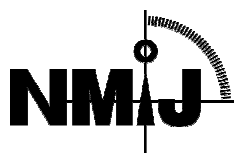


National Institute of Advanced Industrial Science and Technology

National Metrology Institute of Japan

Reference Material Certificate



NMIJ CRM 6204-a
No. +++



Ribonucleic Acid (RNA) Solutions for Quantitative Analysis

This certified reference material (CRM) was produced in accordance with the NMIJ's management system, and in compliance with ISO GUIDE 34:2009 and ISO/IEC 17025:2005. This CRM consists of five kinds of ribonucleic acid (RNA) solutions having different lengths (533 or 1033 bases of single-strand RNA) and sequences. This CRM is principally intended to be used to assign the value of an RNA calibrator for the evaluation and control of the precision of RNA analytical methods such as DNA microarray (DNA chip), quantitative reverse-transcription PCR method, and next-generation DNA sequencer.

Certified Values

The certified values of five solutions (RNA500-A, -B, -C, RNA1000-A, and -B) for the mass concentration of total RNA (whole RNA materials in the sample solution regardless of sequence) at 25 °C are given in the table below. The uncertainty of the certified value is the half-width of the expanded uncertainty interval calculated using a coverage factor (k) of 2, which gives a level of confidence of approximately 95 %. Purity (in mass fraction) of L-arginine is given as follows.

Sample name	Mass concentration of total RNA	
	Certified value (ng/μL)	Expanded uncertainty (ng/μL)
RNA500-A	30.6	3.1
RNA500-B	27.3	2.4
RNA500-C	32.4	3.2
RNA1000-A	58.3	4.9
RNA1000-B	59.5	5.3

Analysis

The certified values for each material were based on the results obtained by the following analytical methods:

(1) Isotope dilution-mass spectrometry (ID-MS)

Ribonucleotides, which were obtained from RNA by using enzymatic digestion, were quantified by liquid chromatography mass spectrometry (LC/MS).

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

Phosphorus in the solution was quantified by ICP-MS after acid digestion.

The mass concentration of the total RNA was calculated from the obtained mass fraction of the total RNA and density of the solution.

Metrological Traceability

Each certified value was traceable to the International System of Units (SI) via ribonucleotides analysis based on ID-MS as the primary method of measurement and phosphorus analysis using the phosphate ion standard solution of the Japan Calibration Service System (JCSS). The standard solution of ribonucleotides, the purity of which was determined by impurity determination and phosphorus analysis using ICP-optical emission spectrometry, was employed.

Indicative Values

The amount of substance concentration of RNA molecules at 25 °C is given in the table below as an indicative value, which

was calculated using the certified value and base content in RNA molecules shown in Technical Information. The uncertainty of the indicative value is the half-width of the expanded uncertainty interval calculated using a coverage factor (k) of 2, which gives a level of confidence of approximately 95 %.

Sample name	Amount of substance concentration of RNA	
	Indicative value (pmol/ μ L)	Expanded uncertainty (pmol/ μ L)
RNA500-A	0.178	0.019
RNA500-B	0.159	0.016
RNA500-C	0.189	0.020
RNA1000-A	0.175	0.015
RNA1000-B	0.179	0.018

Expiration of Certification

This certificate is valid for 3 months from the date of shipment, provided that the material remains unopened and stored in accordance with the instructions given in this certificate.

Sample Form

This CRM is in the form of a clear and colorless liquid at room temperature. Approximately 300 μ L 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.

Homogeneity

The homogeneity of the CRM was evaluated by measuring RNA by spectrophotometric analysis, analyzing 10 vials randomly selected from about 110 vials. The uncertainty of homogeneity was reflected in the uncertainty of the certified value.

Stability

The stability of the CRM was confirmed by NMIJ and the uncertainty of stability was reflected in the uncertainty of the certified value. The stability will be monitored.

Instruction for Storage

This CRM should be kept in a freezer (temperature lower than -20 °C) after delivery.

