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Expanding the chemical space for antiviral discovery: the potential of twistenediones†

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Despite significant progress in drug discovery, there remains an urgent need to identify new structures capable of targeting drug-resistant diseases, as well as novel pathogens, to address the growing challenges in global health. This work highlights the underexplored potential of twistane-like structures as promising candidates for drug development, particularly as antiviral agents. We provide the first comprehensive study of their antiviral activity, in particular against SARS-CoV-2. We report the synthesis of a family of chiral indolyl-twistenediones, with the separation and characterization of both enantiomers *via* chiral semipreparative HPLC. The absolute configurations were determined using experimental and theoretical ECD techniques, supported by DFT calculations. A detailed biological study of their antiviral activity against various pathogenic RNA viruses demonstrates selective efficacy against members of the *Coronaviridae* family, specifically targeting a post-entry step in the viral replication cycle. Further investigation revealed a remarkable chiral distinction in the antiviral activity between the two enantiomers, opening new avenues for research in the 3D space of chiral cage compounds.

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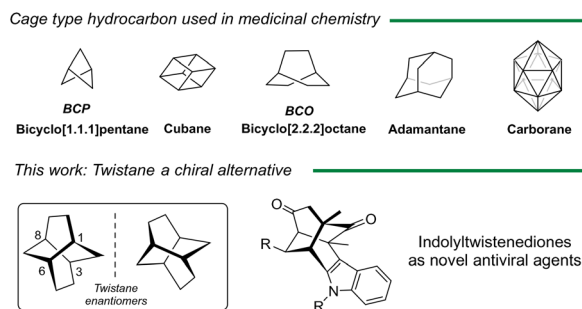‡ These authors contributed equally.

Introduction

The discovery of novel chemical entities is a dynamic field that continually seeks to identify innovative molecules with therapeutic potential. This effort aims to address various medical challenges and improve the effectiveness of treatments. A great deal of interest has been placed on incorporating a three-dimensional (3D) character into screening libraries, as emphasized by the concept of "*Escape from Flatland*".[1] This approach highlights the importance of increasing the fraction of sp³-hybridized carbons in drug discovery, leading to the generation of new saturated molecules with improved medicinal properties.[2] Furthermore, the role of chirality and stereochemistry is vital in drug design, as the 3D arrangement of chiral molecules critically influences their interactions with biological targets. Polycyclic cage-shaped compounds represent a burgeoning area of research and development in medicinal chemistry, materials science, and other interdisciplinary fields. These cage-shaped structures, characterized by their unique geometry and properties, serve as versatile building blocks for creating innovative molecules with diverse applications. Despite the historical focus on certain cage-shaped compounds like adamantane,[3] more recent efforts have expanded the scope to include other derivatives such as bicyclo[1.1.1]pentanes, cubanes or carboranes which have opened the door to unexplored chemical space. Twistane (tricyclo[4.4.0.0³,⁸] decane) is a cage twist-boat isomer of adamantane that exists in two enantiomeric forms due to its inherent chirality (D_2-symmetry) (Fig. 1).[4] In contrast to the extensive attention received by adamantane analogs, twistane derivatives have been the subject of limited research, primarily due to the challenges associated with their synthesis.[5] Twistane was first synthetized by Whitlock *et al.*[6] in 1962 using an intramolecular enolate alkylation as a key step to form the twistane scaffold in a total synthesis of eleven steps. Since then, only a limited number of methodologies,[7–10] including one developed by our group,[11] have described the preparation of these tricyclic cage compounds. Therefore, the development of cage-shape derivatives including chiral derivatives will boost the discover of new biological relevant molecules.[12]

Fig. 1   Main cage hydrocarbon structures as potential drugs and novel indole twistene like structures.

Scheme 1   Synthesis of twistendienones **1** and structures for antiviral evaluation (A) 6/6/6 cage **1a**–**i** and (B) 6/6/9 cage compound **4a,b**.

