



강태준 (Taejoon Kang)
한국생명공학연구원

[Biosens. Bioelectron., Available online 7 July 2023 | https://doi.org/10.1016/j.bios.2023.115522](https://doi.org/10.1016/j.bios.2023.115522)

One-pot, Ultrasensitive, and Multiplex Detection of SARS-CoV-2 Genes Utilizing Self-Priming Hairpin-Mediated Isothermal Amplification

Authors and Affiliations

Yan Li ^a, Taejoon Kang ^{b,c}, Hyun Gyu Park ^a

^aDepartment of Chemical and Biomolecular Engineering (BK21 Four), Korea Advanced Institute of Science and Technology (KAIST), 291 Daehak-ro, Yuseong-gu, Daejeon, 34141, Republic of Korea

^bBionanotechnology Research Center, Korea Research Institute of Bioscience and Biotechnology (KRIBB), 125 Gwahak-ro, Yuseong-gu, Daejeon, 34141, Republic of Korea


^cSchool of Pharmacy, Sungkyunkwan University, 2066 Seobu-ro, Jangan-gu, Suwon, Gyeonggi-do, 16419, Republic of Korea

Corresponding authors: Taejoon Kang, Hyun Gyu Park

Abstract

The global pandemic resulting from the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) and its emerging variants highlights the need for convenient and accurate detection protocols to facilitate timely prevention and management of the disease. Herein, we propose a new self-priming hairpin-mediated isothermal amplification (SIAM) protocol enabling one-pot and ultrasensitive identification of SARS-CoV-2 in a multiplexed way. This approach works by targeting a specific RNA sequence with a self-priming hairpin (SP) probe and promoting continuously repeated extension and nicking reactions to produce numerous trigger molecules, which could specifically bind to molecular beacons (MBs) and produce fluorescent signals. Under an isothermal condition of 37 °C, this technique allowed for the simultaneous identification of the spike (S) and nucleocapsid (N) genes of SARS-CoV-2 down to single copy/μL levels. We further validated the practical diagnostic capabilities of the SIAM method by accurately testing 20 clinical samples with 100% sensitivity and specificity. The SIAM method has a lot of potential to be a reliable nucleic acid testing protocol to identify infections caused by a wide range of pathogens.

논문정보

- 형식: Research article
- 게재일: 2023년 07월 (BRIC 등록일 2023.07.15)
- 연구진: 국내연구진 
- 분야: Biotechnology, Nanobio

Citing URL: https://www.ibric.org/myboard/read.php?Board=hbs_treatise&id=89064&ttype=0&idauthorid=8567