Novel vascular disrupting agents with a cyclohexanedione scaffold identified through a ligand-based virtual screening approach

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TITLE RUNNING HEAD: Vascular disrupting agents based on cyclohexanediones

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Abbreviations: VDA: vascular-disrupting agent; VS: virtual screening; ROCS: rapid overlay of chemical structures; MTC: 2-methoxy-5-(2,3,4-trimethoxyphenyl)-2,4,6-cycloheptatrien-1-one; R-PT: (R)-(+)ethyl 5-amino 2-methyl-1,2-dihydro-3-phenylpyrido[3,4-b]pyrazin-7-yl carbamate.
ABSTRACT.

Vascular Disrupting Agents (VDAs) constitute an innovative anticancer therapy that targets the tumor endothelium, inducing crucial morphological and functional changes that trigger a rapid and dramatic decrease in blood flow leading to tumor necrosis. Our approach for the identification of new VDAs has relied on a ligand 3-D shape similarity virtual screening (VS) approach using as query agents that bind the α,β−tubulin dimer at the colchicine-binding site. The VS campaign, followed by biological testing, afforded an interesting hit with a cyclohexane-1,3-dione structure. This was followed by a synthetic programme of structural analogues. Interestingly, several of these new analogues showed anti-proliferative activity in the sub-µM range, being 100-fold more potent than the initial hit. Additional biological assays showed that they arrest the cell cycle at the G2/M phase and bind at the colchicine-binding site in tubulin with binding affinities in the low-µM range. Moreover, these compounds cause vascular disruption as shown by their ability to destroy an established endothelial tubular network. In conclusion, our approach has led to a new family of promising anti-proliferative compounds with antimitotic and VDA properties.

KEYWORDS: virtual screening; vascular-disrupting agents; colchicine; cyclohexanediones
Introduction

The growth of solid tumors and development of metastasis are highly dependent on the existence of a vascular network that provides oxygen and nutrients. Therefore anti-vascular strategies are increasingly gaining importance among anti-neoplastic therapies and represent a new frontier in the treatment of cancer. Basically, there are two major approaches in anti-vascular therapies: anti-angiogenic drugs, which inhibit new blood vessel formation, and vascular-disrupting agents or VDAs, which affect the existing vasculature. While anti-angiogenic drugs are mostly cytostatic to the endothelial cells and need to be chronically administered, VDAs are typically cytotoxic and cause a quick and dramatic collapse in the blood flow that leads to ischemia and necrosis just 24 h after administration. Thus, they are particularly effective in advanced stages when large tumors are formed. Importantly, one of the main advantages of such anti-vascular approaches is that they target endothelial cells, a cell population less prone to mutations than tumor cells.

The selectivity of VDAs for tumor endothelium versus physiological vessels lies in the crucial differences between them. In fact, tumor vessels are characterized by a higher proliferation rate, absence of pericytes, basal membrane deficiencies and increased vascular permeability. Moreover, tumor vessels are more fragile and tortuous which results in a higher resistance to blood flow. This makes them more sensitive to any decrease in perfusion pressure.

Besides biotechnological approaches, involving immunotoxins and targeted antibodies, there is an increasing interest in low-molecular-weight molecules that behave as VDAs, and some of them are in clinical trials for a variety of solid tumors. The best studied VDAs are microtubule-destabilizing agents that bind the α,β-tubulin dimer at the colchicine-binding site. The most representative compounds include colchicine (1), ZD6126 (2), ABT-751 (3), combrestatins and their prodrugs, CA1P (4), CA4P (5) and AVE-8062 (6), as shown in Figure 1. These compounds inhibit tubulin polymerization in endothelial cells affecting the cellular cytoskeleton and therefore induce changes not only in the shape of endothelial cells but also in their motility, invasion, attachment and proliferation. As a result of a large cascade of events, such as actin stress fiber contraction and the subsequent activation of Rho kinases, VDAs lead to vascular collapse, tumor hypoxia and hemorrhagic necrosis.

It is interesting to mention that colchicine-like VDAs are able to behave as anti-mitotic agents at concentrations slightly higher than those at which the anti-vascular effects are observed. This anti-mitotic effect, closely related to the importance of the cytoskeleton of α,β-tubulin in mitotic spindle formation, when
affecting tumor cells, may induce apoptosis. Therefore, this dual mechanism of action affecting endothelial and tumor cells makes these drugs very promising for anti-cancer therapy.\textsuperscript{14}

Despite the great potential of these drugs, they suffer some drawbacks. In particular, colchicine is too toxic to be considered a suitable anti-cancer drug\textsuperscript{15} and CA-4P presents chemical instability leading to the inactive trans-isomer and shows a short biological half-life.\textsuperscript{16}

Therefore our objective has been to identify new chemical entities able to bind at the colchicine-binding site in α,β-tubulin that may lead to a novel type of compounds with anti-proliferative and anti-vascular properties. To address this aim, we relied on a ligand-based virtual screening (VS) approach.\textsuperscript{17} Such approach represents a quick, efficient and well-documented strategy to access novel active molecules for a particular target. Among the computational methods available for this strategy, we have focused on 3-D shape similarity, a strategy successfully applied on recently reported examples.\textsuperscript{18} A significant advantage of this approach is that no specification of the chemical structure (i.e. types of atoms and/or their bond arrangements) is required, thus chemical scaffolds significantly different from the template can be identified.

Here we report the results obtained after conducting a ligand-based VS campaign using two different queries, identification of the hits and further exploration in terms of improvement of their anti-proliferative properties and determination of the mechanism of action.

**Results and Discussion**

**Virtual screening**

The Virtual Screening protocol employed is summarized in Figure 2 and all details are described in the Experimental Section.

The compounds for the virtual screening were obtained from the freely available ZINC database,\textsuperscript{19} (version 8, which comprises about 8.5 million molecules) in SMILES format.\textsuperscript{20} Prior to VS, the database was filtered on the basis of a customized “drug-like filter” using the software FILTER,\textsuperscript{21} to remove, among others, compounds with reactive functional groups. The filtered database was subjected to OMEGA\textsuperscript{22} to generate the 3D atomic coordinates and conformational ensembles of all the compounds. As tool for the VS, we have used ROCS,\textsuperscript{23} a shape comparison program based on the concept that molecules have a similar shape if their volumes, represented by a Gaussian function, overlay well.\textsuperscript{24,25} Besides shape comparison, an electrostatic complementary term called “color”, which represents chemical similarity, was also applied.
As query against which the comparison was performed, the 3-D structure of two well-known tubulin binders at the colchicine site was used: a colchicine derivative (DAMA-colchicine, \(7\))\(^{26}\) and TN-16 \(8\))\(^{27}\) (Figure 3A). A superposition of their experimental cocrystal structures with \(\alpha,\beta\)-tubulin is shown in Figure 3B. The X-Ray of the complex tubulin-TN-16, reported in 2009, clearly shows that this compound is more deeply buried in the \(\beta\)-monomer than DAMA-colchicine, so that it explores a pocket partially different to that of colchicine.\(^{27}\)

The VS protocol included shape comparison and scoring, “color” comparison and scoring (shape plus color) and finally visual inspection. Based on previously reported examples,\(^{25}\) we established a Tanimoto score of 0.75 for the shape comparison, and a “Combo-score” of 1.4 for the sum of shape and color comparison.

When using DAMA-colchicine as query, the highest hits in the scoring list showed structures that closely resembled colchicine or other described tubulin inhibitors, thus no novel scaffolds were identified. Alternatively, when TN-16 was used as template, the scoring list revealed nine clusters of compounds that had not been previously described in the literature as tubulin binders. Based on compound availability, a total of six representative examples of these clusters (see Table S1 in the Supporting Information) were purchased and evaluated for their anti-proliferative activity against one endothelial cell line (MBEC) and one tumor cell line (L1210). From the compounds tested, only the cyclohexanedione \(9\) (Figure 4) showed significant inhibition of cell proliferation in both cell lines with IC\(_{50}\) values of 13 ± 5 \(\mu\)M (Table S2 in the Supporting Information). Evaluation of \(9\) against other four cell lines indicated similar IC\(_{50}\) values. In addition binding of the hit compound \(9\) to tubulin at the colchicine-site was confirmed by the displacement of 2-methoxy-5-(2,3,4-trimethoxyphenyl)-2,4,6-cycloheptatrien-1-one (MTC, a reversible colchicine-binding site ligand) as determined by fluorescence spectroscopy (see Figure S1 in the Supporting Information). Moreover, and from a chemical point of view, compound \(9\) was considered a suitable hit based on its low molecular weight (335), and synthetic accessibility as described in the next section. Therefore a synthetic-driven medicinal chemistry program was set up to evaluate structural analogues of \(9\).

**Chemistry**

Since the hit compound \(9\) had been acquired from a commercial source, the first task was to devise a synthetic strategy that could allow the exploration of the different fragments of the molecule (Figure 4), in
particular the aromatic rings A and D, and the alkyl chain C, while keeping the cyclohexanedione B as a common structural motif. This strategy consisted in the reaction of the 5-substituted cyclohexane-1,3-diones (representing fragments A and B) with acyl chlorides to obtain the C-acyl derivatives followed by Schiff base formation with anilines (fragment D). Since fragment D was the last structural element incorporated in this synthetic strategy, the first series of modifications were performed on this fragment.