In 1968, Deslongchamps patented the preparation of tricyclo[4.4.0.0]decan-1-amine and reported its potential activity against certain strains of influenza, in particular influenza A and A′.[13] Despite these notable features, no further biological studies have been reported of such derivatives, nor explore the chiral impact on the biological activity. Inspired by these results, and taking into account the pharmacological versatility of indole, we reasoned that indole twistenedienones (Fig. 1) could behave as novel antiviral agents. Herein, we report the preparation of different functionalized twistenedienones structures and their effectiveness against SARS-CoV-2 virus. Moreover, we described the first enantiopure derivative and report the chiral distinction in the viral activity against major human viral pathogens including SARS-CoV-2 and MERS-CoV (middle-east respiratory syndrome) coronavirus, and non-related RNA viruses like West Nile virus (WNV) or a recombinant vesicular stomatitis virus (VSV).

Results and discussion

Our group reported the direct synthesis of indolyl substituted twistenediones **1** by a domino process involving 2-quinonyl boronic acid **2** and (E)-2-arylalkenylindoles **3**.[11] This 6/6/6 cage skeleton are formed via domino sequence that involves three different transformations: the regioselective 1,4-addition of the indole to the C-3 quinone position, followed by an intramolecular regioselective Diels–Alder cycloaddition (IMDA) and protodeboronation step (Scheme 1). The boronic acid on the benzoquinone proved essential for controlling regioselectivity and enhancing reactivity, while also acting as a traceless directing agent.

The initial evaluation of the antiviral activity was conducted with twistenediones **1a**–**c** (R^1 = Ph, ortho-bromophenyl and para-methoxyphenyl) (Scheme 1), which were prepared from the corresponding N-methyl 2-alkenyl-indole **3**, using DCM as solvent at room temperature.[14]

The antiviral activity was initially evaluated using A549-ACE2 cell monolayers. Viral propagation was monitored by quantitation of the viral antigen in cultures treated with the vehicle (100%) or cultures that were not inoculated with the virus (0%). To assess the antiviral activity of **1a**–**c**, serial
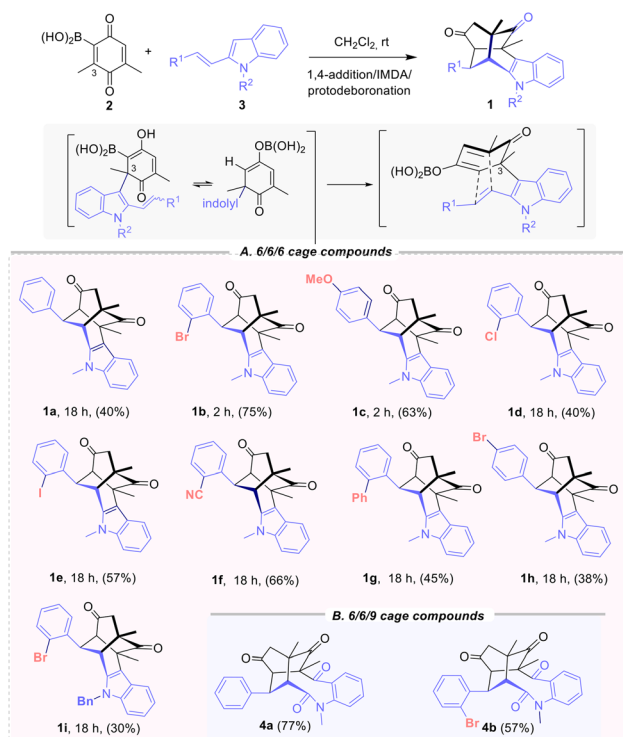
dilutions of the compounds were mixed with infectious SARS-CoV2 before inoculation to the cell monolayers. Table 1 shows potency indexes (EC_{50}, EC_{90} effective concentrations capable of reducing viral propagation by 50% or 90%) of the compounds and their interference with cell viability, as determined by an MTT-based assay (CC_{50} cytotoxic concentration reducing cell viability by 50%). Gratifyingly, compounds **1a**–**c** interfered with viral propagation at concentrations <22 µM for EC50 and <35 µM for EC90 (Table 1, entries 1–3). Moreover, the measurement of the CC_{50}/EC_{50} ratio, which reflects the degree of toxicity versus the level of protection against the virus, displayed some degree of cytotoxicity, resulting in narrow therapeutic windows. Compounds showed substantial antiviral activity at concentrations that reduced viability by more than 20% (CC_{20}/EC_{90} < 1), a cut-off commonly established for the viability assays.

