

# Prevention of the Spread of Virus Infection in Global Societies: Anti-SARS-CoV-2 Agents Learned from HIV Researches

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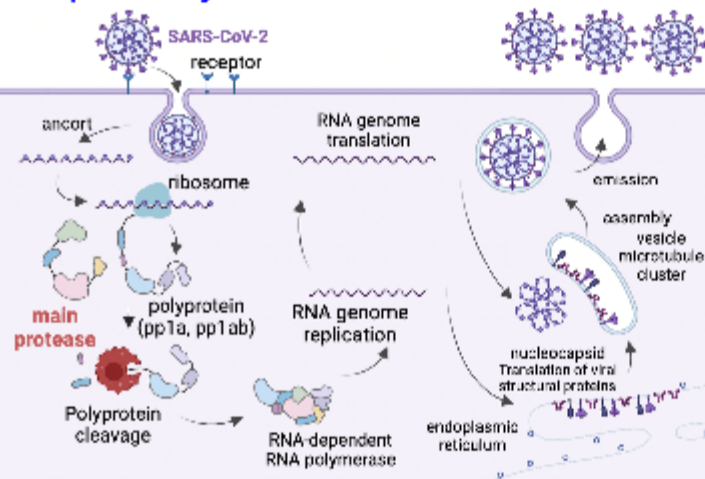


## Introduction

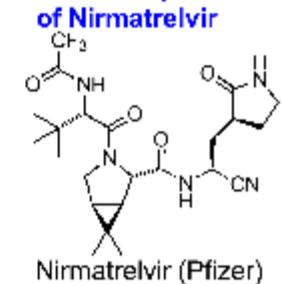
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More than three years have passed since the beginning of the pandemic of the novel emerging infection disease, COVID-19 which is produced by SARS-CoV-2. Since SARS-CoV-2 main protease (M<sup>pro</sup>) is essential for viral replication and there is no human enzyme closely homologous with the M<sup>pro</sup>, it is an important and attractive drug discovery target for treatment of COVID-19. Recently, inhibitors of the SARS-CoV-2 M<sup>pro</sup>, Nirmatrelvir and Ensitrelvir, were developed. However recent studies warned of the possibility of the emergence of drug-resistant SARS-CoV-2 variants against the Nirmatrelvir therefore development of novel drugs and an increase in the repository of the drugs are urgently needed. In order to meet this challenge, we previously reported highly potent M<sup>pro</sup> inhibitors, TKB198 and TKB211, based on YH-53/5h, which was developed by Hayashi et al. in 2013 as a SARS-CoV M<sup>pro</sup> inhibitor. More recently, our further studies led to the development of more potent inhibitors, TKB245 and TKB248. In this study, we report our structure-activity relationship studies and evaluation of the anti-SARS-CoV-2 activity of derivatives of these inhibitors.

### Replication cycle of SARS-CoV-2



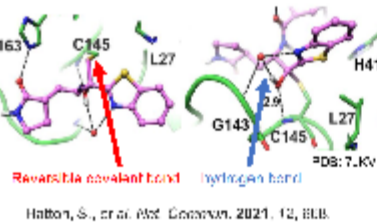
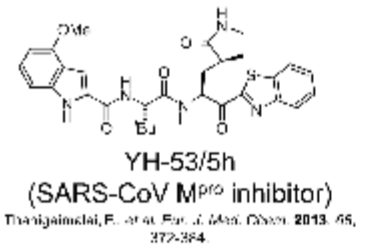
### Potential problems of Nirmatrelvir



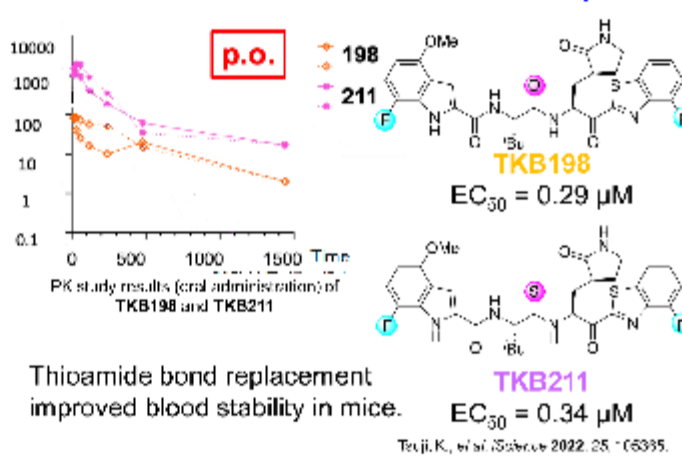
**Problems**  
• antiviral activity  
• metabolic stability