Thus, reaction of the commercially available 5-phenylcyclohexane-1,3-dione (10) with propionyl chloride in dichloromethane in the presence of DMAP and DIPEA at 70 °C (Scheme 1) as described for similar analogues\textsuperscript{28} afforded the O-acyl derivative 11 in 64% yield, but not the C-acyl derivative 12 as expected. Interestingly, similar O-acyl derivatives have been transformed into the corresponding C-isomers by reaction with K$_2$CO$_3$ in acetonitrile in the presence of 1,2,4-triazole, by varying the temperature between -10 and 100 °C, or by phase-transfer catalysis.\textsuperscript{29} When compound 11 was treated with K$_2$CO$_3$ in acetonitrile in the presence of 1,2,4-triazole at 30 °C, only 30% of the C-analogue 12 was obtained after 48 h of reaction. However, with this information, we assayed new reaction conditions meant to perform these two steps in a “one-pot” procedure, and making use of microwave-assisted irradiation to reduce the reaction times. Thus, reaction of 10 with propionyl chloride in CH$_3$CN under microwave conditions in the presence of K$_2$CO$_3$, 1,2,4-triazole and Bu$_4$NBr as a phase transfer catalyst at 70 °C afforded the corresponding C-propionyl derivative 12 in 51% yield. It should be mentioned that the combination of phase transfer catalysis and microwave irradiation afforded the desired C-acyl derivative in just a couple of hours, and to the best of our knowledge, this methodology has never been applied to this type of reaction.

Then reaction of 12 with anilines carrying OH or OMe groups in toluene at 150 °C for 2 h under microwave radiation in the presence of 4Å molecular sieves yielded compounds 14a-j. The structures of 14a-j were determined by $^1$H NMR, $^{13}$C NMR, HMBC and HSQC correlation experiments. It should be emphasized that, in the $^1$H NMR spectra, a broad singlet appeared around 15 ppm corresponding to the NH group. This chemical shift suggested the participation of the NH in a potential hydrogen bond with one of the carbonyl groups of the 1,3-cyclohexanedione. To further explore this issue, the $^1$H NMR spectra of compound 14c was recorded in DMSO-$d_6$ and CDCl$_3$, and in both cases the NH signal remained almost unaffected ($\delta$ 14.7 ppm). Additionally, a ten-fold dilution of the sample with DMSO-$d_6$ did not affect the chemical shift of this signal, strongly suggesting the existence of an intramolecular hydrogen bond.
When tested for anti-proliferative activity (see Table 1, Biological Evaluation section), compounds 14c and 14j afforded 2- to 30-fold lower IC<sub>50</sub> values than the initial hit 9. Therefore, we decided to explore the length of the alkyl chain in fragment C while keeping an OMe or Me in orto at the D aromatic ring (Scheme 2). Thus, reaction of the cyclohexanedione 10 with butyryl or acetyl chloride in the presence of K<sub>2</sub>CO<sub>3</sub>, 1,2,4-triazole, Bu<sub>4</sub>NBr in acetonitrile under microwave conditions afforded the C-acylderivatives 15a and 15b, respectively. Further reaction of these acyl derivatives with o-toluidine or o-anisidine in toluene afforded compounds 16a-d in excellent yields.

As will be later discussed (Biological Evaluation section), those compounds with a methyl at fragment C afforded the best inhibitory values. With this information, and keeping a methyl as side chain in C, a wider battery of anilines was introduced at fragment D with at least one substituent at position 2 of the aromatic ring. Therefore, as shown in Scheme 3, reaction of 2-acetyl-5-phenylcyclohexane-1,3-dione (15b) with commercially available 2-chloro (17a), 2-fluoro (17b), 2-trifluoromethyl (17c), 2,3-difluoro (17d), 2,6-difluoro (17e), 2,5-dimethoxy (17f) and 2,6-dimethoxy (17g) anilines afforded compounds 18a-g.

The next series of modifications affected fragment A by replacing the phenyl ring by other groups, in particular, a cyclohexyl, a benzyl or a gem-dimethyl (Scheme 4). The cyclohexane-1,3-diones with a cyclohexyl or a benzyl at position 5 (21a and 21b) were synthesized through the α,β-unsaturated ketone precursors (20a and 20b) that were obtained following two different approaches. The synthesis of 20a was undertaken by condensation between cyclohexane carbaldehyde (19a) and acetone under basic conditions, while the benzyl derivative 20b was obtained by a Witting reaction of phenylacetaldehyde (19b) with 1-(triphenylphosphoranylidene)-2-propenone in chloroform following described procedures. Then 20a and 20b were allowed to react with diethylmalonate in the presence of sodium ethoxide in ethanol at 60 °C to afford the cyclohexane-1,3-diones 21a and 21b, respectively. C-acylation of 21a and 21b with acetyl chloride, as described in previous examples, and further reaction with o-anisidine afforded 23a and 23b in excellent yields. Similarly, reaction of the commercially available gem-dimethyl derivative 22c with o-anisidine in toluene at reflux afforded 23c. It should be mentioned that these compounds were poorly active, pointing to the importance of the aromatic ring directly attached to the cyclohexanedione for the anti-proliferative activity.

Thus, keeping a phenyl group at A, a variety of substituents were introduced at this ring. The 5-phenylcyclohexane-1,3-diones 24a-h (Scheme 5) were synthesized from the corresponding substituted benzaldehydes following described procedures, in a similar way to that described in Scheme 4. C-
acylation of 24a-h with acetyl chloride and further reaction with o-anisidine or o-toluidine provided compounds 26a-k. Finally, demethoxylation of 26k with BBr₃ in CH₂Cl₂ at room temperature afforded the corresponding hydroxyl derivative 26l.

**Biological Evaluation**

**Anti-proliferative activity**

The synthesized compounds were evaluated for their anti-proliferative activity in six different cell lines: three endothelial cell lines [human microvascular endothelial cells (HMEC-1), mouse brain endothelial cells (MBEC) and bovine aortic endothelial cells (BAEC)] and three cancer cell lines [mouse lymphocytic leukemia (L1210), human lymphoblastic leukemia (CEM) and human cervical carcinoma (HeLa) cells].

Data are expressed as IC₅₀ (50% inhibitory concentration) defined as the concentration at which the compounds reduce cell proliferation by 50% and are shown in Table 1. As reference compound we included colchicine (1), which inhibited proliferation of the different cell lines with IC₅₀ values ranging from 0.0038 to 0.031 µM.

Those compounds with an OH at position orto, meta or para in fragment D (9, 14a or 14b) were moderately active in the different cell lines with IC₅₀ values ranging from 8 to 36 µM in endothelial cells, and from 11 to 79 µM in the tumor cell lines. However, compounds with an OMe at these positions showed a more diverse behavior in terms of anti-proliferative activity. Thus, the p-OMe compound (14e) was marginally active in some cell lines (IC₅₀ ≥ 57 µM), the m-OMe derivative (14d) showed moderate activity (IC₅₀ of 16-66 µM) while the o-OMe isomer (14c) afforded IC₅₀ values of 1.2-4.3 µM, that is, 10- to 30-fold better than the hit compound 9 in endothelial cells. The double or triple substituted compounds (14f, 14g or 14h) were inactive or poorly active. The unsubstituted compound (14i) showed moderate anti-proliferative activity while the o-CH₃ derivative (14j) was as active as the corresponding o-OMe compound 14c.

The length of the alkyl chain in fragment C was varied while keeping an OMe or Me in orto at the D aromatic ring. The resulting propyl derivatives 16a and 16b showed moderate anti-proliferative activity with IC₅₀ values at least 13-fold higher than those of the corresponding ethyl derivatives 14c and 14j. Interestingly, the methyl derivatives 16c and 16d were significantly more active than the corresponding ethyl derivatives, with IC₅₀ values as low as 0.09 µM and 0.18 µM for compound 16c in endothelial and tumor cells, respectively. Depending on the cell line, this represents a 60- (HeLa cells) to 400-fold (BAEC) increase in the
anti-proliferative effect compared with the initial hit. Thus a methyl was selected as the substituent of choice in fragment C.

Keeping a methyl group as the alkyl chain at fragment C, we further explored the orto position in fragment D incorporating different electron withdrawing groups. Thus, introduction of a chlorine (18a) afforded IC\textsubscript{50} values in the sub-\textmu M range (0.3-0.7 \textmu M), whereas a fluorine (18b) resulted in slightly increased IC\textsubscript{50} values (0.5-3.4 \textmu M) and the o-trifluoromethyl derivative 18c lost 10-fold activity (IC\textsubscript{50} of 8-32 \textmu M). Interestingly, the disubstituted derivatives (2,3-difluor (18d), 2,6-difluor (18e) and 2,5-dimethoxy (18f)) were all moderate inhibitors of cell growth, being less potent than the related orto-monomosubstituted compounds (i.e 18b and 16c, respectively) whereas the 2,6-dimethoxy derivative (18g) showed comparable activity as 18b.