In view of these limitations, compounds were subsequently tested for their ability to protect Vero-E6 cell monolayers from SARS-CoV-2-induced cell death. This is a semi-quantitative phenotypic assay that assists in the identification of candidates with robust antiviral activity.[15] A positive protection result requires that the compound does not substantially interfere with cell viability for a period of 72 hours and that it interferes sufficiently with viral propagation so that the monolayer is intact after the same period of time. Among the tested compounds, only **1b** was capable of completely protecting the VeroE6 monolayer in a relatively wide range of concentrations, from 50 to 6.25 µM. These

Table 1 Evaluation of the antiviral activity of the chiral racemic **1a–i**, **4a,b**[a]

| Entry	1	EC_{50}	EC_{90}	CC_{50}	CC_{20}	CC_{50}/EC_{50}	CC_{20}/EC_{90}	VERO[b]
1	**1a**	13.4	34.8	25.7	12.7	1.9	0.4	No
2	**1b**	15.5	35.0	70.4	32.2	4.5	0.9	**Yes**[c]
3	**1c**	21.0	30.3	17.7	5.1	0.8	0.2	No
4	**1d**	13.0	35.2	62.3	20.7	4.8	0.6	No
5	**1e**	10.6	23.4	61.7	10.6	5.8	0.5	No
6	**1f**	18.8	36.0	100	61.7	5.3	1.7	No
7	**1g**	22.0	100	76.7	26.7	3.5	0.3	No
8	**1h**	8.7	48.3	77.6	56.6	8.9	1.2	No
9	**1i**	19.0	66.5	70.7	17.1	3.7	0.3	No
10	**4a**	43.7	89.0	24.8	89.0	2.3	1.0	No
11	**4b**	>100	>100	100	3.7	—	—	No

[a] Antiviral efficacy (EC_{50} and EC_{90}; μM) and cytotoxicity (CC_{50} and CC_{20}; μM) indexes were determined in A549-ACE2 cells and are shown as mean and the SD of a minimum of three biological replicates, see ESI† for details. [b] Cell monolayer integrity experiments were carried out in Vero-E6 cell and are expressed qualitatively. [c] Range of concentration studied 0.8–50 μM, protective at doses ranging from 12.5 to 50 μM.

preliminary results indicated that **1b** exhibits antiviral activity against SARS-CoV-2 both in human and African green monkey cells.

