

**Molecular Physiology Laboratory (2021)**  
**Chief Scientist: Rikiya Watanabe (Ph.D.)**



**(0) Research field**

CPR Subcommittee: Biology

**Keywords:** membrane protein, artificial cell membrane, artificial cell, bioMEMS, digital biology

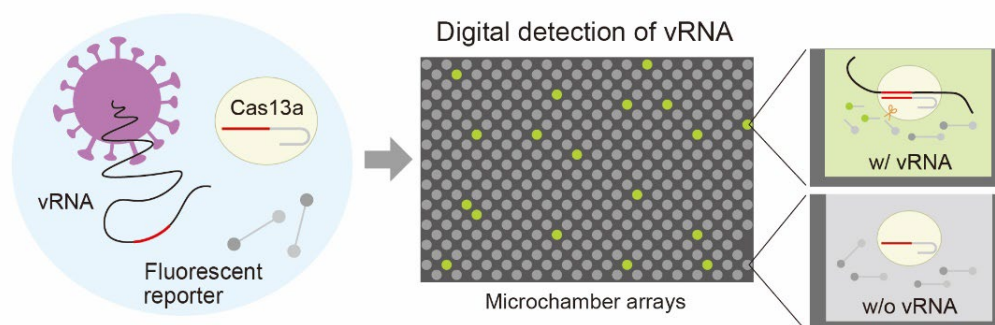
**(1) Long-term goal of laboratory and research background**

Our study aims to understand cellular functions using a bottom-up approach from the single molecule level. To achieve this, we are attempting to elucidate the mechanism by which individual biomolecules or their networks function in a precise manner, by developing novel single-molecule techniques using multidisciplinary approaches, including biophysics, bioMEMS, and chemical biology. In addition, we are developing a methodology to investigate correlations between genetic mutations, dysfunctions, and diseases with single molecule sensitivity, which would provide new insights for biological as well as pharmaceutical studies.

**(2) Current research activities (FY2021)**

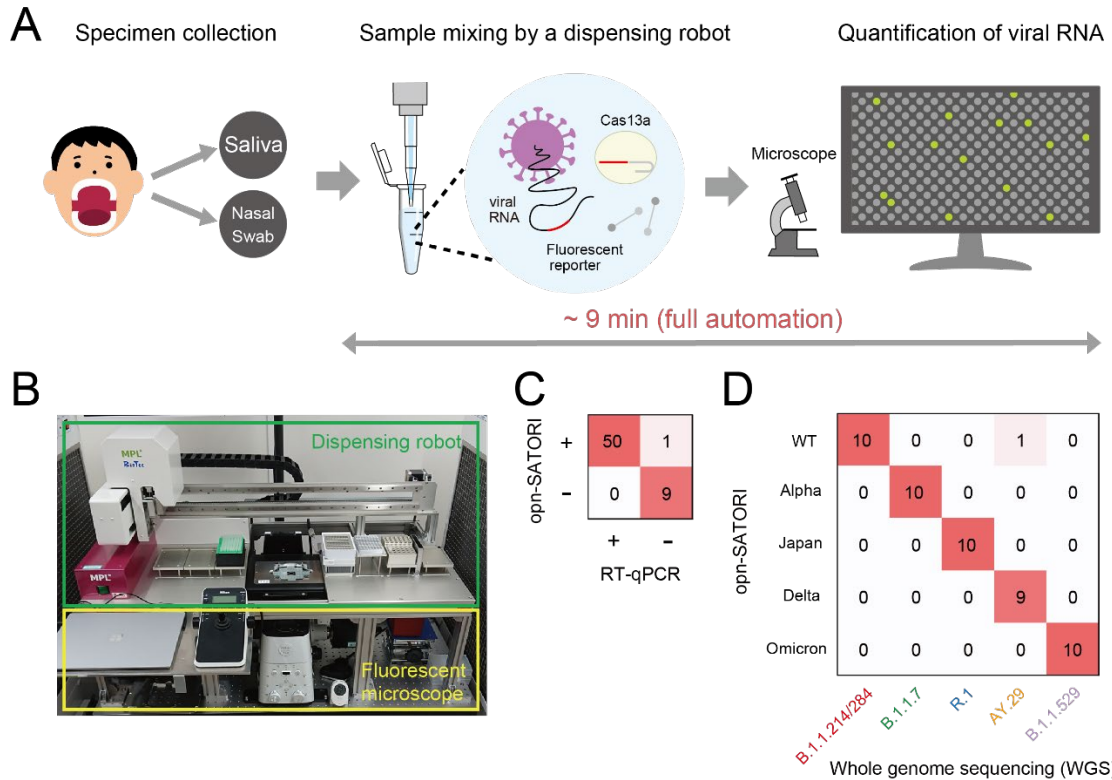
**Development of automated platform for rapid and sensitive COVID-19 diagnosis**

In the ongoing COVID-19 pandemic, there is an urgent need to establish a versatile diagnostic method for viral infections. The PCR and antigen tests are widely used for the diagnosis of viral infections; however, these methods generally have technical drawbacks in terms of sensitivity, accuracy, and throughput, making it difficult to efficiently analyze a large volume of specimens with high sensitivity and accuracy. To address these issues, we have successfully developed an innovative technology using CRISPR-Cas13a (SATORI method)<sup>1</sup> that can detect viral RNA from SARS-CoV-2 at single molecule level with high sensitivity and throughput (the fastest in the world) (Fig. 1). The SATORI method was useful for rapid and sensitive diagnosis of COVID-19; however, full automation of the diagnostic process was essential for its implementation in society.