Concerning fragment A, substitution of the phenyl ring by a cyclohexyl (compare 16c and 23a) resulted in at least 10-fold higher IC\textsubscript{50} values. Replacement by a benzyl or a gem-dimethyl (compounds 23b and 23c) was even more detrimental for activity, suggesting the importance of an aromatic ring directly attached to the cyclohexane-1,3-dione. Finally, we introduced a variety of substituents at this phenyl ring (compounds 26a-l, Table 1). Compounds substituted with Me or F at orto in fragment A (26a and 26b) showed some anti-proliferative activity against the different cell lines, but at IC\textsubscript{50} values 50-fold higher than the unsubstituted phenyl derivative 16c. Introduction of a double substitution at 2 and 6 either with methyl or fluorine (26c and 26d, respectively) reduced the anti-proliferative effect even more. On the other hand, the meta-substituted compounds 26e, 26f and 26k were nicely active in all cell lines with IC\textsubscript{50} values around 0.17-0.79 \textmu M. Only compound 26l, with an OH at meta, was less active with an IC\textsubscript{50} value in the low-\textmu M range. Finally, the para-substituted compounds showed only very moderate anti-proliferative activity, the only exception being 26j, containing a fluorine, that was still 20-fold less active than the unsubstituted compound 16c.

From this evaluation some structural requirements for anti-proliferative activity among this family of compounds can be extracted: (i) the aromatic ring D needs to be substituted at the orto position with either a methoxy, a methyl, a Cl or to a lesser extend a F; (ii) the optimal alkyl chain at fragment C seems to be one carbon atom; (iii) fragment A must be a phenyl, preferentially unsubstituted or with a methyl or methoxy at meta.

A number of in vitro experiments were then carried out to elucidate the mechanism of action of these new anti-proliferative compounds. Colchicine was included as positive control.
Inhibition of Cell Cycle Progression.

To investigate at which phase of the cell cycle the compounds exert their anti-mitotic effect, we performed cell cycle analysis on HMEC-1 treated with 6 of the most potent compounds (16c, 16d, 18a, 18b, 26e and 26f).35 Thus, HMEC-1 were incubated in the presence of different concentrations of colchicine or tested compound for 24 hours, after which cell cycle distribution was measured by flow cytometry. The results obtained for compounds 16c, 16d and colchicine (1) are shown in Fig 5. Untreated cells showed a classical pattern of proliferating cells distributed in the G1, S and G2/M phase. When these cells were treated with 0.03 µM colchicine, an accumulation in G2/M phase and an increase in subG1 cells undergoing apoptosis, was observed. The same effect was noted in the presence of the tested compounds, although at different concentrations. Compound 16c caused G2/M phase arrest and apoptosis at 0.3 µM, while at least 1 µM of the other inhibitors was needed to exert a comparable activity. Therefore, 16c was the compound of choice for subsequent experiments.

Tubulin binding

VDAs, including colchicine, arrest cell cycle progression at G2/M phase by interfering with tubulin polymerization. In order to evaluate whether our compounds bind to tubulin we carried out a recently described competition assay with N,N’-ethylene-bis(iodoacetamide) (EBI) in human breast cancer MDA-MB-231 cells.36 EBI is an alkylating agent that specifically binds to two cysteine residues present in the colchicine-binding site of tubulin, Cys239 and Cys354. This β-tubulin adduct can be detected by western blot as a second immunoreactive β-tubulin band that migrates faster than β-tubulin itself. So, if EBI is added to cells previously treated with a colchicine-site binder, the binding site is already occupied and the EBI adduct cannot be observed. Since the distance between the different bands depends on the cell type, we used MDA-MB-231 cells in which the two bands can be easily distinguished. As shown in Figure 6, addition of EBI led to the appearance of a second band below the β-tubulin band. Addition of colchicine (1) (0.5 µM) or 16c (10 µM) prevented the covalent binding of EBI in living MDA-MB-231 cells, resulting in the absence of the adduct band. This effect was also observed in the presence of the other compounds (data not shown). Thus, these data indicate that our compounds induce their anti-mitotic effects by binding to the colchicine-binding site in β-tubulin.
Binding of some representative inhibitors (14c, 16c and 26f) to tubulin was confirmed by competition with 2-methoxy-5-(2,3,4-trimethoxyphenyl)-2,4,6-cycloheptatrien-1-one (MTC) and (R)-(+)ethyl 5-amino 2-methyl-1,2-dihydro-3-phenylpyrido[3,4-b]pyrazin-7-yl carbamate (R-PT), two well characterized, reversible colchicine-binding site ligands.37-40 Indeed all three compounds displace both MTC and R-PT as determined by fluorescence spectroscopy (as shown in Fig. S3 for 16c in the Supporting Information).

In order to determine their binding affinities, competition experiments with MTC, the lower affinity compound (Ka 4.7x10^5 M^-1),41 were performed. The data indicated a very high affinity making it difficult to precisely quantify the binding constant.42 Therefore, the displacement of R-PT was used to measure the binding constants. First, we determined the binding constant of R-PT at 25º C (5.1x10^6 M^-1) which was found to be 10 times higher than that of MTC, and similar to the previously reported value (3.2x10^6 M^-1).43 Then, R-PT displacement experiments were carried out with our three compounds allowing the determination of their binding affinities (Figure 7) in the sub-micromolar range (Table 2). Interestingly, the binding constants obtained are higher than those previously determined for other classical colchicine-binding site ligands, such as Nocodazole (4×10^5 M^-1),44 or Podofilotoxin (1.8×10^6 M^-1).45

**Inhibition of mitotic spindle formation.**

Next, we investigated whether the interaction of the compounds with tubulin affects the organization of tubulin into mitotic spindles during cell division.47 Therefore, MDA-MB-231 cells were treated with colchicine (0.1 µM) or 16c (1 µM) for 8 h after which the microtubules were visualized by immunofluorescence microscopy (Fig. 8). Whereas in the control cells (treated with 0.1% DMSO) chromosomes lined up along the metaphase plate and normal bipolar mitotic spindles extended from the cell poles toward the midpoint, highly aberrant, multipolar spindles were formed in the presence of colchicine or 16c. Thus, the inability of the cells to proceed through the cell cycle after treatment with 16c may be explained by the lack of DNA organization into a metaphase plate, which is required for proper cell division.

**Anti-vascular Activity.**

Finally, we studied the capacity of this new family of anti-proliferative compounds to destroy a preexisting vasculature network formed by endothelial cells. HMEC-1 were seeded on top of matrigel, which induces their reorganization into tube-like structures within 3 h.34 Then, colchicine (0.1-10 µM) or 16c (0.1-10 µM)
were added. After 90 min of treatment the plates were photographed and the extent of tube formation was evaluated. As shown in Figure 9, colchicine displayed vascular disrupting effects at 0.1 µM with total destruction of the cord-like structures at 0.3 µM. Similar effects were observed in the presence of 16c, but at 10-fold higher concentrations.

**Conclusions**

Vascular disrupting agents show a great potential in cancer therapies and the limitations observed with those prototypes in clinical trials demand new chemical entities that could be useful as VDAs. Herein we describe the identification of a new family of anti-proliferative compounds with potential VDA-like properties.

Starting from a ligand 3-D shape similarity virtual screening approach using as queries compounds that target the α,β−tubulin dimer at the colchicine-binding site, a cyclohexanedione derivative 9 was identified as a moderate cell growth inhibitor, both in tumor and endothelial cells. By performing a synthetic programme of structural analogues, structure-activity relationships could be established and six derivatives showed anti-proliferative activities in the sub-micromolar range, being 100-fold more potent than the initial hit, and only 10-fold less effective than colchicine itself, a well-known, but toxic, VDA. The compounds described herein inhibit endothelial and tumor cell proliferation at sub-micromolar concentrations, doses at which also their inhibitory effects on the cell cycle and endothelial morphology were observed. Binding at the colchicine site was confirmed by means of fluorescence measurements of MTC and R-PT displacement, and the binding constants obtained are in the µM range. Thus, these colchicine-binding compounds represent a very promising family of anti-proliferative and anti-vascular agents that are being further explored.

**Experimental section**

**Computational methods.**

All VS calculations have been performed in a Dell Precision T7400 workstation. The software used for the VS (FILTER, OMEGA, ROCS and VIDA) were obtained from OpenEye Scientific Software (www.eyesopen.com) by an academic license.

**Database preparation.** The database compounds were obtained from the ZINC database, version 8 and contained about 8.5 million compounds in SMILE string format. The initial filter was performed with the
“drug-like filter” implemented in the tool FILTER v2.1.1 and only selected terms were customized from the default file. The filtered dataset was thus reduced to 2.8 million compounds. 2D to 3D conversion and generation of conformational ensembles was carried out with OMEGA v2.3.2. This tool is controlled by a configuration file in which three parameters were changed as compared to the default values. The maximum number of output conformations of each molecule was set to 300; the energy window was set to 20 kcal/mol in order to discard high-energy conformations; and finally, a rmsd cut-off of 0.75 Å below which two conformations are considered to be the same. When the data set was then submitted to OMEGA, a total of 510 million chemical entities (in mol2 format) were obtained and submitted to the VS protocol.

**Virtual Screening with ROCS.** ROCS v3.1.2 (Rapid Overlay of Chemical Structures) method was used for the virtual screening. ROCS overlays the multiconformer compound’s database in shape and chemistry (“color”) with respect to the reference ligand. As query molecules, a single conformation found in the α,β-tubulin-ligand X-Ray structures of DAMA-colchicine (pdb 1sa0)\(^{26}\) and TN-16 (pdb 3hkd)\(^{27}\) were used. ROCS was run using the default settings. The output conformations of ROCS were ranked according to their shape similarity with the query using a Tanimoto coefficient of 0.75 and a ComboScore (shape plus color) of 1.4. Final results were visualized and analyzed with VIDA v4.1.1. VS carried out with TN-16 as query, resulted in a total of 170 compounds, that were clustered by visual inspection into 9 chemical families. Only the best ranked compound of each family was selected as hit for the antiproliferative assay, of which, only six were available at the time and subsequently tested.