A comparison of derivatives **1a–c** revealed that *ortho* substitution has a positive influence on antiviral activity. Subsequently, various *ortho*-phenyl substituted 2-(aryl) vinylindoles were synthetized to prepare the corresponding twistenediones. Thus, derivatives with pedant *ortho*-chloro (**1d**), iodo (**1e**), cyano (**1f**) or phenyl (**1g**) substituents were prepared (Scheme 1). Moreover, to assess the impact of the bromo substituent at the *para* position, derivative (**1h**) was also synthetized. Additionally, the *N*-benzyl *ortho*-bromophenyl derivative **1i** was also prepared. The new 6/6/6 cage-derivatives **1d–i** and expanded 6/6/9 tricyclic derivative **4a–b** were tested for their ability to interfere with viral propagation of SARS-CoV-2 in A549-ACE2 cells (Table 1, entries 4–11). All twistenedienones **1d–i** interfered at concentrations <22 μM (Table 1, EC50, EC90). The *o*-cyano **1f** and the *p*-bromo **1h** derivatives presented the best values (Table 1, entries 6 and 8 respectively). Most compounds showed antiviral activity at concentrations that reduced viability by more than 20% (CC_{20}/EC_{90} < 1). To assess the impact of the 6/6/6 cage skeleton, expanded 6/6/9 tricyclic derivative **4a** and *ortho*-bromo **4b**, lacking the twistene framework, were also evaluated. The ketolactams derivatives **4a–b**[14] were prepared through the oxidative cleavage of the C-2–C-3 indole bond (Witkop type oxidation)[16] promoted by MCPBA and catalytic *p*-TsOH. Both 6/6/9 tricyclic cage derivatives **4a** and **4b** showed a complete lack of reactivity, pointing out the importance of the twistene framework. Further evaluation of the Vero-E6 cell monolayers protection from SARS-CoV-2-induced cell death, revealed that exclusively the *ortho*-bromo derivative **1b** provided protection with a sufficient therapeutic window (Table 1, entry 2). Therefore, compound **1b** was selected for further biological characterization.

Next, we investigated the relationship between the configuration of the twistenedione **1b** and the antiviral activity. Both e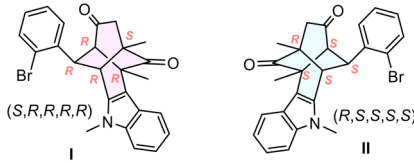nantiomers of **1b** (namely **I** and **II**) were separated from the racemic mixture by semipreparative scale chiral stationary phase HPLC (CSP-HPLC) (Fig. 2a). The absolute configuration was established by comparison of the experimental ECD spectra of pure enantiomers with the simulated ECD spectra. The experimental ECD spectra exhibits two main absorption bands centered at 275 and 320 nm with opposite sign for each enantiomer (Fig. 2c, black and blue solid lines). The geometry of the enantiomer (*S,R,R,R,R*) was optimized by DFT calculations (Fig. 2b), and the corresponding UV-vis and ECD spectra were simulated with the aid of TD-DFT (Fig. 2c, red dash line),
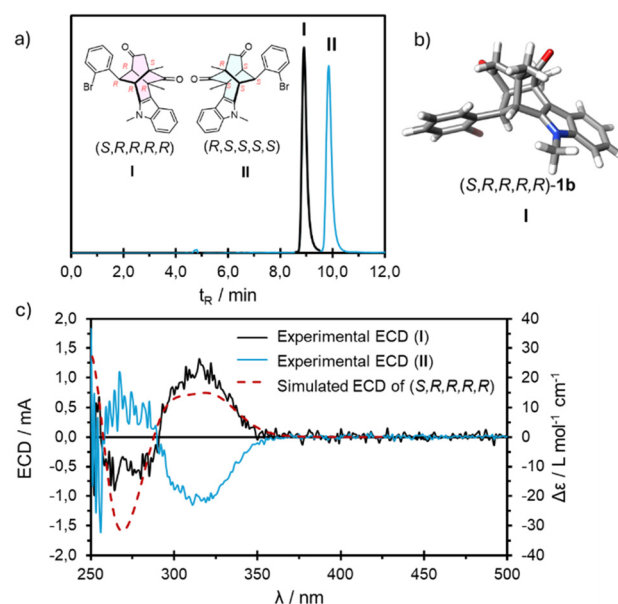
Fig. 2 a) CSP-HPLC separation chromatogram of chiral racemic **1b** using a Chiralpak-IA column, hexane/EtOAc (7:3), 3.4 mL min⁻¹, retention time I = 8.90 min and II = 9.85 min and assignment of absolute configuration; b) optimized structure of (*R,R,R,R,R*)-**1b** (CAM-B3LYP/6-31G++(d,p)); c) simulated ECD spectrum of (*S,R,R,R,R*) (CAM-B3LYP/6-31G++(d,p)/PCM, dashed red line) and experimental ECD spectra of both enantiomers of I and II in DCM at *ca*. 10⁻⁵ M (black and blue lines).

