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Short communication

Insights into the development of effective materials to suppress replication of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)

Fumiaki Uchiumi^{1,*}, Yoko Ogino¹, Akira Sato² and Sei-ichi Tanuma³

- ¹ Department of Gene Regulation, Faculty of Pharmaceutical Sciences, Tokyo University of Science, Noda-shi, Chiba-ken 278-8510, Japan
- ² Department of Biochemistry, Faculty of Pharmaceutical Sciences, Tokyo University of Science, Noda-shi, Chiba-ken 278-8510, Japan
- ³ Department of Genomic Medicinal Science, Research Institute for Science and Technology, Organization for Research Advancement, Tokyo University of Science, Noda-shi, Chiba-ken 278-8510, Japan
- * Correspondence: Email: f_uchiumi@rs.tus.ac.jp; Tel: +81471213616; Fax: +81471213608.

Abstract: The life cycle of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), which is the etiologic agent for coronavirus disease-19 (COVID-19), has been revealed, and the molecular mechanism of how this virus replicates has been studied. Effective drugs could be designed to target one of the replication processes. This virus, which contains plus-strand RNA as a genome, is characteristic because it synthesizes its genome and other transcripts with minus-strand template RNA. These viral RNAs, including its genome, are transcribed by an RNA-dependent RNA polymerase (RdRp). In this article, we discuss whether novel materials from RNA derivatives could be applied to COVID-19 patients to prevent or disturb viral transcription.

Keywords: COVID-19; miRNA; RdRp; SARS-CoV-2; transcription

Molecular biologists have learned that the transcription of genes is executed by proteins or protein complexes, including TBP, TFIID, and an RNA polymerase II [1]. Elongation of the RNA chain requires ribonucleotide triphosphates as substrates, and an RNA primer is essentially required in in vitro reactions [1]. However, how the primers for the transcription of genes are provided in the

cells remains unclear because it is quite obvious that they are present in every cell from its beginning as an egg.

Coronavirus consists of several proteins and a positive-strand RNA (+RNA) as a genome [2]. The complete sequence of the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), which causes COVID-19, has been revealed (https://www.ncbi.nlm.nih.gov/nuccore/1798174254). Whole genome sequencing of SARS-CoV-2 indicated that it is most closely related to the bat coronavirus SL-CoVZC45 [3]. It contains several open reading frames that encode viral proteins [2,3]. To replicate the viral genome, the +RNA chain needs to be transcribed first by an RNA-dependent RNA polymerase (RdRp), which is a non-structural protein 12 (nsp12) encoded on the viral genome [4–6]. The transcripts, including minus-strand RNA (-RNA) and templates for translation of the viral proteins, are thought to be produced by the RNA polymerizing enzyme RdRp.

The RdRp enzyme needs a primer to start transcription [4]. Although it has been speculated that the nsp7-nsp8 complex confers RNA-primer synthesis [7], the kind of primers required in the first step to polymerize ribonucleotides has not been precisely indicated or described. We, as molecular biologists, have learned that life has been evolved through the RNA world [8,9]. Cells must contain short RNAs from the beginning. Presumably, the nucleotide sequences of the viral genome will give us a hint in developing a new strategy to suppress or disturb its replication system. Previously, based on the surveillance of the SARS-coronavirus genome, it was reported that a specific nucleotide sequence 5'-ACGAAC-3' is present near the translation start sites or first AUG of the open reading frames [10].

To predict possible competitors that could affect replication of the +RNA strand, the 5'-UTR sequence (1 to 265) or the most 5'-upstream of the +RNA was converted to the complimentary -RNA strand, and it was surveyed by the miRBase program (miRbase.org/index.shtml). A few miRNAs were retrieved from this surveillance; they are hairpin RNAs, including hsa-mir-2053 and hsa-mir-668. Of them, hsa-mir-2053 has a sequence in the loop that may hybridize with the most 5'-upstream of the SARS-CoV-2 +RNA. Interestingly, hsa-mir-2053 has recently been reported to be a tumor suppressor miRNA [11]. The hairpin RNA hsa-mir-668 plays a role in the suppression of mitochondrial fragmentation by suppressing MTP18 gene expression to protect from acute kidney injury [12,13]. We should also remember that some of the miRNAs have antiviral functions [14]. These observations have inspired us to hypothesize that miRNAs may prevent or hinder the initiation or elongation of the +RNA strand on the template -RNA. The clinical data show that worsening of the COVID-19 viral infectious disease appears to occur in older people [15]. This might be due to lower amounts of transcription suppressor miRNAs because the expression profile of miRNAs in lung is altered in accordance with aging [16].