**Chemistry procedures**

Melting points were obtained on a Reichert-Jung Kofler apparatus and are uncorrected. The elemental analysis was performed with a Heraeus CHN-O-RAPID instrument. The elemental compositions of the compounds agreed to within ±0.4% of the calculated values. For all the tested compounds, satisfactory elemental analysis was obtained supporting >95% purity. Electrospray mass spectra were measured on a quadrupole mass spectrometer equipped with an electrospray source (Hewlett-Packard, LC/MS HP 1100). \(^1\)H and \(^{13}\)C NMR spectra were recorded on a Varian INNOVA 300 operating at 299 MHz (\(^1\)H) and 75 MHz (\(^{13}\)C), respectively, a Varian INNOVA-400 operating at 399 MHz (\(^1\)H) and 99 MHz (\(^{13}\)C), respectively, and a VARIAN SYSTEM-500 operating a 499 MHz (\(^1\)H) and 125 MHz (\(^{13}\)C), respectively.
Analytical TLC was performed on silica gel 60 F<sub>254</sub> (Merck) precoated plates (0.2 mm). Spots were detected under UV light (254 nm) and/or charring with ninhydrin or phophomolibdic acid. Separations on silica gel were performed by preparative centrifugal circular thin-layer chromatography (CCTLC) on a Chromatotron<sup>R</sup> (Kiesegel 60 PF<sub>254</sub> gipshaltig (Merck)), with layer thickness of 1 and 2 mm and flow rate of 4 or 8 mL/min, respectively. Flash column chromatography was performed in a Biotage Horizon instrument.

Microwave reactions were performed using the Biotage Initiator 2.0 single-mode cavity instrument from Biotage (Uppsala). Experiments were carried out in sealed microwave process vials utilizing the standard absorbance level (400 W maximum power). The temperature was measured with an IR sensor on the outside of the reaction vessel.

The synthesis of the different series of compounds has been performed applying the here described general procedures. A few key compounds are described in this Experimental section. Still full details and analytical and spectroscopic data for all the compounds are included in the Supporting Information.

5-oxo-1,2,5,6-tetrahydro-[1,1'-biphenyl]-3-yl propionate (11).

To a solution of 5-phenylcyclohexane-1,3-dione (300 mg, 1.59 mmol) and DMAP (59 mg, 0.48 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (1.6 mL), Hunig’s base (DIPEA) (277 µL, 1.59 mmol) was added. Then, propionyl chloride (147 µL, 1.59) was added dropwise and the reaction mixture was refluxed for 2 h. After cooling, HCl 1N (10 mL) was added and the crude extracted with ethyl acetate (10 mL x 3). The organic layer was washed with brine (15 mL), dried over Na<sub>2</sub>SO<sub>4</sub>, concentrated, and purified by flash chromatography (hexane/ethyl acetate 3:1) to yield 250 mg (64 %) of 11 as an oil. EM (ES, positive mode): m/z 245 (M+H)<sup>+</sup>. <sup>1</sup>H NMR (DMSO-d<sub>6</sub>, 300 MHz) δ: 1.07 (t, 3H, J = 7.4 Hz, CH<sub>3</sub>), 2.57 (s, 3H, CH<sub>3</sub>), 2.66 (m, 2H, H-4, H-6), 3.28 (m, 1H, H-5), 5.91 (d, 1H, J = 2.1 Hz, H-2), 7.24 (m, 1H, Ar), 7.35 (m, 4H, Ar).

5-Phenyl-2-propionylcyclohexane-1,3-dione (12).

To a solution of 5-phenylcyclohexane-1,3-dione (300 mg, 1.59 mmol) and DMAP (59 mg, 0.48 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (1.6 mL), Hunig’s base (DIPEA) (277 µL, 1.59 mmol) was added. Then, propionyl chloride (147 µL, 1.59) was added dropwise and the reaction mixture was refluxed for 2 h. After cooling, HCl 1N (10 mL) was added and the crude extracted with ethyl acetate (10 mL x 3). The organic layer was washed with brine (15 mL), dried over Na<sub>2</sub>SO<sub>4</sub>, concentrated, and purified by flash chromatography (hexane/ethyl acetate 3:1) to yield 250 mg (64 %) of 11 as an oil. EM (ES, positive mode): m/z 245 (M+H)<sup>+</sup>. <sup>1</sup>H NMR (DMSO-d<sub>6</sub>, 300 MHz) δ: 1.07 (t, 3H, J = 7.4 Hz, CH<sub>3</sub>), 2.57 (s, 3H, CH<sub>3</sub>), 2.66 (m, 2H, H-4, H-6), 2.81 (m, 2H, H-4, H-6), 3.28 (m, 1H, H-5), 5.91 (d, 1H, J = 2.1 Hz, H-2), 7.24 (m, 1H, Ar), 7.35 (m, 4H, Ar).
m/z 245 (M+H). 1H NMR (DMSO-d$_6$, 300 MHz) δ (enol form): 1.05 (t, 3H, J = 7.3 Hz, CH$_3$), 2.62-2.71 (m, 2H, H-4, H-6), 2.87-2.95 (m, 2H, H-4, H-6), 3.01 (q, 2H, J = 7.2 Hz, CH$_2$), 3.46 (m, 1H, H-5), 7.23-7.38 (m, 5H, Ar). Although this compound was mentioned in ref 28 no analytical or spectroscopic data were provided.

**General procedure for the reaction of 2-acyl-5-phenylcyclohexane-1,3-diones with anilines**

A microwave vial was charged with 2-acyl-5-phenylcyclohexane-1,3-dione (1.0 mmol), the appropriate aniline (1.5 mmol) and 4 Å molecular sieves in toluene (2 mL). The reaction vessel was sealed and heated in a microwave reactor at 150 ºC for 2 h. After cooling, the solvent was evaporated. The resulting residue was purified as specified.

2-(1-((3-Hydroxyphenyl)amino)propylidene)-5-phenylcyclohexane-1,3-dione (14a).

Following the general procedure for the reaction of 2-acyl-5-phenylcyclohexane-1,3-diones with anilines, a microwave vial was charged with 5-phenyl-2-propionylcyclohexane-1,3-dione (12) (40 mg, 0.16 mmol) and 3-aminophenol (26 mg, 0.24 mmol) in toluene. The residue was worked up and purified by CCTLC in the Chromatothron (dichloromethane/methanol, 40:1) to yield 50 mg (93%) of 14a as a white solid. Mp 199-201 ºC. EM (ES, positive mode): m/z 336 (M+H). 1H NMR (DMSO-d$_6$, 500 MHz) δ: 1.06 (t, 3H, J = 7.3 Hz, CH$_3$), 2.60-2.64 (m, 2H, H-4, H-6), 2.80-2.89 (m, 4H, H-4, H-6, CH$_2$), 3.33 (m, 1H, H-5), 6.65 (t, 1H, J = 2.2 Hz, Ar), 6.70 (m, 1H, Ar), 6.81 (ddd, 1H, J = 8.2, 2.4, 0.9 Hz, Ar), 7.18–7.25 (m, 1H, Ar), 7.27 (t, 1H J = 8.0 Hz, Ar), 7.32 (d, 4H, J = 4.4 Hz, Ar), 9.90 (br s, 1H, OH), 14.94 (br s, 1H, NH). 13C NMR (DMSO-d$_6$, 125 MHz) δ: 12.8 (CH$_3$), 23.4 (CH$_2$), 36.0 (C-5), 46.0 (C-4, C-6), 106.7 (NHC=), 112.9, 115.8, 116.4, 126.5, 126.7, 128.5, 130.4, 136.8, 143.4, 158.2 (Ar), 177.0 (NHC=C). Anal. calc. for (C$_{21}$H$_{21}$NO$_3$): C, 75.20; H, 6.31; N, 4.18. Found: C, 75.33; H, 6.12; N, 4.36.

2-(1-((4-Hydroxyphenyl)amino)propylidene)-5-phenylcyclohexane-1,3-dione (14b).

Following the general procedure for the reaction of 2-acyl-5-phenylcyclohexane-1,3-diones with anilines, a microwave vial was charged with 5-phenyl-2-propionylcyclohexane-1,3-dione (12) (40 mg, 0.16 mmol) and 4-aminophenol (26 mg, 0.24 mmol) in toluene. The residue was worked up and purified by CCTLC in the Chromatothron (dichloromethane/methanol 40:1) to yield 30 mg (56%) of 14b as a white solid. Mp 203-205 ºC. EM (ES, positive mode): m/z 336 (M+H). 1H NMR (DMSO-d$_6$, 300 MHz) δ: 1.02 (t, 3H, J = 7.3 Hz, CH$_3$), 2.59-2.63 (m, 2H, H-4, H-6), 2.78-2.88 (m, 4H, H-4, H-6, CH$_2$), 3.32 (m, 1H, H-5), 6.85 (m, 2H, Ar), 7.11 (m, 2H, Ar), 7.23 (m, 1H, Ar), 7.33 (d, 4H, J = 4.4 Hz, Ar), 9.81 (br s, 1H, OH), 14.73 (br s, 1H, NH). 13C NMR (DMSO d$_6$, 100 MHz) δ: 13.0 (CH$_3$), 23.7 (CH$_2$), 36.5 (C-5), 46.5 (C-4, C-6), 107.1 (NHC=C),
2-(1-((2-Methoxyphenyl)amino)propylidene)-5-phenylcyclohexane-1,3-dione (14c).