Table 2 Antiviral activity of chiral racemic **1b** and enantiopure **I** and **II**[a]

(S,R,R,R,R) **I** (R,S,S,S,S) **II**

Entry	1	EC_{50}	EC_{90}	CC_{50}	EC_{20}	CC_{50}/EC_{50}	CC_{20}/EC_{90}	VERO[b]
1	**1b**[c]	15.5	35.0	70.4	32.2	4.5	0.9	Yes[d]
2	**I**	5.8	8.6	100	36	17.2	4.2	**Yes**[e]
3	**II**	28.9	47.0	100	71.3	3.5	1.5	No

[a] Antiviral efficacy (EC_{50} and EC_{90}; μM) and cytotoxicity (CC_{50} and CC_{20}; μM) indexes were determined in A549-ACE2 cells and are shown as mean; SD of a minimum of three biological replicates, see ESI† for details. [b] Cell monolayer integrity experiments were carried out in Vero-E6 cell and are expressed qualitatively. Range of concentration studied. [c] Racemic mixture. [d] 50–12.5 μM. [e] 25–6.25 μM.

matching the experimental ECD of **I** enantiomer, thereby proving the (S,R,R,R,R) configuration to **I** and (R,S,S,S,S) to **II**.

Enantiomers **I** and **II** were then subjected to the same biological characterization using A549-ACE2 cell monolayers, and the results indicate that antiviral activity segregates with **I**, improving all EC_{50} and EC_{90} values, as well as the therapeutic window (CC_{20}/EC_{90}) compared to **II** (Table 2, entries 2–3).

Moreover, only the enantiomer **I** provided protection of the Vero-E6 cell monolayers from SARS-CoV-2-induced cell death, at reasonable lower concentrations of the racemic sample, demonstrating a clear selectivity of the virus for this enantiomer (Table 2, VERO column). This phenomenon was verified in three different cell lines (Fig. 3) (Vero-E6, A549-ACE2 and human lung epithelial cell line Calu 3). Both compounds also show some degree of associated cytotoxicity with a measurable impact on cell viability (CC_{20} = 36 ± 10 μM for **I** and 71 ± 8 μM for **II**). All subsequent characterization experiments were performed below 30 μM to ensure a minimal impact on cell viability of either compound.

To determine if the antiviral activity of both enantiomers **1b** was restricted to SARS-CoV-2, cells were infected with a pathogenic middle-east respiratory syndrome coronavirus (MERS-CoV) in the presence of increasing doses of **I** and **II** (Fig. 4). Moreover, both enantiomers were also tested against non-related RNA viruses like West Nile virus (WNV) or a

recombinant vesicular stomatitis virus (VSV), bearing a reporter green fluorescent protein (GFP) (Fig. 4). Viral propagation efficiency was determined by measuring the relative viral RNA load at 48 hours post-inoculation in cells infected with SARS-CoV-2, MERS-CoV or WNV (MOI – multiplicity of infection experiment – 0.01) by real-time quantitative PCR. VSV–GFP propagation efficiency was determined by the relative GFP expression at 16 hours post-inoculation (MOI 0.01) using automated fluorescence microscopy. The results indicated that **I** caused more than 3 log viral load reduction in SARS-CoV-2 and MERS-infected cells, while the same concentration only marginally interfered with WNV or VSV replication (Fig. 4). On the contrary, **II** did not significantly interfere with infection by any of these viruses, confirming that antiviral activity segregates with **I**. These results together with the mock control experiments and solvent (DMSO) assays and experiments suggest a selective antiviral mechanism and strongly reduces the concern that antiviral activity is solely due to compound cytotoxicity.