Instructions for Use

Prior to use, the frozen solution to be analyzed should be allowed to stand at room temperature (about 20 °C) until it thaws (heating is strictly forbidden). After confirming that the cap of the vial is tightly closed, the vial is turned upside down gently several times for complete mixing. The thawed solution should be used immediately and is for single use only. The thawed solution should be sampled using low-adsorption and RNase-free pipette tips and vials. NMIJ CRM 6204 is intended for *in vitro* laboratory use only.

Precaution for Handling

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

Preparation Method

This CRM was designed, synthesized, purified, and bottled by Biomedical Research Institute, National Institute of Advanced Industrial Science and Technology (AIST). RNAs in five solutions had different lengths (533 or 1033 bases of single-strand RNA) and sequences. These random sequences, which were not coding-specific genes, were inserted into a plasmid and the plasmid 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.

Information

(1) Sequence analysis and molecular weight

The sequence, molecular weight, and accession number of the database concerning the sequence (DDBJ/GenBank/EMBL) of this CRM is 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 of all kinds of materials 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 GAAUCCGGGA GUGCAGUUC CGAUUCUCAC AUCAAUCGCC AAUAAGGCCU UGUCGCAUA
 121 UAGACUCAAC GGUUCUAGUA GCUGAUCGGU AUUACGUGAC GCAACCGAUU AGACAUGCAC
 181 AAUCCUUGG UCGCUAUACU ACGGAAUUCG UCAGGUACUA UAACCCGUCG CAGGCCUUAU
 241 ACGUGUCGUC ACAUCGCCAA CCUAUCGUCA GUCGGAAAGA CGUUGCUGUC UACCAUCGAA
 301 ACUAUUUACC GCUCCGAGAU UCACGAGUAC GAACUCACGA GGAAGUUGCC CUAUGUAAGG
 361 UAUACAUCCA GGUACUUGCG CGAUAGUACC AGGUGAUCAA ACGGUUGCAA GAAGGCCACG
 421 ACGUAUCGGG CUCUUUAGAC GUACGCUCGA GAUUAACGC GCACUGAUUC ACUUUAGCCC
 481 GAAUGUCUC GGUGCGAUGU AGAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAA

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Molecular weight: 171 603.8

Accession number: AB610939 (registered as 500-1)

Fig. 1(a). RNA500-A

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BASE COUNT      163 a    130 c    130 g    110 u
ORIGIN
   1 GGGAGACUAA AUCUCGGCGU CGGUUCAUAC GCGCGAUCGU UUGCUGUCAG GCGAUACUCG
  61 AAUCCGGACU CCGACAUAUA UAGGCCAUCC UGAAUAGCCG AUCAUGCGAG UCACGUAUAG
 121 GCAGGCUCUG CGAUAUCCCG AUUAUCUGGA GAAGCUGAAU CCCACCUAGA GCGAACUGUC
 181 AGAGGAUCGA CCUCAGGCUC GCUAUCAUCA UAACGGCGGA CGACCUUGU CACAUUCCGA
 241 ACGCUACGUG ACGAUUUUAU CUGUCGAAAG GCAUAGAACG CCGGUAUAU UCUGCGGCA
 301 UUCUCUUUAU CACCGGCUAU AACUACUAGG UUCCGCAGAU AUAGACUGCG CACGGAACAU
 361 GUAGAUAGAU CGAGUAGGGU AGCGAUUUAA CGACUCGACU UACAGACAGA GACGUAGAAC
 421 GUCAGACGAG UGUUAGGCC ACCAGAGGCG AUACAGGCUG UACCUUGCUA GCACUAGAGU
 481 CGUGCGUCAU GCGGACCCUA UCUAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAA

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Molecular weight: 171 906.1

Accession number: AB610940 (registered as 500-2)