→ Solve these problems

### YH-53/5h

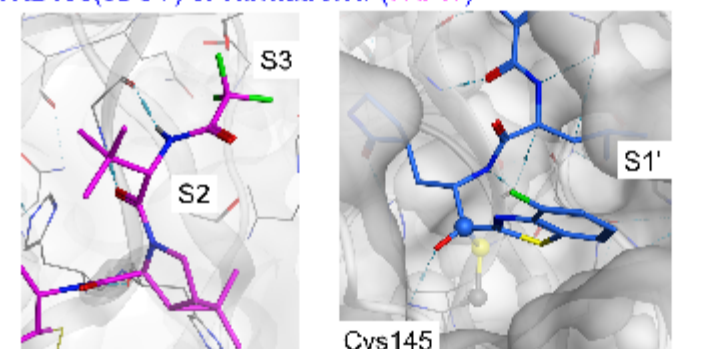


### Fluorination and thioamidation of the lead compound



## TKB245 & TKB248

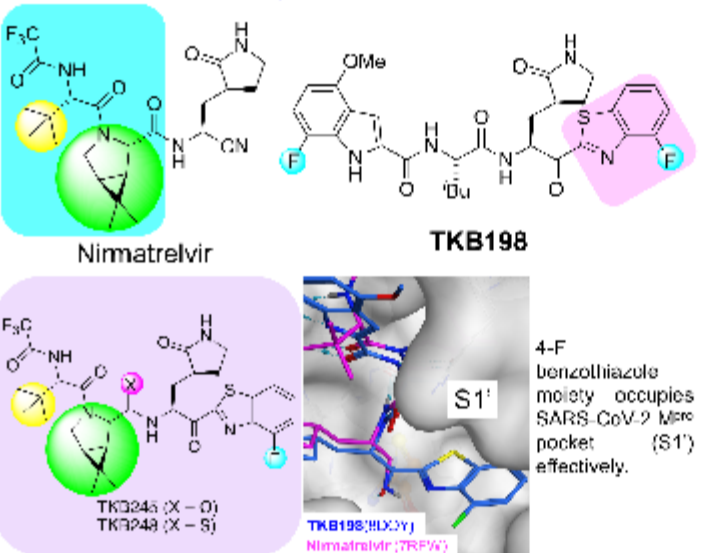
### Co-crystal structures of SARS-CoV-2 M<sup>pro</sup> with TKB198(8DOY) or Nirmatrelvir (7RFW)



Nirmatrelvir has numerous interactions in the S2-S3 region of SARS-CoV-2 M<sup>pro</sup>.

TKB198 forms a covalent bond with Cys145, the active center of SARS-CoV-2 M<sup>pro</sup>. Also, it inactivates the enzymatic activity of M<sup>pro</sup>.