Finally, we set out to determine if both enantiomers interfered with viral *entry* or with a *post-entry* event of coronaviruses. To this aim, we performed a *time-of-addition* relative to human coronavirus hCoV-229E inoculation and determine the relative efficiency of infection in the presence of antiviral doses of **I** or **II**. For these experiments, Huh7 cells
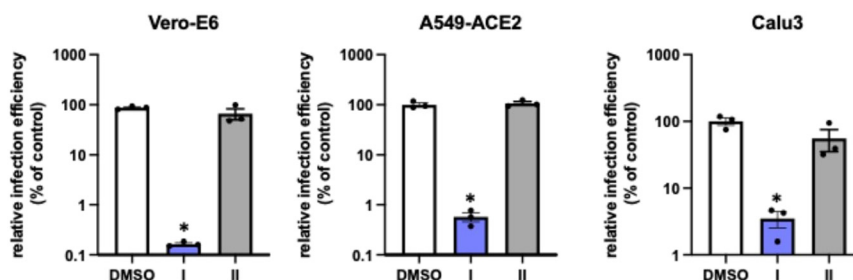
Fig. 3 Stereoselective evaluation of enantiopure **I** and **II** in different cell lines. Target cells were inoculated with SARS-CoV-2 in multiple cycle infection experiments in the presence of **I** (blue) or **II** (grey) at 10 μM, using solvent (DMSO, white) as a control. Relative infection efficiency was determined as described in the Experimental section (ESI†). Data is shown as mean and SD of three biological replicates. Statistical significance was determined by one-way ANOVA and a Dunnett's *post hoc* analysis (*$p < 0.05$).
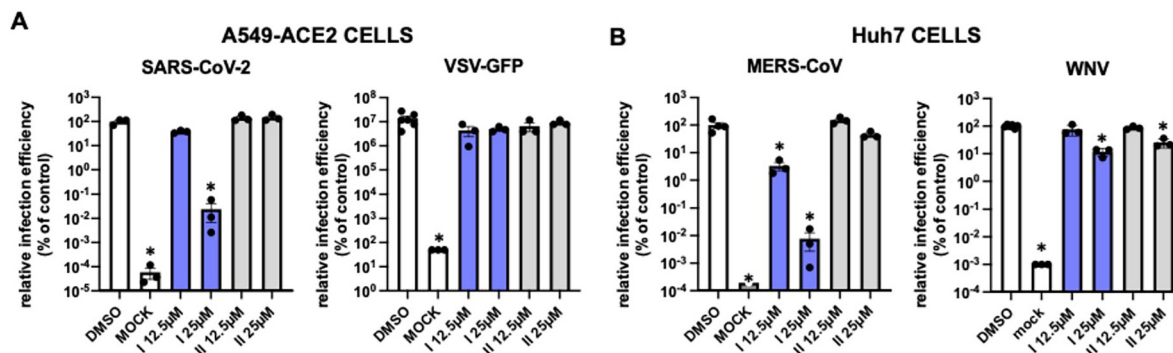
Fig. 4 Evaluation of the selectivity of chiral enantiomers I and II for coronaviruses. Target cells were inoculated with different viruses in multiple cycle infection experiments in the presence of the candidate compounds (white, controls; blue I; grey, II). Relative infection efficiency was determined as described in the Experimental section (ESI†). A) SARS-CoV-2 (MOI 0.001) or VSV-GFP (MOI 0.01) were used to inoculate A549-ACE2 cells. B) MERS-CoV (MOI 0.01) or WNV (MOI 0.01) were used to inoculate Huh7 cells. Data is shown as mean and SD of three biological replicates. Statistical significance was determined by one-way ANOVA and a Dunnett's *post hoc* analysis (*$p < 0.05$).

