

Manidipine is not a potential inhibitor against SARS-CoV-2 main protease

Rui Zhang,¹ Jiahao Zhou,¹ Haohao Yan,¹ Xiaoping Liu,¹ Chao Shang,² Yunyu Chen¹

AUTHOR AFFILIATIONS See affiliation list on p. 2.

KEYWORDS COVID-19, SARS-CoV-2, main protease inhibitor, fluorescence polarization, manidipine

The rapid spread of the coronavirus disease 2019 (COVID-19) pandemic caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection highlights an urgent need for developing antivirals against coronavirus. An attractive drug target for COVID-19 is the main protease (Mpro), as this conserved enzyme plays a key role in virus replication and host immune evasion (1, 2). Given the urgency of identifying antivirals, repurposing approved drugs as effective Mpro inhibitors represent a promising strategy for COVID-19 treatment (3, 4). Recently, manidipine was identified as a potent Mpro inhibitor with the half-maximal inhibitory concentration (IC₅₀) of 10.4 ± 1.6 μM using fluorescence resonance energy transfer (FRET) assay (5). Nonetheless, a recent study suggested that manidipine is likely to be a promiscuous Mpro inhibitor because it is prone to colloidal aggregation in the enzymatic assay, resulting in the inactivation of Mpro enzyme (6). Considering the potential of manidipine in COVID-19 treatment, a rigorous validation for its Mpro inhibition is necessary.

For this purpose, we evaluated the *in vitro* inhibition of Mpro by manidipine using a robust high-throughput screening (HTS) platform, including FRET, fluorescence polarization (FP), and dimerization-dependent red fluorescent protein (ddRFP) assays (7–9). When we repeated the FRET assay developed by the authors (5), the IC₅₀ value of manidipine was 24.14 ± 3.41 μM (Fig. 1A and B). In clear contrast, manidipine showed no inhibition in the FRET assay when Tween-20 was present in the reaction buffer (IC₅₀ > 200 μM), suggesting that its Mpro inhibition is false positive (Fig. 1C). To further test this possibility, we separately evaluated Mpro inhibition by manidipine using FP and ddRFP assays in the reaction buffer containing 0.05% Tween-20. As expected, manidipine did not inhibit Mpro in both assays (IC₅₀ > 200 μM) (Fig. 1D and E). Consistently, manidipine exhibited no detectable inhibition of Mpro using SDS-PAGE because the ddRFP biosensor could be cleaved into RFP-A₁ and RFP-B₁ fragments in the presence of manidipine (Fig. 1F). Thus, our results invalidated manidipine as a potential Mpro inhibitor *in vitro*. Intriguingly, a recent study independently confirmed our results using cell-based assays (10). In the presence of detergent, the aggregation caused by compounds is minimized in the enzymatic assay (11). Therefore, our results demonstrated the necessity for the addition of detergent such as Tween-20 or Triton X-100 into the assay buffer when assessing Mpro inhibitors, which could disrupt colloids and keep Mpro enzyme stable. Moreover, manidipine might have other targets to achieve the anti-SARS-CoV-2 effect, which can be further studied.

In summary, our data indicate that manidipine is not a potential inhibitor against Mpro based on the results from a set of *in vitro* assays. These results suggest that the colloidal aggregation caused by compounds should be considered when evaluating Mpro inhibitors. It is necessary to verify the inhibitory activity of candidate Mpro inhibitors using diverse biochemical assays due to the limitations of the FRET approach.

Editor Miguel Angel Martinez, IrsiCaixa Institut de Recerca de la Sida, Badalona, Barcelona, Spain

Address correspondence to Chao Shang, shangchao1290@126.com, or Yunyu Chen, chenyunyu1984@163.com.

Rui Zhang and Jiahao Zhou contributed equally to this article. Author order was determined by drawing straws.

The authors declare no conflict of interest.

See the funding table on p. 3.

Ed. Note: The authors of the published article did not respond at the time of publication.

See the original article at <https://doi.org/10.1128/aac.02577-20>.

Published 31 January 2024

Copyright © 2024 American Society for Microbiology. All Rights Reserved.