### Hybridization of SARS-CoV-2 inhibitors with Nirmatrelvir and TKB198

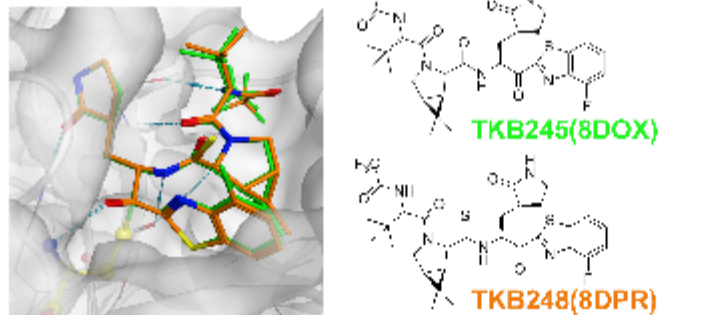


### Biological evaluation of TKB245 and TKB248

| Compound     | EC <sub>50</sub> (μM) | EC <sub>90</sub> (μM) | IC <sub>50</sub> (μM) | IC <sub>90</sub> (μM) | T <sub>1/2</sub> (h) | Oral F (%) |
|--------------|-----------------------|-----------------------|-----------------------|-----------------------|----------------------|------------|
| YH-53        | 2.6                   | 9.47                  | 0.13                  | 1.5                   | 0.27                 | n.d.       |
| TKB198       | 0.27                  | 0.83                  | 0.023                 | 0.75                  | 0.83                 | 1.98       |
| TKB245       | 0.03                  | 0.53                  | 0.007                 | 0.14                  | 3.82                 | 48         |
| TKB248       | 0.22                  | 0.87                  | 0.074                 | 6                     | 4.34                 | 72         |
| Nirmatrelvir | 0.94                  | 7.81                  | 0.013                 | 0.77                  | 1.03                 | 56         |

\*Oral F = AUC oral data after oral administration / AUC per unit dose after intravenous administration.  
\*PK test was performed using human liver primary mice (TKB245, TKB248, and Nirmatrelvir) and Jc11CR mice (YH-53 and TKB198).  
\*EC<sub>50</sub>: Drug concentration to inhibit half of the multiplication of SARS-CoV-2.

### Co-crystal structures of SARS-CoV-2 M<sup>pro</sup> with TKB245 or TKB248



## Results & Discussion

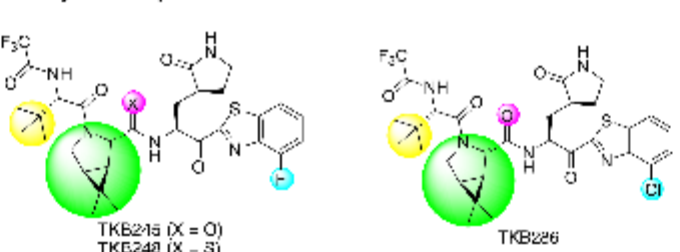
### Structure-activity relationship studies

| Compound | R <sup>1</sup> | R <sup>2</sup> | R <sup>3</sup> | X | EC <sub>50</sub> (μM) | CC <sub>50</sub> (μM) |
|----------|----------------|----------------|----------------|---|-----------------------|-----------------------|
| TKB245   | tert-butyl     | isopropyl      | F              | O | 0.012                 | > 100                 |
| TKB248   | tert-butyl     | isopropyl      | F              | S | 0.31                  | > 100                 |
| TKB269   | tert-butyl     | isopropyl      | F              | O | 0.36                  | > 100                 |
| TKB255   | tert-butyl     | isopropyl      | F              | O | 0.095                 | > 100                 |
| TKB263   | tert-butyl     | isopropyl      | F              | O | 2.4                   | > 100                 |
| TKB253   | tert-butyl     | isopropyl      | F              | O | 0.25                  | > 100                 |
| TKB267   | tert-butyl     | isopropyl      | F              | O | 2.8                   | > 100                 |
| TKB265   | tert-butyl     | isopropyl      | F              | O | 0.59                  | > 100                 |
| TKB286   | tert-butyl     | isopropyl      | Cl             | O | 0.019                 | > 100                 |
| TKB287   | tert-butyl     | isopropyl      | Br             | O | 0.063                 | > 100                 |

### Short summary

R<sup>1</sup>: L-tert-Leu residue is suitable.  
R<sup>2</sup>: is the best amino acid among the tested derivatives.  
R<sup>3</sup>: F and Cl is tolerable.

TKB245 and TKB248 are still the most promising compounds. In addition, we found TKB286, which the activity is comparable to that of TKB245.