Following the general procedure for the reaction of 2-acyl-5-phenylcyclohexane-1,3-diones with anilines, a microwave vial was charged with 5-phenyl-2-propionylcyclohexane-1,3-dione (12) (35 mg, 0.14 mmol) and o-anisidine (24 µL, 0.21 mmol) in toluene. The residue was worked up and purified by CCTLC in the Chromatothron (hexane/ethyl acetate, 5:1) to yield 48 mg (98%) of 14c as a white solid. Mp 93-95 ºC. EM (ES, positive mode): m/z 350 (M+H)+. 1H NMR (DMSO-d6, 400 MHz) δ: 0.98 (t, 3H, J = 7.3 Hz, CH3), 2.59-2.64 (m, 2H, H-4, H-6), 2.80-2.84 (m, 4H, H-4, H-6, CH2), 3.36 (m, 1H, H-5), 3.82 (s, 3H, OCH3), 7.06 (td, 1H, J = 7.6, 1.2 Hz, Ar), 7.19–7.27 (m, 2H, Ar), 7.30 (dd, 1H, J = 7.7, 1.5 Hz, Ar), 7.33 (d, 2H, J = 1.4 Hz, Ar), 7.34 (s, 2H, Ar), 7.39–7.46 (m, 1H, Ar), 14.69 (br s, 1H, NH). 13C NMR (DMSO-d6, 100 MHz) δ: 12.2 (CH3), 25.5 (CH2), 36.0 (C-5), 46.1 (C-4, C-6), 55.8 (OCH3), 106.9 (NHC=C), 112.4, 120.7, 124.2, 126.5, 126.7, 127.5, 128.5, 129.7, 143.5, 153.5 (Ar), 178.0 (NHC=C). Anal. calc. for (C22H23NO3): C, 75.48; H, 6.52; N, 4.00.

Details of the synthesis and analitical and spectroscopic data for compounds 14d-j are provided in the Supporting Information.

2-Butyryl-5-phenylcyclohexane-1,3-dione (15a)

Following the described procedure for the synthesis of 12, a microwave vial was charged with 5-phenylcyclohexane-1,3-dione (10) (500 mg, 2.66 mmol), butyrylchloride (548 µL, 5.31 mmol), anhydrous K2CO3 (809 mg, 5.85 mmol), 1,2,4-triazole (73 mg, 1.06 mmol) and tetrabutylammonium bromide (429 mg, 1.33 mmol) in anhydrous acetonitrile (10 mL) to yield 200 mg (29%) of 15a as a white solid. Mp 68-70 ºC. EM (ES, positive mode): m/z 259 (M+H)+. 1H NMR (DMSO-d6, 300 MHz) δ (enol form): 0.92 (t, 3H, J = 7.3 Hz, CH3), 1.58 (m, 2H, CH2), 2.69 (m, 2H, H-4, H-6), 2.92 (m, 2H, H-4, H-6), 2.97 (t, 2H, J = 7.4 Hz, CH2), 3.37-3.45 (m, 1H, H-5), 7.25 (m, 1H Ar), 7.33 (m, 4H, Ar). Note: Although this compound was mentioned in ref 29, no analitical data were provided.

2-Acetyl-5-phenylcyclohexane-1,3-dione (15b).

Following the described procedure for the synthesis of 12, a microwave vial was charged with 5-phenylcyclohexane-1,3-dione (10) (500 mg, 2.66 mmol), acetylchloride (417 µL, 5.31 mmol), anhydrous K2CO3 (809 mg, 5.85 mmol), 1,2,4-triazole (73 mg, 1.06 mmol) and tetrabutylammonium bromide (429 mg,
1.33 mmol) in anhydrous acetonitrile (10 mL) to yield 350 mg (54%) of 15b as a white solid. Mp 100-102 °C.

EM (ES, positive mode): m/z 231 (M+H)+. 1H NMR (DMSO-d6, 300 MHz) δ (enol form): 2.55 (s, 3H, CH3), 2.66-2.72 (m, 2H, H-4, H-6), 2.94 (m, 2H, H-4, H-6), 3.36-3.47 (m, 1H, H-5), 7.22-7.34 (m, 5H, Ar). Note: Although this compound was mentioned in ref 29, no analitical data were provided.

2-(1-((2-Methoxyphenyl)amino)butylidene)-5-phenylcyclohexane-1,3-dione (16a).

A mixture of 2-butyryl-5-phenylcyclohexane-1,3-dione (15a) (60 mg, 0.23 mmol) and o-anisidine (30 µL, 0.22 mmol) in 1.5 mL of toluene was heated for two days under reflux in a flask equipped with a Dean-Stark trap. Volatiles were removed and the crude reaction mixture was purified by CCTLC in the Chromatothron (hexane/ethyl acetate, 5:1) to yield 70 mg (84%) of 16a as a pale yellow solid. Mp 108-110 °C. EM (ES, positive mode): m/z 364 (M+H)+. 1H NMR (DMSO-d6, 400 MHz) δ: 0.74 (t, 3H, J = 7.3 Hz, CH3), 1.40 (m, 2H, CH2), 2.59-2.63 (m, 2H, H-4, H-6), 2.77 (m, 4H, H-4, H-6, CH2), 3.29-3.37 (m, 1H, H-5), 3.82 (s, 3H, OCH3), 7.05 (m, 1H, J = 7.6, 1.2 Hz, Ar), 7.21 (dd, 1H, J = 8.4, 1.2 Hz, Ar), 7.25 (m, 1H, Ar), 7.28 (dd, 1H, J = 7.7, 1.6 Hz, Ar), 7.33 (d, 2H, J = 1.3 Hz, Ar), 7.33 (s, 2H, Ar), 7.42 (dddd, 1H, J = 8.6, 7.6, 1.6 Hz, Ar), 14.73 (br s, 1H, NH). 13C NMR (DMSO-d6, 100 MHz) δ: 14.1 (CH3), 21.3 (CH2), 31.9 (CH2), 36.0 (C-5), 46.0, 46.5 (C-4, C-6), 55.8 (OCH3), 107.2 (NHC=C), 112.3, 112.7, 124.4, 126.5, 126.5, 127.5, 128.5, 129.6, 143.4, 153.5 (Ar), 176.4 (NHC=C). Anal. calc. for (C23H25NO3): C, 76.01; H, 6.93; N, 3.85. Found: C, 75.93; H, 6.68; N, 4.10.

5-Phenyl-2-(1-(o-tolylamino)butylidene)cyclohexane-1,3-dione (16b).

Following the general procedure for the reaction of 2-acyl-5-phenylcyclohexane-1,3-diones with anilines, a microwave vial was charged with 2-butyryl-5-phenylcyclohexane-1,3-dione (15a) (30 mg, 0.12 mmol) and o-toluidine (18 µL, 0.17 mmol) in toluene. The residue was worked up and purified by CCTLC in the Chromatothron (hexane/ethyl acetate, 5:1) to yield 42 mg (99%) of 16b as a white solid. Mp 98-100 °C. EM (ES, positive mode): m/z 348 (M+H)+. 1H NMR (DMSO-d6, 500 MHz) δ: 0.74 (t, 3H, J = 7.3 Hz, CH3), 1.40 (m, 2H, CH2), 2.17 (s, 3H, CH3), 2.63 (ddd, 2H, J = 16.7, 3.7 Hz, H-4, H-6), 2.75 (t, 2H, J = 7.3 Hz, CH2), 2.85 (m, 2H, H-4, H-6), 3.32-3.39 (m, 1H, H-5), 7.22-7.26 (m, 1H, Ar), 7.27 (dd, 1H, J = 7.2, 2.1 Hz, Ar), 7.34 (m, 6H, Ar), 7.41 (m, 1H, Ar), 14.91 (br s, 1H, NH). 13C NMR (DMSO-d6, 125 MHz) δ: 14.2 (CH3), 17.5 (CH3), 21.3 (CH2), 31.8 (CH2), 36.0 (C-5), 46.0 (C-4, C-6), 107.0 (NHC=C), 126.5, 126.7, 126.8, 126.9, 128.3, 128.5, 131.0, 133.6, 135.0, 143.5 (Ar), 176.1 (NHC=C). Anal. calc. for (C23H23NO2): C, 79.51; H, 7.25; N, 4.03. Found: C, 79.27; H, 6.98; N, 4.11.
2-(1-((2-Methoxyphenyl)amino)ethylidene)-5-phenylcyclohexane-1,3-dione (16c).