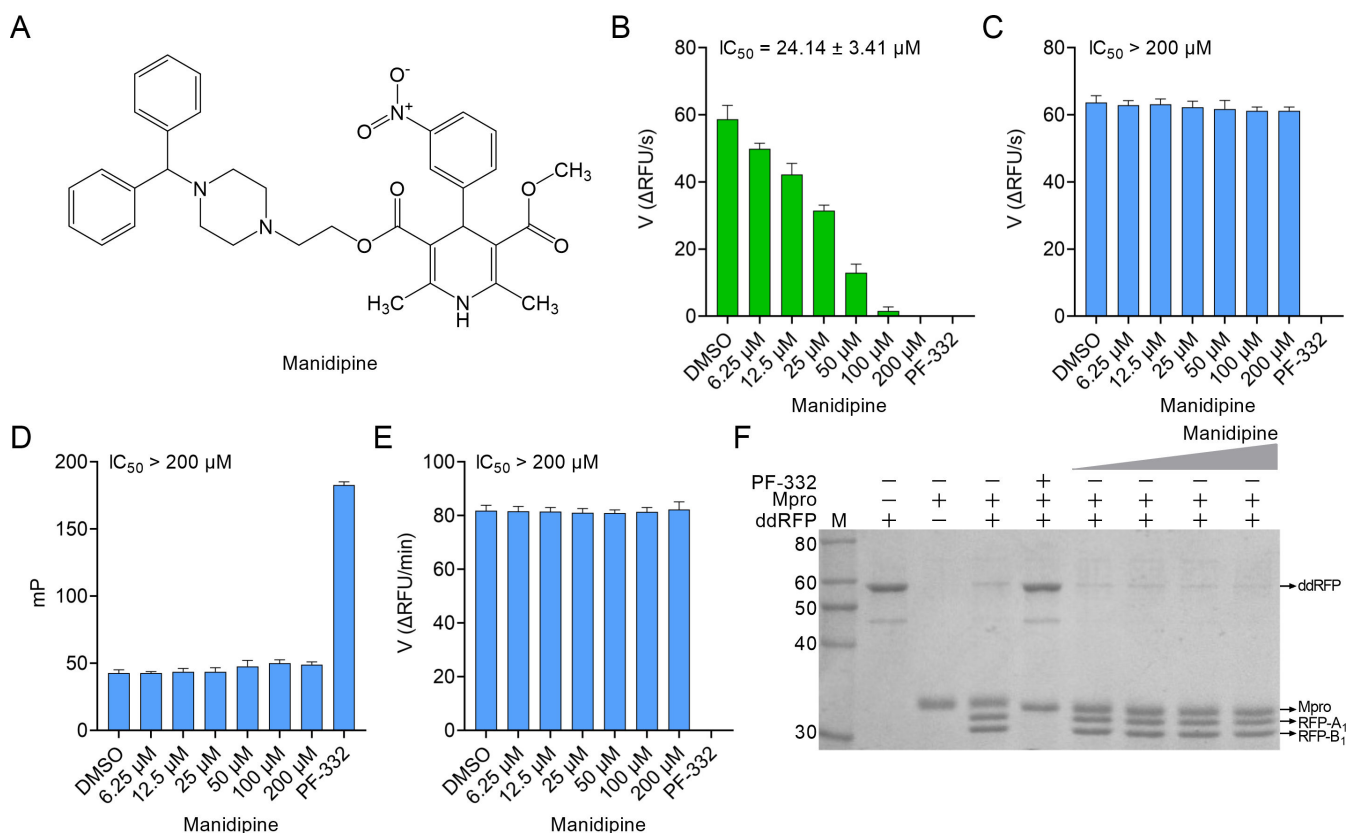


FIG 1 Inhibition of SARS-CoV-2 main protease (Mpro) by manidipine *in vitro*. (A) The chemical structure of manidipine. (B) Evaluation of the activity of manidipine against Mpro using FRET assay developed by authors (5). (C and D) Evaluation of the activity of manidipine against Mpro using FRET and FP assays in the reaction buffer containing 0.05% Tween-20. In these biochemical assays, nirmatrelvir (PF-332, 1 μ M) and DMSO served as the positive and negative controls, respectively. The IC_{50} value of manidipine was shown. (E) Inhibition of Mpro by manidipine through monitoring the initial velocity of Mpro enzyme reaction initiated by ddRFP biosensor. The IC_{50} value of manidipine was shown. (F) Gel-based assay of ddRFP biosensor cleavage by manidipine *in vitro*. In the absence of Mpro inhibitors, the ddRFP biosensor (55 kDa) can be cleaved by Mpro (34 kDa) to generate RFP-A₁ (top band, 29 kDa) and RFP-B₁ fragments (bottom band, 26 kDa) in the gels. Nirmatrelvir (PF-332) and manidipine were purchased from TargetMol (Shanghai, China). The concentrations of manidipine used in this experiment were 25, 50, 100, or 200 μ M. In the ddRFP assay, nirmatrelvir (PF-332, 10 μ M) and DMSO served as the positive and negative controls, respectively. All assays were conducted as previously described (7–9, 12). The reaction buffer was 10 mM HEPES, 50 mM NaCl, 1 mM EDTA, 1 mM dithiothreitol (DTT), and 0.05% Tween-20 (vol/vol), pH7.0.