(human hepatoma cells expressing ACE2) were used, as they are susceptible to a recombinant hCoV-229E expressing GFP as reporter gene.[17] Three different evaluations were run: i) I or II present at all times (Fig. 5, *ALL*), ii) I or II added only during virus inoculation (Fig. 5, *ENTRY*), and iii) I or II added only after the virions had effectively penetrated the cells in single-cycle infection experiments (Fig. 5, *POST-ENTRY*). For the *ENTRY* and *ALL* experiments, compounds were mixed with sufficient virions to achieve MOI 2 and produce a single cycle infection experiment. After 1 h at 37 °C, cells were washed and replenished with media devoid of antivirals (Fig. 5, *ENTRY*) or left with the media until the end of the

experiment (Fig. 5, *ALL*). To determine if the compounds interfered with infection *POST-ENTRY*, target cells were inoculated with the virions (MOI 2) for 1 h at 37 °C in the absence of compounds. Then, the inoculum was removed, cells were washed, and the inoculated cultures were subsequently treated with the compounds I and II.

The relative infection efficiency was measured 36 hours post-inoculation using automated fluorescence microscopy. 2mAde (2′-*c*-methyladenosine) (Boc Sciences) was used as a positive post-entry replication inhibitor in the experiment and DMSO as a negative inhibition control. Enantiomers I or II did not interfered with infection when added during viral *ENTRY*
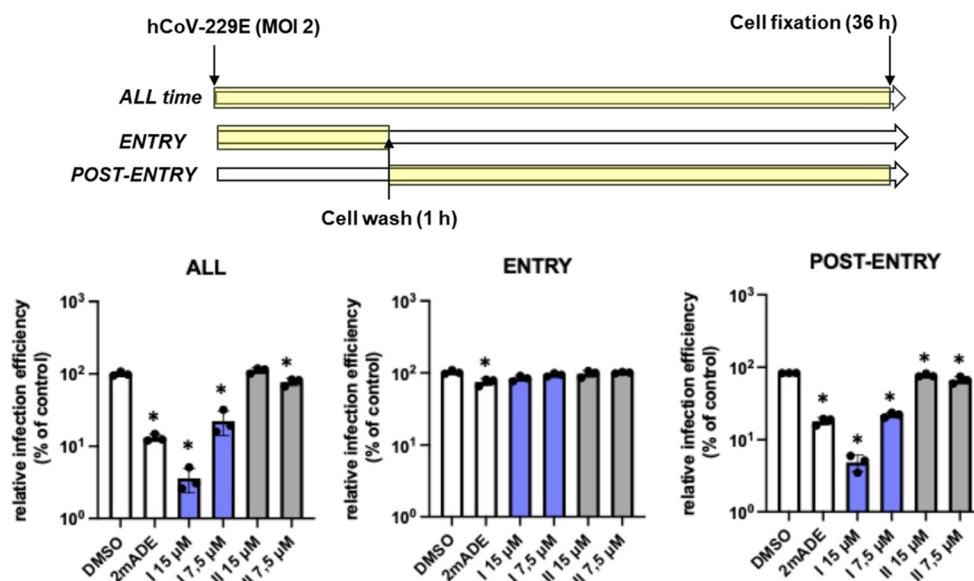
Fig. 5 Time-of-addition experiments: schematic diagram of the times where enantiomers I or II were present in the assay (yellow fill arrows) Huh7 cells were inoculated at MOI 2 with a recombinant human alphacoronavirus expressing GFP (hCoV-229E-GFP) in the presence or absence of the compounds. After 1 hour at 37 °C, media were either left until the end of the experiment (*ALL*) or removed and replenished with normal media (*ENTRY*). A set of untreated samples was treated after the inoculation period (*POST-ENTRY*) until the end of the experiment. 2mAde (25 µM) (Boc Sciences) was used as a positive post-entry replication inhibitor in the experiment. Samples were fixed 36 hours post-inoculation and infection efficiency was determined by quantitative fluorescence microscopy. Data is shown as mean and SD of three biological replicates. Statistical significance was determined by one-way ANOVA and a Dunnett's *post hoc* analysis (*$p < 0.05$).

