## **Coronavirus Screening Tools**

Cayman has developed a SARS-CoV-2 Screening Library in which over 70,000 unique drugs and small molecules have been screened *in silico* for binding to COVID-19 targets, Spike Glycoprotein, Peptidases, Non-structural proteins (Nsps), and human ACE2. These are available as both a data pack and a library of the chemical entities in a 96-well format.



Cayman's unique set of compounds

FDA-approved drugs

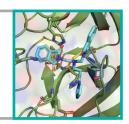
Diverse commercial compounds



# Virtual screening against SARS-CoV-2 targets

#### Hit selection criteria:

- · Docking scores
- · Intermolecular hydrogen bonds within the amino acid residues of the binding pockets





#### Library Data Package

Parameter	HTS02812	JFD03677
GlideScore/Glide Emodel	-9.38/-85.955	-9.60/-98.831
MW	369.463	452.509
Number of H Donors	3	4
Number of H Acceptors	7.4	6
PSA	82.08	97.04
ClogS	-2.18	-5.376
ClogPo/w	2.57	4.867
ClogBB	-0.56	-1.69
ClogHERG	-6.29	-7.64

Docking score units are in kcal/mol.

#### **Library Preparation**



- Can be customized to your selected compounds/targets
- · Supplied in a 96-well Matrix™ format



Hit-to-Lead and Lead Optimization Services

- · In vitro/In vivo Screening
- · SBDD/MedChem

## SARS-CoV-2 Screening Library

Item No. 9003509

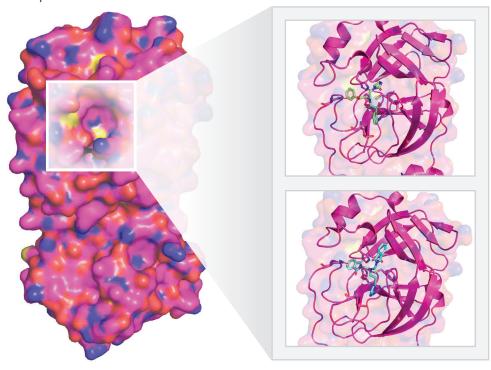
The SARS-CoV-2 Screening Library is a custom, made-to-order library that features a diverse set of FDA-approved and drug-like compounds identified from *in silico* modelling using Maestro (Schrödinger Suite) software. Choose from the entire library of 2,000+ compounds or a custom built library to your preferred target and specifications. The data package for all SARS-CoV-2 targets is also available. SARS-CoV-2 targets include:

- Main protease (3CL<sup>pro</sup>)
- · Spike glycoprotein
- · ACE2 (human)
- · RdRP (Nsp12)
- · Endoribonuclease (Nsp15)
- Guanine-N7 methyltransferase (Nsp14) | ExoN domain (Nsp14)
- PL<sup>pro</sup> (Nsp3)
- · ADP-ribose phosphatase (Nsp3)
- · Nucleocapsid protein

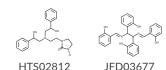
## SARS-CoV-2 Library Data Package

The SARS-CoV-2 Library Data Package includes compound characteristics, docking scores (per target), and predicted physicochemical properties for nine SARS-CoV-2 targets. It is available for purchase, and the cost can be applied towards your next screening, lead optimization, or compound synthesis project with Cayman.

#### Sample Data:



Key Residues in 3CL<sup>pro</sup> SARS-CoV-2: His41 motif, His163-Glu166 motif, and Catalytic Cys145



HTS02812 Parameter JFD03677 GlideScore/ -9.38/ -9.60/ Glide Emodel -85.955 -98.831 MW 369.463 452.509 Number of 4 H Donors Number of 7.4 6 **H** Acceptors PSA 97.04 82.08 ClogS -2.18 -5.376 ClogPo/w 2.57 4.867 ClogBB -0.56 -1.69

Docking score units are in kcal/mol.

-6.29

-7.64

ClogHERG

To learn more about our Medicinal Chemistry & Structure-Based Drug Design services, visit www.caymanchem.com/medchem

3CLpro (PDB ID 6LU7)