Fig. 1(b). RNA500-B

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BASE COUNT      171 a    126 c    116 g    120 u
ORIGIN
   1 GGGACUAAAC GCACUGAAUA CCGUACUACA ACAGACGAAG UUGUAAAUAG CCGUGGUAU
  61 UAUGAACGAA UAUGGCCAUG UGUCCGCUAA UCCGCGGUAC UAGCCAGUUA GCAACUGCAC
 121 CAUUCGCUCA CGUCAGUGGU UCUAUGCAAU AUGCUCCAGU ACCCUGUAAG UUCGCAAUCA
 181 AUAGACGGC CUUACUCCUC UCAAGAAGGG UAUCUGCAUG AGCCGACACA UCAAGACCCA
 241 AUGGACGUUU GAGCGAGUGG CUUGGAGAGU AUUAACGCAC UAACUCUUCG AAGGCUUACU
 301 UCGGCAAAUC CGCGAGCUCC ACUAUUUACA UGCCAAUACG ACAGGAUCA UUCUGCGACU
 361 GCAGACCGA AUUAUGCACC UACUUUGUGA GGCACGAGAU UCGUCUUGCA GCUUUUUAAA
 421 GGGUCCAGC UUAUGGAUAG GCGACUCUUC AGUGCGUAU AAAGCAACGC CCAUUCGGCA
 481 UGUUACCGGA UAGUACGGC GAUAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAA

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Molecular weight: 171 547.8

Accession number: AB610942 (registered as 500-4)

Fig. 1(c). RNA500-C

BASE COUNT 283 a 258 c 257 g 235 u
 ORIGIN
 1 GGGCGAUUCG AAGAGGUACG AGUGGACGCG UAAGCGAAUG ACCUAGACCU CGGCGUUAU
 61 UAGGACCCUC UAAUCGCAA CUCGACUCUC GUCCCAAUCC AAUGGAUGUC CAGUGCUCGG
 121 UAGCAUGAUC GUAUGAUGCG UAUCGCUGCG AGUAGAGGCC GACAAGUAGA CCGGUGCGAA
 181 UUGGAGGUA CUUAGCCUCA UAUGAGAGCG CCUUGAAAUC ACCCAGUGCC GAUCGUAGCG
 241 GAAGAUUACU AGACUCCGCA GGGAAAUCCC ACCUGUAACG ACGGAAGAGC GUCACGAUAG
 301 CCUCUACUA UCCGGUUCGC GACUAUCCGC UUAUGUGCCU CCACCUAAUG UGAGAGUUA
 361 CCGAGGCAA UGAUCUGUCA ACCGGUGUGA UCAGGACAUU CGCUJAAUGC CGUAGAAGCC
 421 CGUAAGCUCU CCGCCUUAU AGAGGUUGUA GACGGCAGUU CUAAGGUCGU CGGGUCUAUG
 481 CCUUGCGACC JAAUAAUACG ACCGUGUGCU UAUGCGGACU GUCCUCUAAU GAAUAUCGCU
 541 UGUCCUAGC UGGCGGUACU AGUGCUUAGG AUCGCACACC UCACCACAGU GCGCAUUUAA
 601 CCCUGUAGAU AACAUUGUAG ACACCGUAA AUCGCGUUCG AAUUUCGCC AAUCGAAGGC
 661 CCACAUACU ACGUCGCCUG UAUUCUGAAC CUUGCGUCG ACGUAGCAUA UAGAGCGUAC
 721 AUUCAUACU CCAGUUGCCU CCGACUGAAG UCGCUAGCG UAUGACAUAG CGAGCUCUUA
 781 GUUCGGUGAC UACUUCUAGC ACUCCAAU CAAGCUCUG GUUAUCAGGG UCGGAAGGU
 841 AGGUUCGAU UUCGACAGG UAACAGAGCG AUAAGUGAUG AAUCGCUCC GGGAGCAUCU
 901 AGACAAUAC CGCGUUAAG AGAAGGGCGA CAUAAGCGCG GGUGUCAACG UUCAACCAG
 961 UUGUAGCCAU CGCGAUUACC CGUUGGAAU CUGAGGCGAC CUAAAAAAAAA AAAAAAAAA
 1021 AAAAAAAAA AAA