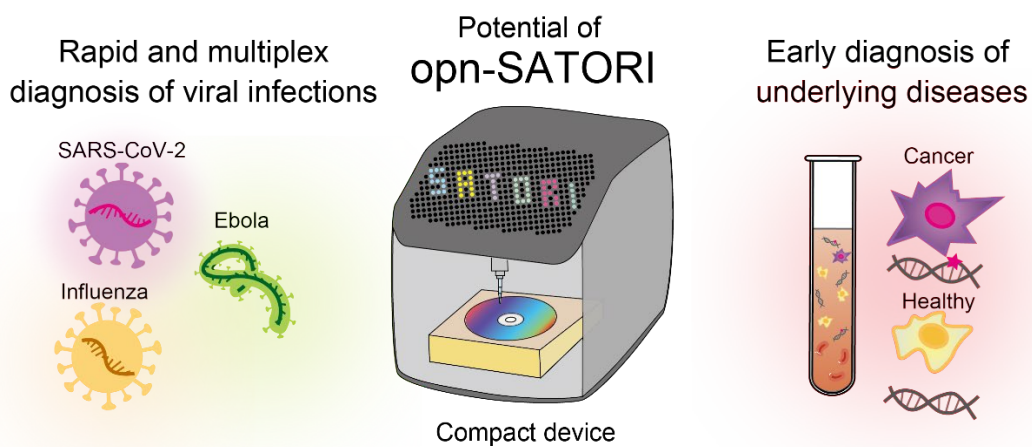
**Fig. 1 Amplification-free rapid detection platform of viral RNA (SATORI method)**

Last fiscal year (FY2021), we developed an automated platform on SATORI (opn-SATORI)<sup>2</sup> for fully automated diagnosis of viral infections using the SATORI method by incorporating the custom-made dispensing robot (Fig. 2). Using clinical specimens of COVID-19, we demonstrated that opn-SATORI can detect viral RNA from SARS-CoV-2 with high sensitivity (1.4 copies/ $\mu$ L) comparable to that of PCR, and discriminate between SARS-CoV-2 variants of concern in  $\sim$  9 min with an accuracy of 98%. In addition, by developing plastic microchips and optimizing assay reagents, we have succeeded in reducing the cost of opn-SATORI to less than \$2 per test, making it ideal for social implementation.



**Fig. 2 Automated platform on SATORI (opn-SATORI) for COVID-19 diagnosis**  
 (A) Schematics of COVID-19 diagnosis, (B) Overall view, (C, D) Comparison of opn-SATORI and PCR/WGS results in SARS-CoV-2 detection and variant discrimination.

We demonstrated that opn-SATORI is a sensitive, rapid, and automated platform for viral RNA detection that can precisely diagnose SARS-CoV-2 and its variants. Given that CRISPR-Cas13a-based methods have been used to detect RNA biomarkers for cancer diagnosis, opn-SATORI may serve as a versatile platform for diverse applications, including diagnosis of viral infections and evaluation of disease-related biomarkers for liquid biopsy (Fig. 3). We believe that opn-SATORI will be a key technology for high-throughput genetic diagnostics.



**Fig. 3 Potential of opn-SATORI for liquid biopsy**

### (3) Members

#### (Chief Scientist)

Rikiya Watanabe

#### (Research Scientist)

Jun Ando, Hajime Shinoda

#### (Special Postdoctoral Researcher)

Yoshiaki Kinoshita

#### (Technical Staff)

Asami Makino, Chiharu Takahashi,

Tatsuya Iida, Mami Yoshimura

### (4) Representative research achievements

1. “Amplification-free RNA detection with CRISPR-Cas13”, Shinoda, H., Taguchi, Y., Nakagawa, R., Makino, A., Okazaki, S., Nakano, M., Muramoto, Y., Takahashi, C., Takahashi, I., Ando, J., Noda, T., \*Nureki, O., \*Nishimasu, H., & \*Watanabe, R., *Commun. Biol.*, 4, 476 (2021)
2. “Automated amplification-free digital RNA detection platform for rapid and sensitive SARS-CoV-2 diagnosis”, Shinoda, H., Iida, T., Makino, A., Yoshimura, M., Ishikawa, J., Ando, J., Murai, K., Sugiyama, K., Muramoto, Y., Nakano, M., Kiga, K., Cui, L., Nureki, O., Takauechi, H., Noda, T., \*Nishimasu, H., & \*Watanabe, R., *Commun. Biol.*, 5, 473 (2022)

### Laboratory Homepage

[https://www.riken.jp/research/labs/chief/mol\\_physiol/index.html](https://www.riken.jp/research/labs/chief/mol_physiol/index.html)

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