ACKNOWLEDGMENTS

This work was supported by University Natural Science Research Project of Anhui Province, China (No. KJ2021A0839); The Young Fellow Program of Wannan Medical College, China (No. wyqnyx202104); and Postgraduate Academic Innovation Program of Anhui Province, China (No. 2022xscx129).

The authors thank Prof. Yanchang Wang (Department of Biomedical Sciences, College of Medicine, Florida State University, Tallahassee, USA) for his insightful comments of the manuscript.

AUTHOR AFFILIATIONS

¹Institute for Drug Screening and Evaluation, Wannan Medical College, Wuhu, China
²Changchun Veterinary Research Institute, Chinese Academy of Agricultural Sciences, Changchun, China

AUTHOR ORCIDs

Chao Shang <http://orcid.org/0000-0002-9006-8492>

Yunyu Chen  <http://orcid.org/0000-0002-2750-8221>

FUNDING

Funder	Grant(s)	Author(s)
University Natural Science Research Project of Anhui Province	KJ2021A0839	Yunyu Chen
The Young Fellow Program of Wannan Medical College, China	wyqnyx202104	Yunyu Chen
Postgraduate Academic Innovation Program of Anhui Province, China	2022xscx129	Haohao Yan

REFERENCES

- Anirudhan V, Lee H, Cheng H, Cooper L, Rong L. 2021. Targeting SARS-CoV-2 viral proteases as a therapeutic strategy to treat COVID-19. *J Med Virol* 93:2722–2734. <https://doi.org/10.1002/jmv.26814>
- Zhu W, Shyr Z, Lo DC, Zheng W. 2021. Viral proteases as targets for coronavirus disease 2019 drug development. *J Pharmacol Exp Ther* 378:166–172. <https://doi.org/10.1124/jpet.121.000688>
- Li G, Hilgenfeld R, Whitley R, De Clercq E. 2023. Therapeutic strategies for COVID-19: progress and lessons learned. *Nat Rev Drug Discov* 22:449–475. <https://doi.org/10.1038/s41573-023-00672-y>
- Aherfi S, Pradines B, Devaux C, Honore S, Colson P, Scola BL, Raoult D. 2021. Drug repurposing against SARS-CoV-1, SARS-CoV-2 and MERS-CoV. *Future Microbiol* 16:1341–1370. <https://doi.org/10.2217/fmb-2021-0019>
- Kuo C-J, Chao T-L, Kao H-C, Tsai Y-M, Liu Y-K, Wang L-C, Hsieh M-C, Chang S-Y, Liang P-H. 2021. Kinetic characterization and inhibitor screening for the proteases leading to identification of drugs against SARS-CoV-2. *Antimicrob Agents Chemother* 65:e02577-20. <https://doi.org/10.1128/AAC.02577-20>
- O'Donnell HR, Tummino TA, Bardine C, Craik CS, Shoichet BK. 2021. Colloidal aggregators in biochemical SARS-CoV-2 repurposing screens. *J Med Chem* 64:17530–17539. <https://doi.org/10.1021/acs.jmedchem.1c01547>
- Yan G, Li D, Lin Y, Fu Z, Qi H, Liu X, Zhang J, Si S, Chen Y. 2021. Development of a simple and miniaturized sandwich-like fluorescence polarization assay for rapid screening of SARS-CoV-2 main protease inhibitors. *Cell Biosci* 11:199. <https://doi.org/10.1186/s13578-021-00720-3>
- Zhang J, Yan H, Yan G, Liu X, Wang Y, Chen Y. 2022. Protocol for high-throughput screening of SARS-CoV-2 main protease inhibitors using a robust fluorescence polarization assay. *STAR Protoc* 3:101794. <https://doi.org/10.1016/j.xpro.2022.101794>
- Yan H, Zhang R, Yan G, Liu Z, Liu X, Liu X, Chen Y. 2023. Production of a versatile SARS-CoV-2 main protease biosensor based on a dimerization-dependent red fluorescent protein. *J Med Virol* 95:e28342. <https://doi.org/10.1002/jmv.28342>
- Ma C, Tan H, Choza J, Wang Y, Wang J. 2022. Validation and invalidation of SARS-CoV-2 main protease inhibitors using the Flip-GFP and protease-Glo luciferase assays. *Acta Pharm Sin B* 12:1636–1651. <https://doi.org/10.1016/j.apsb.2021.10.026>
- McGovern SL, Helfand BT, Feng B, Shoichet BK. 2003. A specific mechanism of nonspecific inhibition. *J Med Chem* 46:4265–4272. <https://doi.org/10.1021/jm030266r>
- Zhang R, Yan H, Zhou J, Liu X, Chen Y. 2023. Invalidation of geraniin as a potential inhibitor against SARS-CoV-2 main protease. *Nat Prod Res*:1–4. <https://doi.org/10.1080/14786419.2023.2241973>