Following the general procedure for the reaction of 2-acyl-5-phenylcyclohexane-1,3-diones with anilines, a microwave vial was charged with 2-acetyl-5-phenylcyclohexane-1,3-dione (15b) (40 mg, 0.17 mmol) and o-anisidine (30 µL, 0.26 mmol) in toluene. The residue was worked up and purified by CCTLC in the Chromatotron (hexane/ethyl acetate, 5:1) to yield 49 mg (86%) of 16c as a white solid. Mp 139-141 °C. EM (ES, positive mode): m/z 336 (M+H)+. 1H NMR (DMSO-d6, 500 MHz) δ: 2.41 (s, 3H, CH3), 2.60-2.63 (m, 2H, H-4, H-6), 2.82 (m, 2H, H-4, H-6), 3.36 (m, 1H, H-5), 3.84 (s, 3H, OCH3), 7.04 (td, 1H, J = 7.6, 1.2 Hz, Ar), 7.20 (dd, 1H, J = 8.5, 1.2 Hz, Ar), 7.21-7.26 (m, 1H, Ar), 7.32 (dd, 1H, J = 7.8, 1.6 Hz, Ar), 7.33 (d, 2H, J = 0.8 Hz, Ar), 7.34 (s, 2H, Ar), 7.39 (m, 1H, Ar), 14.78 (br s, 1H, NH). 13C NMR (DMSO-d6, 125 MHz) δ: 19.7 (CH3), 36.1 (C-5), 46.4 (C-4, C-6), 55.8 (OCH3), 108.4 (NHC=), 112.3, 120.6, 124.5, 126.5, 126.7, 128.9, 128.5, 129.2, 143.5, 153.1 (Ar), 172.4 (NHC=). Anal. calc. for (C21H21NO3): C, 75.20; H, 6.31; N, 4.31. Found: C, 74.98; H, 6.32; N, 4.20.

5-Phenyl-2-(1-((o-tolylamino)ethylidene)cyclohexane-1,3-dione (16d).

Following the general procedure for the reaction of 2-acyl-5-phenylcyclohexane-1,3-diones with anilines, a microwave vial was charged with 2-acetyl-5-phenylcyclohexane-1,3-dione (15b) (30 mg, 0.13 mmol) and o-toluidine (22 µL, 0.20 mmol) in toluene. The residue was worked up and purified by CCTLC in the Chromatotron (hexane/ethyl acetate, 5:1) to yield 25 mg (60%) of 16d as a white solid. Mp 128-130 °C. EM (ES, positive mode): m/z 320 (M+H)+. 1H NMR (DMSO-d6, 300 MHz) δ: 2.20 (s, 3H, CH3), 2.38 (s, 3H, CH3), 2.59-2.66 (m, 2H, H-4, H-6), 2.79-2.89 (m, 2H, H-4, H-6), 3.36 (m, 1H, H-5), 7.23-7.40 (m, 9H, Ar), 14.91 (br s, 1H, NH). 13C NMR (DMSO-d6, 100 MHz) δ: 17.4 (CH3), 19.6 (CH3), 36.1 (C-5), 45.8 (C-4, C-6), 108.2 (NHC=), 126.5, 126.7, 126.9, 128.1, 128.49, 131.0, 133.3, 135.1, 143.5, 159.9 (Ar), 172.7 (NHC=). Anal. calc. for (C21H21NO2): C, 78.97; H, 6.63; N, 4.39. Found: C, 78.69; H, 6.54; N, 4.27.

Biological methods

Cell proliferation.

Endothelial cells. Mouse brain endothelial cells (MBEC), bovine aortic endothelial cells (BAEC) and human dermal microvascular endothelial cells (HMEC-1) were seeded in 48-well plates at 10,000 cells/well (except HMEC-1 at 20,000/well). After 24 h, 5-fold dilutions of the compounds were added. The cells were allowed to proliferate 3 days (or 4 days for HMEC-1) in the presence of the compounds, trypsinized, and counted by means of a Coulter counter (Analis, Belgium).
**Tumor cells.** Human cervical carcinoma (HeLa) cells were seeded in 96-well plates at 15,000 cells/well in the presence of different concentrations of the compounds. After 4 days of incubation, the cells were trypsinized and counted in a Coulter counter. Suspension cells (Mouse leukemia L1210 and human lymphoid Cem cells) were seeded in 96-well plates at 60,000 cells/well in the presence of different concentrations of the compounds. L1210 and Cem cells were allowed to proliferate for 48 h or 96 h, respectively and then counted in a Coulter counter. The 50% inhibitory concentration (IC$_{50}$) was defined as the compound concentration required to reduce cell proliferation by 50%. Colchicine was added as reference compound.

**Cell cycle analysis.** HMEC-1 cells were seeded in 6-well plates at 125,000 cells/well in DMEM with 10% FCS. After 24 h, the cells were exposed to different concentrations of the compounds. After 24 h, the DNA of the cells was stained with propidium iodide using the CycleTEST PLUS DNA Reagent Kit (BD Biosciences, San Jose, CA). The DNA content of the stained cells was assessed by flow cytometry on a FACSCalibur flow cytometer and analyzed with CellQuest software (BD Biosciences) within 3h after staining. Cell debris and clumps were excluded from the analysis by appropriate dot plot gating. Percentages of sub-G1, G1, S, and G2/M cells were estimated using appropriate region markers.$^{35}$ Colchicine was added as reference compound.

**Tube formation.** Wells of a 96-well plate were coated with 70 µl matrigel (10 mg/ml, BD Biosciences, Heidelberg, Germany) at 4°C. After gelatinization at 37°C during 30 min, HMEC-1 cells were seeded at 60,000 cells/well on top of the matrigel in 200 µl DMEM containing 10% FCS. After 3 h of incubation at 37°C, when the endothelial cells had reorganized to form tube-like structures, the compounds were added. Ninety minutes later, colchicine, which was added as a reference compound, had destroyed the endothelial tubes. The cultures were photographed at 100 x magnification.

**Tubulin binding.** Human breast carcinoma MDA-MB-231 cells were seeded in 6-well plates at 500,000 cells/well. After 48 h, compounds were added to the cells for 16 h before adding EBI (N,N’-ethylene-bis(iodoacetamide) at 100 µM. After 1.5 h, the cells were harvested and cell extracts were prepared for western blot analysis. Twenty µg of proteins were subjected to gel electrophoresis using 0.1% SDS (85% purity) and 10% polyacrylamide gels. After electrophoresis, proteins were transferred to pretreated Hybond-P polyvinylidene difluoride (PVDF) membranes (Amersham Biosciences). The membranes were incubated for 1 h at room temperature in blocking buffer (2.5 % non-fat dry milk in PBS containing 0.1% Tween) and subsequently for 16 h at 4°C in blocking buffer with primary antibodies raised against β-tubulin. After washing, the membranes were incubated with the corresponding horseradish peroxidase-conjugated
secondary antibody in blocking buffer for 25 min at room temperature. Next, the membranes were washed extensively. Immunoreactive proteins were detected by chemiluminescence (ECLplus, Bio-Rad).

In living cells, EBI cross-links the cysteine residues at positions 239 and 354 of β-tubulin. This β-tubulin adduct formed by EBI is easily detectable by western blot as a second immunoreactive band that migrates faster than β-tubulin. Colchicine was added as a reference compound.

**Determination of binding constants.**

Proteins and ligands. Calf brain tubulin was purified as described.48 2-methoxy-5-(2,3,4-trimethoxyphenyl)-2,4,6-cycloheptatrien-1-one (MTC)49 was a kind gift of Prof. T.J. Fitzgerald School of Pharmacy, Florida A & M University. (R)-(+)ethyl 5-amino 2-methyl-1,2-dihydro-3-phenylpyrido[3,4-b]pyrazin-7-yl carbamate (R-PT) ENREF_5350 was a kind gift of Prof. G.A. Rener, Organic Chemistry Research Department, Southern Research Institute, Birmingham, Alabama. The compounds were diluted in 99.8% D6-DMSO (Merck, Darmstadt, Germany) to a final concentration of 10 mM and stored at -80 ºC.

Determination of binding constants. Competition of the compounds with MTC was tested by the change in the intensity of fluorescence of MTC upon binding to tubulin. The fluorescence emission spectra (excitation at 350 nm) of 10 µM Tubulin and 10 µM MTC in 10 mM Sodium Phosphate, 0.1 mM GTP pH 7.0 were measured in the presence or the absence of 20 µM of the desired ligand with 5 nm excitation and emission slits using a Jobin-Ybon SPEX Fluoromax-2 (HORIBA, Ltd. Kyoto, Japan). The decrease in the intensity of the fluorescence in the presence of the competitor ligand indicated competition for the same binding site.

The binding constant of R-PT for dimeric tubulin was determined using the competition method in 10 mM sodium phosphate, 0.1 mM GTP pH 7.0 at 25ºC. To do so 0.2 µM of R-PT was incubated with increasing amounts of tubulin up to 10µM and vice versa, 0.2 µM of tubulin was incubated with increasing amounts of R-PT up to 10µM, the fluorescence emission spectra (excitation 374 nm) of the samples (5 nm excitation and emission slits) were determined using a Jobin-Ybon SPEX Fluoromax-2 (HORIBA, Ltd. Kyoto, Japan). Using these spectra it is possible to calculate the free and the bound R-PT concentration for each sample and thus to determine the binding constant of R-PT for tubulin.