experiment (Fig. 5, *ENTRY*). Interestingly, enantiomer **I** demonstrated full antiviral activity at a *POST-ENTRY* step (Fig. 5, *POST-ENTRY*), with antiviral activity comparable to that observed when the compound was kept for the entire duration of the experiment (Fig. 5, *ALL*). These experiments validate the antiviral activity of (S,R,R,R,R)-**1b** (**I**) while demonstrating the absence of activity for **II** (Fig. 5, *ALL*), further extending the breadth of the anti-coronavirus spectrum.

Conclusions

This study has brought to light the potential of indolyl twistenedione cage compounds, both in their racemic form and as enantiopure chiral entities, as novel drug candidates, particularly as antiviral agents. In particular, a series of indolyl twistenediones has been prepared and their activity against SARS-CoV-2 has been evaluated for the first time. The results highlighted the potential of the *ortho*-bromo phenyl derivative **1b** providing the best results as the antiviral candidate. Both enantiomers of twistenedione **1b** were separated from the racemic mixture by semipreparative chiral HPLC (CSP-HPLC) and the absolute configuration was established by comparison of the experimental ECD spectra of pure enantiomers with the simulated ECD spectra. The antiviral activity of a chiral enantiopure 6/6/6 cage derivative has been evaluated for the first time, showing an outstanding chiral differentiation using the (S,R,R,R,R)-enantiomer (**I**), while demonstrating the absence of activity for (R,S,S,S,S)-**II** enantiomer. The efficiency of both enantiomers was tested in three different cell lines (A549-ACE2, Vero-E6 and Calu3), and with three different coronavirus (SARS-CoV-2, MERS-CoV or hCoV-229-GFP) and two non-related RNA viruses (West Nile virus or a recombinant vesicular stomatitis virus). Our results indicate that enantiomer (S,R,R,R,R)-(**I**) showed a remarkable antiviral activity in all the cell lines evaluated. In particular, enantiomer **I** caused more than 3 log viral load reduction in SARS-CoV-2 and MERS-infected cells, while the same concentration slightly interfered with WNV or VSV replication. Regarding the interaction with the virus replication cycle, the time-of-addition experiments showed a selective post-entry event of enantiomer **I** with the virus. Our results highlight the potential use of chiral enantiopure twistene-like compounds, with unique geometries, for antiviral therapy, demonstrating its selectivity against coronaviruses. One of the main limitations of the study is the current therapeutic window (CC50/EC90), of the lead compound. The differential efficacy of the compounds against different viruses is probably due to their differential replication strategies and host cell requirements. We ruled out that the host cell line contributes to the differential efficacy by testing VSV and SARS-CoV2 in A549-ACE2 cells and MERS and WNV in Huh7 cells, further supporting the notion for a selective antiviral activity and a family-specific mode-of-action. Thus, further endeavours should aim at preparing derivatives with larger therapeutic window, which will constitute valuable tools to determine the antiviral spectrum quantitatively. Additionally, such compounds could also be used for genetic resistance profiling as well as other mode of action and target deconvolution approaches. This study paves the way for further applications of enantiopure cage-shape compounds in antiviral research and drug development.

Data availability

Cell culture, viruses, evaluation of the antiviral activity, cytotoxicity, viral quantitation, general synthetic procedures and analytical data including copies of ^{1}H and ^{13}C-NMR spectra are provided in the ESI.†

Author contributions

The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript. Conceptualization was done by U. G., P. G., J. A. and M. R.; methodology and characterization was performed by A. J., D. G.-G. and A. P.; separation of enantiomers, photophysical characterization and theoretical computation was performed by A. M. O., C. M. C., J. M. C.; biological evaluations was done by P. B. V. C. U. G. P. G. Manuscript preparation was done by J. A., P. G. and M. R.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

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