### Antiviral activity in various target cells

| Cell line              | TKB245 | TKB248 | Nirmatrelvir | Inhibitor controls |
|------------------------|--------|--------|--------------|--------------------|
| Ver0L6                 | 0.03   | 0.22   | 1.02         | 0.11               |
| Ver0E6 + 2 μM CP100356 | 0.001  | 0.03   | 0.025        | -                  |
| HeLa-ACE2-TMPRSS2      | 0.021  | 0.16   | 0.21         | 0.2                |
| A549-ACE2-TMPRSS2      | 0.0027 | 0.199  | 0.017        | -                  |

\*CP100356 is P-glycoprotein inhibitor

### Stability against acid treatment

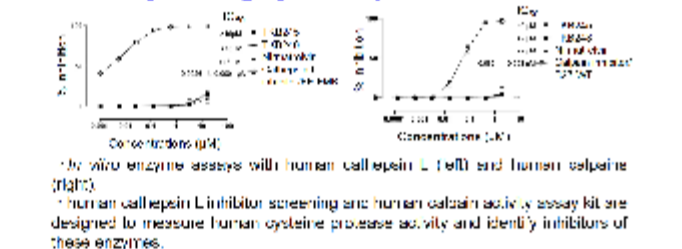
| Compound     | EC <sub>50</sub> (μM) |          | CC <sub>50</sub> (μM) |
|--------------|-----------------------|----------|-----------------------|
|              | acid (-)              | acid (+) |                       |
| TKB245       | 0.074                 | 0.087    | >100                  |
| TKB248       | 0.31                  | 0.32     | >100                  |
| Nirmatrelvir | 1.4                   | 2.5      | >100                  |

\*The acid (-) means test compounds dissolved in PBS without acid treatment.  
\*The acid (+) means test compounds treated with 0.1 M HCl aq. in PBS, incubated for 1h at 37 °C, and then neutralized with 0.1 M NaOH aq. in the mixture.

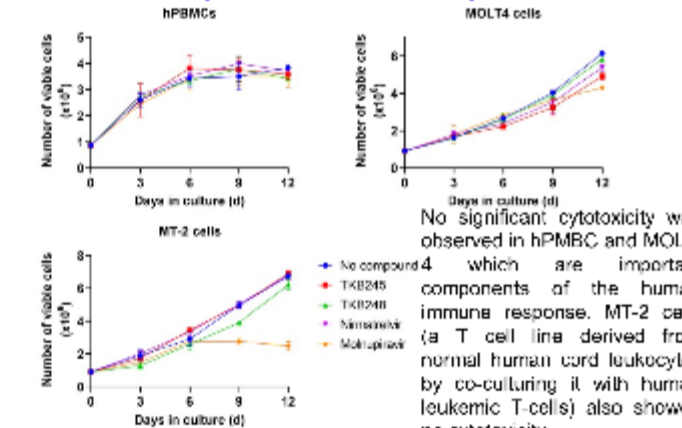
### Antiviral activity against mutant strains

| Compound     | EC <sub>50</sub> (μM) |       |       |       |          |
|--------------|-----------------------|-------|-------|-------|----------|
|              | Wk-521                | oBA.1 | oBA.2 | oBA.5 | oBA.2.75 |
| TKB245       | 0.03                  | 0.014 | 0.052 | 0.016 | 0.049    |
| TKB248       | 0.22                  | 0.282 | 0.16  | 0.066 | 0.43     |
| Nirmatrelvir | 0.94                  | 1.047 | 1.3   | 1.126 | 0.955    |

### Selectivity among cysteine proteases



### Evaluation of potential immune response



Higashi-Kuwata, N., et al. Nat. Commun. 2023, 14, 1079.  
Tsuji, K., et al. J. Med. Chem. 2023, 66, 13516-13526.

## Conclusion & Acknowledgement

As the results of the SAR studies of TKB245, a 4-fluoro or 4-chlorobenzothiazole ketone was suitable as the P1' unit. Biological evaluation demonstrated that TKB245 and TKB248 showed preferable acid stability in the test conditions and no significant cytotoxicity in hPMBCs, MOLT-4 cells and MT-2 cells. These data supported that TKB245 and TKB248 can be orally administrable drug candidates for COVID-19 therapy as M<sup>pro</sup> inhibitors. We deeply thank to Prof. Yoshio Hayashi and Dr. Sho Konno from Tokyo University of Pharmacy and Life Sciences for the valuable discussions.