Once K_b of R-PT is determined (5.1x10^6 M^-1) this compound could be used as a reference ligand as described in .For that purpose, the fluorescence emission of a previous mixed sample of 0.2µM of R-PT and 0.2 µM of tubulin was evaluated in the presence of increasing concentrations of studied ligand in a black 96-well plate (0; 0.05; 0.2; 0.5; 2; 5; 10; 30; 50; 70 µM). The samples were incubated 30 minutes at 25ºC in a
Varioskan plate reader (Thermo Scientific Waltham, Massachusetts, USA) before the fluorescence emission intensity at 456 nm (excitation 374 nm) was measured. The data were analyzed and the binding constants determined using Equigra V5.0.42

**Immunocytochemistry.** MDA-MB-231 cells were grown on poly-L-Lysine pre-coated 8-well chamber slides (Lab-Tek, Nunc, Roskilde, Denmark) in DMEM containing 10% FBS and exposed to compound (DMSO, 16c or colchicine). After 8 h, the cells were fixed in 4% PFA for 15 min at room temperature, washed three times with PBS and permeabilized for 10 min at room temperature with 0.25% Triton X-100 (Sigma-Aldrich). Nonspecific binding sites were blocked for 30 min at room temperature with 0.5% BSA in PBS. The cells were then incubated with a monoclonal anti-β-tubulin antibody (2 µg/ml, Sigma-Aldrich) for 2 h at room temperature, washed three times and incubated for 1 h at room temperature with goat anti-mouse Alexa Fluor 488 (4 µg/ml; Molecular Probes, Invitrogen) in 0.5% BSA. After three washes, nuclei were stained with 300 nM 4′,6-diamidino-2-phenylindole (DAPI; Sigma-Aldrich). Fluorescent microscopic analysis was done with an Axiovert 200 M inverted microscope (Zeiss, Göttingen, Germany), using an EC Plan-Neofluar 40x/1.30 oil objective. Pictures were taken with an AxioCamMRm camera and processed with AxioVision Release 4.6 software (Zeiss).

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SUPPORTING INFORMATION AVAILABLE. The synthesis and spectroscopic data of all the synthesized compounds is included. Molecular structures (SMILES strings) of the virtual screening hits tested are included in Table S1. Anti-proliferative activity of the VS hits in endothelial and tumor cell lines is included in Table S2. Dose-response curves of 9 in endothelial and tumor cell lines is included as Figure S2. Fluorescence emission spectra for the displacement of MTC by 9 and R-PT and MTC by 16c are included as Figures S1 and S3, respectively. This material is available free of charge via the Internet at http://pubs.acs.org

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39. Barbier, P.; Peyrot, V.; Leynadier, D.; Andreu, J. M. The active GTP- and ground CDP-Liganded states of tubulin are distinguished by the binding of chiral isomers of ethyl 5-amino-2-methyl-1,2-dihydro-3-phenylpyrido 3,4-b pyrazin-7-yl carbamate. *Biochemistry (Mosc)*. 1998, 37, 758-768.


**Figure 1.** Chemical structures of selected colchicine-site binders

Colchicine (1)  
ZD8126 (2)  
ABT-751 (3)  
CA-1P (4)  
CA-4P (5)  
AVE8022 (6)
Figure 2. Schematic view of the virtual screening protocol
A. Chemical structures of DAMA-colchicine (7) and TN-16 (8). B. Colchicine binding-site in the α,β-tubulin dimer. C. Superposition of the ligands used in the VS: DAMA-colchicine (green, pdb 1sa0) and TN-16 (cyan, pdb 3hkd).
Figure 4. Structure of the hit compound 9 identified from the VS campaign.
Cell cycle distribution. HMEC-1 were treated with DMSO (A), colchicine (0.03 µM, B), 16c (1 µM, C or 0.3 µM, D) or 16d (1 µM, E or 0.3 µM, F) for 24 h. Next, the cells were harvested, stained with propidium iodide, and cell cycle distribution was evaluated by flow cytometry. Percentages of cells in the different phases of the cell cycle are indicated (G).
Figure 6.

\[ \text{EBI} \quad 0.5 \quad 40 \quad 10 \quad 2.5 \]

\[ \text{colchicine} \quad 0.5 \quad 40 \quad 10 \quad 2.5 \]

\[ \text{16c} \quad 0.5 \quad 40 \quad 10 \quad 2.5 \]

**Inhibition of tubulin binding by 16c.** MDA-MB-231 cells were treated with DMSO, colchicine (0.5 µM) or 16c (40, 10 or 2.5 µM) for 16 h. Next EBI (100 µM) was added and after 1.5 h, the cells were harvested and cell extracts were prepared for western blot analysis using anti-β-tubulin antibody. EBI cross-links cysteine residues in β-tubulin resulting in the formation of a β-tubulin adduct (second immunoreactive band) that migrates faster than β-tubulin. Compounds that bind to the colchicine-binding site in β-tubulin prevent the formation of the EBI:β-tubulin adduct.
Displacement of the fluorescent probe R-PT (0.2 µM) bound to tubulin (0.2 µM) by 14c (black lines and circles), 16c (red lines and squares) and 26f (green lines and triangles) at 25 ºC. The solid lines were generated with the best fit value of the binding equilibrium constant of the competitors, assuming a one-to-one binding to the same site.
Figure 8.

**Inhibition of mitotic spindle formation.** MDA-MB-231 cells were treated for 8 h with DMSO (A), colchicine (0.1 µM, B) or 16c (1 µM, C) fixed and stained with anti-β-tubulin antibody (green) to visualize the microtubules and DAPI for DNA (blue). Column 1: double staining; column 2: DNA staining only; column 3: β-tubulin staining only.
Vascular disrupting effects of 16c. HMEC-1 cells were cultured on matrigel for 3 h to allow the formation of tube-like structures. Then DMSO 0.1% (control), colchicine (0.1 or 0.3 µM) or 16c (0.3, 1 or 3 µM) were added. Images show the vascular network after 90 min of treatment.
Scheme 1a

Reagents and conditions: (a) DMAP, DIPEA, CH₂Cl₂, 70 ºC, 2 h, 64% yield; (b) K₂CO₃ anh., 1,2,4-triazole, CH₃CN, 30 ºC, > 48 h, 30% yield; (c) K₂CO₃ anh., 1,2,4-triazole, Bu₄NBr, CH₃CN, MW, 70 ºC, 2 h, 51% yield. (d) Toluene, 4Å molecular sieves, MW, 150 ºC, 2 h, 30-98% yields.
Scheme 2a

Reagents and conditions: (a) R'COCl, K₂CO₃ anh, 1,2,4-triazole, Bu₄NBr, CH₃CN, MW, 70 °C, 2 h (15a, 29% yield; 15b, 54% yield); (b) substituted aniline, toluene, molecular sieves, MW, 150 °C, 2 h, 60-84% yields.
Scheme 3a

![Chemical structure](image)

\[15b + 17a-g \xrightarrow{(a)} 18a-g\]

- **a** \(R^1=\text{Cl}, R^2=R^3=R^4=\text{H}\)
- **b** \(R^1=\text{F}, R^2=R^3=R^4=\text{H}\)
- **c** \(R^1=\text{CF}_3, R^2=R^3=R^4=\text{H}\)
- **d** \(R^1=R^2=\text{F}, R^3=R^4=\text{H}\)
- **e** \(R^1=R^4=\text{F}, R^2=R^3=\text{H}\)
- **f** \(R^1=R^3=\text{OCH}_3, R^2=R^4=\text{H}\)
- **g** \(R^1=R^4=\text{OCH}_3, R^2=R^3=\text{H}\)

\[^a\text{Reagents and conditions: (a) toluene, molecular sieves, MW, 150 °C, 2 h, 23-78% yields.}\]
Scheme 4a

Reagents and conditions: (a) NaOH (1%), acetone, H₂O, rt, overnight (for 20a, 65% yield); (b) Ph₃P=CHCOCH₃, CHCl₃, 60 °C, 4 h (for 20b, 77% yield); (c) (i) Diethyl malonate, EtONa, EtOH, Δ; (ii) NaOH, 80 °C, 2 h; (iii) HCl, Δ, 1 h (21a 69% yield, 21b 33% yield); (d) ClCOCH₃, K₂CO₃ anh, 1,2,4-triazole, Bu₄NBr, CH₃CN, MW, 70 °C, 2 h (22a 36% yield, 22b 38% yield, 22c was commercially available); (e) toluene, molecular sieves, pressure tube, 110 °C, overnight (for 23a 89% yield, for 23b 73% yield); (f) toluene, molecular sieves, MW, 150 °C, 2 h (for 23c 79% yield).
Scheme 5a

Reagents and conditions: (a) CH3COCl, K2CO3 anh, 1,2,4-triazole, Bu4NBr, CH3CN, MW, 70 °C, 2 h, 14-74% yields; (b) For 26e-j: toluene, molecular sieves, MW 150 °C, 2 h, 39-99% yields; (c) For 26a-d,k: toluene, molecular sieves, pressure tube, 110 °C, overnight, 39-75% yields; (d) BBr3, CH2Cl2, rt, 12 h, 23% yield.
Table 1. Anti-proliferative activity of compounds 14a-j, 16a-d and 18a-g in endothelial and tumor cell lines