Molecular biologists have learned that DNA sequences can be read by the Sanger method [17]. The DNA sequencing analysis achieved by this method has not only contributed to the scientific field but also to society. Although the analytical equipment has been greatly improved from the original procedure, the sequence of SARS-CoV-2 was revealed by this method. The reaction for the DNA sequencing uses dideoxy nucleotides ATP, GTP, CTP, and TTP. The elongation of the DNA strand stops by incorporation of the dideoxy nucleotide into the 3'-ends. The Honorable Dr. Frederic Sanger might have left a hint for us to develop a new strategy against viral infectious disease. If transcription is initiated by RNA polymerases, including by RdRp, it must require short RNAs that are originally present in host cells (Figure 1A). Moreover, the inhibitory effect of 3'-deoxyribofuranosyl nucleotides (3'-deoxy UTP and 3'-deoxy TTP) could be synthesized

chemically [18] to be added to the 3'-end of the RNA primers. Therefore, 3'-end deoxy RNA may prevent elongation of the transcripts.

For example, 5'-auuaaagguuuauaccuucccagguaacaa(deoxy-a)-3', which could be hybridized with hsa-mir-2053, would prevent the initiation of +RNA transcription. If other 3'-end deoxy RNA primers were annealed to the -RNA strand, no elongation would occur (Figure 1B). The 3'-end deoxy-RNA primers that hybridize to the adjacent sites of the transcription start sites for viral RNAs should be examined (Figure 1B). These 3'-end deoxy-RNA primers would not be so harmful compared with the use of RNA polymerase inhibitory nucleotide derivatives, including favipiravir [19], because the 3'-end deoxy-RNA primers specifically target the transcription start sites of the viral genome. However, the length of the 3'-end deoxy-RNA primers should be determined. Moreover, not only the very first 5'-end of the genome but also the most effective sites at which the artificial aberrant RNA primers anneal on the -RNA strand must be investigated. Additionally, to improve stability, incorporation of modified ribonucleotides should be examined. Such modified ribonucleotides have been already studied [20].

Antisense octaguanidinium dendrimer-conjugated morpholino oligomers showed an inhibitory effect on enterovirus 71, the genome of which is like that of coronavirus, a single +RNA [21]. Because the 3'-end deoxy (modified) RNA primers are short unelongatable sense strands, they are not equal to the antisense RNAs. The false bait RNA primers rather resemble short 5'-capped oligomer RNAs with 3'-phosphates, which inhibit elongation of the +RNA strand on the minus strand template genome of the influenza virus [22]. However, it would be essential to identify the most frequently occurring transcription initiation site(s) on the -RNA strand of the coronavirus. Quantitative analysis or read count plots [3] of the viral transcripts could provide a way to find the precise sequence. It should be noted that the exo- and endoribonucleases nsp14 and 15, respectively, are encoded on the coronaviral genome [23]. As for the influenza virus, it was suggested that endo-RNA nucleases detach inappropriate ribonucleotide to elongate the RNA strand from the nick [22]. Therefore, the length of the false bait RNAs should be examined. Moreover, to effectively suppress elongation, the development of specific inhibitors for nsp14 and 15 is required.

Next, how should these RNAs be applied clinically to patients? Presently, some of the RNAs or miRNAs have been clinically tested to treat lung cancer [24] and could be applied by inhalation. The most difficult impediment to overcome would be the current pharmaceutical regulations, which, of course, must be obeyed. Because these RNAs originally exist in cells, they might be approved as processed nutrients or supplements. However, because such artificial RNAs have never been tested clinically, we cannot predict the consequences in the future. The discovery of antibiotics greatly contributed to the treatment of bacterial infectious diseases, but their repeated use has resulted in the emergence of antibiotic-resistant bacteria [25]. This kind of problem must be considered and avoided.





Figure 1. 3'-deoxy (modified) RNA primers disturb the initiation and elongation of viral RNA strands. (A) Minus-strand RNA (-RNA) is synthesized by RdRp. (B) (Left) Plus-strand RNAs (+RNAs) are transcribed from the template -RNA. (Right) Excess amounts of 3'-deoxy (modified) RNAs that specifically bind to the -RNA prevent elongation of the viral genome and +RNAs that should be translated to viral proteins.

Not only the 3'-deoxy ribonucleotides but also 3'-modified nucleotides could be applied to the inelongatable RNA primers. For example, 3'-fluoro-3'-deoxyadenosine has been shown to have a suppressive effect on double-stranded RNA viruses [26]. The essential tasks are to construct several 3'-deoxy (or modified)-RNA primers and to examine whether they restrict viral replication and are not harmful to normal healthy cells. Further, the conditions for inhalation must be followed according to established procedures [24]. The sooner this can begin, the better. The authors hope that some of the scientists and physicians involved in drug development will focus on RNA-based therapeutics, and we emphasize that studies on transcription will greatly contribute to the development of effective medicines in the future [27,28]. We further hope that scientists, researchers, medical institutions, chemical and pharmaceutical companies, governments, and anyone who has been inspired by this article take full advantage of all available resources at hand to develop medications to fight COVID-19.

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Conflict of interest

The authors declare no conflict of interest.

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