC UAAUCGAAA CUCGACUCUC GUCCCAAUCC AAUGGAUGUC CAGUGCUCGG
121 UAGCAUGAUC GUAUGAUGCG UAUCGUCGCG AGUAGAGGCC GACAAGUAGA CCGGUGCGAA
181 UUUGGAGGUA CUUAGCCUCA UAUGAGAGCG CCUUGAAAUC ACCCAGUGCC GAUCGUAGCG
241 GAAGAUUACU AGACUCCGCA GGGAAAUCCC ACCUGUAACG ACGGAAGAGC GUCACGAUAG
301 CCUCUAAUA UCCGGUUCGC GACUAUCCGC UUAUGUGCCU CCACCUAAUG UGAGAGUUA
361 CCGAGGCAA UGAUCUGUCA ACCGGUGUGA UCAGGACAU CGCUUAAUG CGUAGAAGCC
421 CGUAAGCUCU CCGCCUUA AGAGGUUGUA GACGGCAGUU CUAAGGUCGU CGGGUCUAUG
481 CCUUGCGACC UAAUAAUACG ACCGUGUGCU UAUGCGGACU GUCCUCUAAU GAAUAUCGCU
541 UGUCCUAAAG UGGCGUACU AGUGCUUAGG AUCGCACACC UCACCACAGU GCGCAUUUA
601 CCCUGUAGAU AACAUUGUAG ACACCGGUA AUCGCGUUCG AAUUUCGCC AAUCGAAGGC
661 CCACAUCACU ACGUCGCCUG UAUCUGAAC CUUGCGUCG ACGUAGCAUA UAGAGCGUAC
721 AUUCAAUUA CCAGUUGCCU CCGACUGAAG UCGGCUAGCG UAUGACAUAG CGAGCUCUUA
781 GUUCGGUGAC UACUUCUAGC ACUCCCAAU CAAGCUCUGC GUUAUCAGGG UCGGAAGGUU
841 AGGUUCGAU UUCGACAGGC UAACAGAGCG AUAGUGAUG AAUCCGUCC GGGAGCAUCU
901 AGACAAUAC CGCGUUAAG AGAAGGGCG CAUAAGCGCG GGUGUCAACG UUCAACCAG
961 UUGUAGCCAU CGCGAUUACC CGUUGGAAU CUGAGGCGAC CUAAAAAAAA AAAAAAAAAA
1021 AAAAAAAAA AAA

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Molecular weight: 332 585.9

Accession number: AB610946 (registered as 6204-a-1000-3)

Fig. 1(d). Sequence of RNA molecule in RNA1000-A

BASE COUNT	267 A	262 C	245 G	259 U
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ORIGIN

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1 GGGAUUCCUA GGACUGUACU CUCGGUGCGU UGACCAUACG UAAGGCGAUC CUUUGAGUGG
61 AUCCAAUAC UACGCGUCAC ACCUGCUUAC CCUCCAAUA GUUGGUUCAG UAGCUCUCAG
121 CGGUUCUGGC AGAGUUCGGA UGAGUUUCUG CCUAUCAGUU CAUAGGUGCC CACGCAUUGG
181 GUCCACUCCU CGCCAGAAU UGCGCAUUGC ACCAUUACUA CAGGCGGCUU UGGUUGUACG

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241 UCUAACGUUC GCACCAACAG GAGUCUCAGC UGAUCAUAGG CCCGGACCCU CAAUGUUCGA
301 UGCGAUUCGU AAGAGGGUGU UCGUGUAAGG CCCAAUACGU UGUCAUGCCG GCUUAGAAAC
361 CCAGUCGGAC GCGUCUCUAA CACUCGGAUG UGCAGGUAU AGCCUUUACC AGCGCUUCUG
421 UACGACCAUA CUUAGAGCUC GAGAUGCCGA CAUGAAAGGA UUCCGGAGUA CUGACCUGAA
481 UACACGUUA UAGCGUAAU CGGCCGAGAU UCAACUUUAC GGCACGGAUA CAGCUCUCU
541 ACCUAUUUCC GUCGAAGUCU CUCACGAUAG UCGCGUACAU UUAGUGGGCG GUACACACAG
601 CACGUCAACG CCAUCGCACU CUGAGUJCCC ACUCCACGGU ACGUUCACAG CACGUUGCCU
661 UAAUAAGCUA CUUCGGUJCC GAGCAGUCAA CCUACUGUUU CCGGGUJAGC GCUCUGAUCA
721 GCACCCGUUU ACUGACACGA ACCGCUAUCG AAUACUGAGU AGGUCGUGUG CCAAUAAUJ
781 UGGUUGCAGC UAAGCUAAUJ GGACGGCGAC UJUAGCAAGU AACUCAGCCG UAUUGUJACG
841 CUGACCGUAA ACGACGUGAG CGAUUGUCGU AGGUUAGCCA UAACAUAJAG GUJUJCCGAA
901 CGGUAGCAUA GUUAGGCCUG UGUCCAGUCA GGUAAUJACGA GAGAGUAJAU AACGCGAUJU
961 AAUGAGAAGC CGUGCAUGUC GAUCCUJGUU ACGGGUGUGA AAUAJAAAAA AJAAAAAJAA
1021 AJAAAAAJAA AJA
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Molecular weight: 331 744.9

Accession number: AB610947 (registered as 6204-a-1000-4)

Fig. 1(e). Sequence of RNA molecule in RNA1000-B

## (2) Gel electrophoresis

The CRM material gave single bands near 533 or 1033 base using microchip gel electrophoresis.

## (3) Density measurement

The density of these CRM solutions at 25 °C is 0.9971 g/cm<sup>3</sup>.

**NMIJ Analysts**

The technical manager is for this CRM is TAKATSU A., the production manager is FUJII S., and the analysts are FUJII S., SHIBAYAMA S., INAGAKI K., YAMAZAKI T., SEKIGUCHI Y., NODA N., MATSUKURA S., SASAKI A. and YOSHIOKA M.

**Information**

If substantive technical changes occur that affect the certification before the expiration of this certificate, NMIJ will notify the registered customer. Customer registration on the NMIJ Website (given below) will facilitate notification. Technical reports regarding this CRM can be obtained from the contact details given below.

**Reproduction of Certificate**

In reproducing this certificate, it should be clearly indicated that the document is a copy.

April 1, 2020

ISHIMURA Kazuhiko

President

National Institute of Advanced Industrial Science and Technology

If you have any questions about this CRM, please contact:  
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Sample