<table>
<thead>
<tr>
<th>Compound</th>
<th>HMEC-1</th>
<th>MBEC</th>
<th>BAEC</th>
<th>L1210</th>
<th>CEM</th>
<th>HeLa</th>
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<tbody>
<tr>
<td></td>
<td>IC₅₀ (µM)</td>
<td></td>
<td></td>
<td>IC₅₀ (µM)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Colchicine</td>
<td>0.0038 ± 0.0011</td>
<td>0.031 ± 0.015</td>
<td>0.0069 ± 0.0008</td>
<td>0.010 ± 0.001</td>
<td>0.013 ± 0.001</td>
<td>0.0087 ± 0.0001</td>
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<td>9</td>
<td>22 ± 13</td>
<td>13 ± 5</td>
<td>35 ± 1</td>
<td>13 ± 1</td>
<td>15 ± 9</td>
<td>11 ± 8</td>
</tr>
<tr>
<td>14a</td>
<td>13 ± 6</td>
<td>12 ± 3</td>
<td>7.8 ± 0.9</td>
<td>15 ± 12</td>
<td>48 ± 26</td>
<td>75 ± 25</td>
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<tr>
<td>14b</td>
<td>28 ± 21</td>
<td>18 ± 0</td>
<td>36 ± 2</td>
<td>14 ± 7</td>
<td>34 ± 0</td>
<td>79 ± 42</td>
</tr>
<tr>
<td>14c</td>
<td>1.4 ± 0.2</td>
<td>1.4 ± 0.1</td>
<td>1.2 ± 0.6</td>
<td>1.5 ± 0.2</td>
<td>2.3 ± 1.7</td>
<td>4.3 ± 0.7</td>
</tr>
<tr>
<td>14d</td>
<td>18 ± 9</td>
<td>16 ± 4</td>
<td>64 ± 14</td>
<td>32 ± 9</td>
<td>45 ± 12</td>
<td>66 ± 9</td>
</tr>
<tr>
<td>14e</td>
<td>≥ 100</td>
<td>57 ± 7</td>
<td>≥ 100</td>
<td>58 ± 13</td>
<td>67 ± 11</td>
<td>≥ 100</td>
</tr>
<tr>
<td>14f</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
</tr>
<tr>
<td>14g</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
<td>&gt; 100</td>
</tr>
<tr>
<td>14h</td>
<td>53 ± 11</td>
<td>50 ± 18</td>
<td>43 ± 6</td>
<td>38 ± 0</td>
<td>53 ± 22</td>
<td>76 ± 5</td>
</tr>
<tr>
<td>14i</td>
<td>15 ± 6</td>
<td>8.3 ± 2.1</td>
<td>16 ± 11</td>
<td>22 ± 17</td>
<td>21 ± 11</td>
<td>56 ± 3</td>
</tr>
<tr>
<td>14j</td>
<td>2.1 ± 2.0</td>
<td>2.8 ± 0.2</td>
<td>2.2 ± 0.2</td>
<td>6.9 ± 1.5</td>
<td>6.3 ± 1.2</td>
<td>9.2 ± 5.0</td>
</tr>
<tr>
<td>16a</td>
<td>31 ± 3</td>
<td>27 ± 4</td>
<td>16 ± 3</td>
<td>36 ± 8</td>
<td>41 ± 1</td>
<td>59 ± 23</td>
</tr>
<tr>
<td>16b</td>
<td>44 ± 2</td>
<td>47 ± 3</td>
<td>46 ± 4</td>
<td>41 ± 19</td>
<td>≥ 100</td>
<td>&gt; 100</td>
</tr>
<tr>
<td>16c</td>
<td>0.09 ± 0.01</td>
<td>0.12 ± 0.01</td>
<td>0.09 ± 0.01</td>
<td>0.16 ± 0.08</td>
<td>0.18 ± 0.05</td>
<td>0.18 ± 0.00</td>
</tr>
<tr>
<td>16d</td>
<td>0.26 ± 0.08</td>
<td>0.32 ± 0.04</td>
<td>0.17 ± 0.03</td>
<td>0.30 ± 0.18</td>
<td>0.68 ± 0.09</td>
<td>0.48 ± 0.26</td>
</tr>
<tr>
<td>18a</td>
<td>0.56 ± 0.16</td>
<td>0.47 ± 0.05</td>
<td>0.29 ± 0.03</td>
<td>0.70 ± 0.62</td>
<td>0.64 ± 0.18</td>
<td>0.71 ± 0.20</td>
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<tr>
<td>18b</td>
<td>0.93 ± 0.46</td>
<td>1.0 ± 0.1</td>
<td>0.51 ± 0.01</td>
<td>1.9 ± 0.7</td>
<td>1.1 ± 0.3</td>
<td>3.4 ± 2.4</td>
</tr>
<tr>
<td>18c</td>
<td>7.7 ± 2.1</td>
<td>8.7 ± 0.1</td>
<td>8.1 ± 0.5</td>
<td>14 ± 10</td>
<td>18 ± 2</td>
<td>32 ± 18</td>
</tr>
<tr>
<td>18d</td>
<td>10 ± 2</td>
<td>12 ± 0</td>
<td>14 ± 2</td>
<td>19 ± 1</td>
<td>25 ± 18</td>
<td>36 ± 3</td>
</tr>
<tr>
<td>18e</td>
<td>6.6 ± 0.5</td>
<td>9.2 ± 1.8</td>
<td>8.4 ± 1.5</td>
<td>11 ± 2</td>
<td>21 ± 6</td>
<td>17 ± 4</td>
</tr>
<tr>
<td>18f</td>
<td>2.6 ± 0.2</td>
<td>3.2 ± 0.2</td>
<td>2.6 ± 0.5</td>
<td>36 ± 22</td>
<td>6 - &gt;100*</td>
<td>46 ± 3</td>
</tr>
<tr>
<td>18g</td>
<td>1.4 ± 0.8</td>
<td>1.2 ± 0.2</td>
<td>0.46 ± 0.29</td>
<td>1.0 ± 0.0</td>
<td>1.4 ± 0.7</td>
<td>1.1 ± 0.2</td>
</tr>
<tr>
<td>------</td>
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<td>-----------</td>
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</tr>
</tbody>
</table>

*Indicates no dose response in the indicated concentration range, making accurate IC$_{50}$ calculation impossible.
Table 1 (cont). Anti-proliferative activity of compounds 23a-c and 26a-l in endothelial and tumor cell lines.

<table>
<thead>
<tr>
<th>Compound</th>
<th>Endothelial cells</th>
<th>Tumor cells</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>HMEC-1</td>
<td>MBEC</td>
</tr>
<tr>
<td>Colchicine</td>
<td>0.0038 ± 0.0011</td>
<td>0.031 ± 0.008</td>
</tr>
<tr>
<td>23a</td>
<td>12 ± 1</td>
<td>21 ± 5</td>
</tr>
<tr>
<td>23b</td>
<td>22 ± 2</td>
<td>87 ± 15</td>
</tr>
<tr>
<td>23c</td>
<td>76 ± 18</td>
<td>71 ± 41</td>
</tr>
<tr>
<td>26a</td>
<td>5.8 ± 0.4</td>
<td>6.8 ± 0.8</td>
</tr>
<tr>
<td>26b</td>
<td>4.3 ± 0.3</td>
<td>5.4 ± 0.6</td>
</tr>
<tr>
<td>26c</td>
<td>35 ± 1</td>
<td>39 ± 2</td>
</tr>
<tr>
<td>26d</td>
<td>39 ± 8</td>
<td>60 ± 9</td>
</tr>
<tr>
<td>26e</td>
<td>0.33 ± 0.12</td>
<td>0.42 ± 0.01</td>
</tr>
<tr>
<td>26f</td>
<td>0.30 ± 0.08</td>
<td>0.41 ± 0.08</td>
</tr>
<tr>
<td>26g</td>
<td>11 ± 3</td>
<td>9.2 ± 1.1</td>
</tr>
<tr>
<td>26h</td>
<td>&gt;100</td>
<td>&gt;100</td>
</tr>
<tr>
<td>26i</td>
<td>55 ± 16</td>
<td>46 ± 1</td>
</tr>
<tr>
<td>26j</td>
<td>1.8 ± 0.1</td>
<td>2.1 ± 0.1</td>
</tr>
<tr>
<td>26k</td>
<td>0.67 ± 0.07</td>
<td>0.64 ± 0.05</td>
</tr>
<tr>
<td>26l</td>
<td>2.2 ± 0.6</td>
<td>2.5 ± 0.2</td>
</tr>
</tbody>
</table>

*Indicates no dose response in the indicated concentration range, making accurate IC₅₀ calculation impossible
Table 2. Association constants for compounds 14c, 16c and 26f and other colchicine-binding site ligands

<table>
<thead>
<tr>
<th>Compound</th>
<th>$K_{assoc}$ ($M^{-1}$) 25°C</th>
</tr>
</thead>
<tbody>
<tr>
<td>Colchicine</td>
<td>$1.16 \times 10^7$ (at 37°C)$^a$</td>
</tr>
<tr>
<td>MTC</td>
<td>$4.7 \times 10^5$ $^b$</td>
</tr>
<tr>
<td>R-PT</td>
<td>$3.2 \times 10^6$ $^c$</td>
</tr>
<tr>
<td>Podofilotoxin</td>
<td>$1.8 \times 10^6$ $^d$</td>
</tr>
<tr>
<td>Nocodazole</td>
<td>$4 \times 10^5$ $^e$</td>
</tr>
<tr>
<td>14c</td>
<td>$(7.1 \pm 1.2) \times 10^6$</td>
</tr>
<tr>
<td>16c</td>
<td>$(9.6 \pm 1.2) \times 10^6$</td>
</tr>
<tr>
<td>26f</td>
<td>$(7.5 \pm 0.76) \times 10^6$</td>
</tr>
</tbody>
</table>

$^a$Data from ref. 46, $^b$Data from ref. 40, $^c$Data from ref. 43, $^d$Data from ref. 45, $^e$Data from ref. 44.
Novel vascular disrupting agents with a cyclohexanedione scaffold identified through a ligand-based virtual screening approach

Maria-Dolores Canela, María-Jesús Pérez-Pérez, Sam Noppen, Gonzalo Sáez-Calvo, J. Fernando Díaz, María-José Camarasa, Sandra Liekens and Eva-María Priego.