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Molecular weight: 332 585.9

Accession number: AB610946 (registered as 1000-3)

Fig. 1(d). RNA100-A

BASE COUNT 267 a 262 c 245 g 259 u
 ORIGIN
 1 GGGAUUCCUA GGACUGUACU CUCGGUGCGU UGACCAUACG UAAGGCGAUC CUUUGAGUG
 61 AUCCAAUAC UACGCGUCAC ACCUGCUUAC CCUCCAAUA GUUGGUUCAG UAGCUCUCAG
 121 CGGUUCUGG AGAGUUCGGA UGAGUUUCUG CCUAUCAGU CAUAGGUGCC CACGCAUUGG
 181 GUCCACUCCU GCAGAAAU UGCGCAUUGC ACCAUUACUA CAGGCGGCUU UGGUUGUACG
 241 UCUAACGUUC GCACCAACAG GAGUCUCAGC UGAUCAUAGG CCCGGACCCU CAAUUGUCGA
 301 UGCGAUUCGU AAGAGGGUGU UCGUGUAAG CCCAAUACGU UGUCAUGCCG GCUUAGAAAC
 361 CCAGUCGGAC GCGUCUCUAA CACUCGGAUG UGCAGGUAU AGCCUUUACC AGCGCUUCUG
 421 UACGACCAUA CUUAGAGCUC GAGAUCCGA CAUGAAAGGA UUCGCGGUA CUGACCUGAA
 481 UACACGUUCA UAGCGUAAU CGGCCGAGAU UCAACUUUAC GGCACGGAUA CAGCUCUCU
 541 ACCUAAUUC GUCCGAAGUCU CUCACGAUAG UCGCGUACAU UUAGUUGGCG GUACACACAG
 601 CACGUCAACG CCAUCGCACU CUGAGUJCCC ACUCCACGGU ACGUUCACAG CACGUUGCCU
 661 UAAUAAGCUA CUUCGGUUC GAGCAGUCA CUAUCUGUU CCGGGUAGC GCUCUGAUCA
 721 GCACCCGUU ACUGACACGA ACCGCUAUCG AAUACUGAGU AGGUCGUGUG CCAAUAACU
 781 UGGUUGCAGC UAAGCUAAU GACGCGGAC UUUAGCAAG AACUCAGCCG UAUUGUUCG
 841 CUGACCGUAA ACGACGUGAG CGAUUGUCGU AGGUUAGCCA UAACAUAAG GUUUCGGA
 901 CGGUAGCAUA GUUAGGCCUG UGUCCAGUCA GGUAAUACGA GAGAGUAAU AACGCGAUCU
 961 AAUGAGAAGC CGUGCAUGUC GAUCCUUGU ACGGGUGUGA AAUAAAAAAAA AAAAAAAAA
 1021 AAAAAAAAA AAA

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Molecular weight: 331 744.9

Accession number: AB610947 (registered as 1000-4)

Fig. 1(e). RNA100-B

(2) Gel electrophoresis

The single bands near 533 or 1033 bases were obtained using polyacrylamide gel electrophoresis and microchip gel electrophoresis.

(3) Density measurement

The density of these solutions at 25 °C was 0.9971 g/cm³.

NMIJ Analysts

For this CRM, the technical manager is A. Takatsu, the production manager is S. Fujii, and the analysts are S. Fujii, S. Shibayama, K. Inagaki, T. Narukawa, Y. Sekiguchi, M. Kawaharasaki, and M. Yoshioka.

Technical Information

Customer registration on the NMIJ Website (given below) will facilitate notification of any revision of the information given above. 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, 2015

Ryoji Chubachi
President

National Institute of Advanced Industrial Science and Technology

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

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National Metrology Institute of Japan,

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Phone: +81-29-861-4059; Fax: +81-29-861-4009; <https://www.nmij.jp/english/service/C/>

Revision history

April 1, 2015: “Metrology Management Center” was renamed to “Center for Quality Management of Metrology.”