

Abstract # AAVGV0025

Evaluation of SARS-CoV-2 3C-like protease inhibitors using selfassembled monolayer desorption ionization mass spectrometry

Zachary A. Gurard-Levin¹, Cheng Liu², Andreas Jekle², Ruchika Jaisinghani², Suping Ren², Koen Vandyck³, Dirk Jochmans⁴, Pieter Leyssen⁴, Johan Neyts⁴, Lawrence M. Blatt², Leonid Beigelman², Julian A. Symons², Pierre Raboisson³, Michael D. Scholle¹, and <u>Jerome D</u>eval²

Background: Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is the cause of the COVID-19 pandemic that began in 2019. The coronavirus 3-chymotrypsin-like cysteine protease (3CLpro) controls replication and is therefore considered a major target for antiviral discovery. Developing robust and sensitive in vitro screening assays is key to rapidly identify the most promising inhibitors of SARS-CoV-2 3CLpro. Methods: SARS-CoV-2 3CLpro inhibitors were evaluated in a novel self-assembled monolayer desorption ionization mass spectrometry (SAMDI-MS) in parallel with a standard FRET enzymatic assay, with and without reducing agent. Antiviral potency of control compounds was also determined in cells infected with OC-43 or SARS-CoV-2. Potential for compound cytotoxicity was conducted in parallel with antiviral assessment ⁽¹⁾.



Buffer optimization of SAMDI-MS assay



A) The initial velocity calculated from triplicate data utilizing the linear range of 3CLpro activity over a range of distinct conditions: NaCl concentration from 0 to 100 nM, buffer (Hepes, Tris, and Bicine) and pH (6.8–8.0), reducing agent (DTT and TCEP at 1 mM), bovine skin gelatin (BSG)—a carrier protein concentration), and detergent Tween-20 and Triton-X 100. (B) K_M values were determined for the SAMDI Velocity of 3CLpro is linear over a range of concentrations in the SAMDI-MS assay using 10 µM substrate.





IC₅₀ measurements of six reported 3CLpro inhibitors were calculated in the SAMDI-MS (red) and FRET (grey) assay formats. Experiments were performed in duplicate and error bars represent standard deviation.

¹SAMDI Tech, Inc., Chicago, USA; ²Aligos Therapeutics, Inc., South San Francisco, USA; ³Aligos Belgium; ⁴Rega Institute, KU Leuven, Belgium; ⁴Rega Institute, KU Leuven, Belgium BV, Leuven, Belgium; ⁴Rega Institute, KU Leuven, Belgium; ⁴Re

Robustness of the SAMDI-MS assay



IC₅₀ measurements of 6 reported 3CLpro inhibitors measured by SAMDI-MS in the absence of reducing agent (grey), in the presence of 1 mM DTT (red), and in the presence of 1 mM glutathione (blue). Experiments were performed in duplicate and error bars represent standard deviation.

Summary of antiviral activity against OC43-CoV and SARS-CoV-2



The label-free SAMDI-MS assay was optimized and validated with known inhibitors of coronavirus 3CLpro such as GC376 (IC_{50} = 0.060 μ M), calpain inhibitors II and XII (IC_{50} ~20-25 μM). The FDA-approved drugs shikonin, disulfiram, and ebselen did not inhibit SARS-CoV-2 **3CLpro** activity in the SAMDI-MS assay under physiologically relevant reducing conditions. The three drugs did not directly inhibit human β-coronavirus OC-43 or SARS-CoV-2 in vitro, but instead induced cell death. Compared with a traditional FRET readout, the SAMDI-MS assay offers greater sensitivity and eliminates false positive inhibition from compound interference with the optical signal. In conclusion, the SAMDI-MS 3CLpro assay, combined with antiviral and cytotoxic assessment, provides a robust platform to evaluate antiviral agents directed against SARS-CoV-2.

References: Gurard-Levin ZA, Liu C, Jekle A, Jaisinghani R, Ren S, Vandyck K, Jochmans D, Leyssen P, Neyts J, Blatt LM, Beigelman L, Symons JA, Raboisson P, Scholle MD, Deval J. Antiviral Res. 2020 Sep 5;182:104924. doi: 10.1016/j.antiviral.2020.104924.

$\mathbf{AII(-(-)S)}$ **FHERAPEUTICS**

OC-43 Hela Cells		OC-43 MRC-5 Cells		SARS-CoV-2 VeroE6 Cells	
EC ₅₀ (μM)	СС ₅₀ (µМ)	EC ₅₀ (μM)	СС ₅₀ (µМ)	EC ₅₀ (μM)	СС ₅₀ (µМ)
0.114±0.009	>1	0.21	≥50	n.d.	n.d.
n.d.	n.d.	n.d.	n.d.	<0.8	55±18
0.42±0.033	>10	0.83	>50	10±4.2	>100
20.7±3.3	>100	20.95	>100	27±1.4	>100
15.2±2.4	60.3±8.3	6.93	>50	1.3±0.57	27±0.0
>100	15.9±2.4	>100	8.23	>100	37.5±9.2
>100	75	>100	8.35	>100	13.5±0.70
>100	1.4±0.001	>100	0.32	>100	1.55±0.07

Results and Conclusions