New 6- and 7-heterocyclyl-1H-indole derivatives as potent tubulin assembly and cancer cell growth inhibitors

Giuseppe La Regina, a Ruoli Bai, b Antonio Coluccia, a Valentina Naccarato, a Valeria Famiglini, a Marianna Nalli, a Dotomiza Mascl, a Annalisa Verrico, c Paola Rovella, c Carmela Mazzoccoli, c Elena De Pozzo, c Chiara Cavallini, a Claudia Martin, a Stefania Vultaggio, f Giulio Dondio, f Mario Varasi, f Ciro Mercurio, f Ernest Hamel b, Patrizia Lavia, c, Romano Silvestri a, *

a Department of Drug Chemistry and Technologies, Sapienza University of Rome, Laboratory Affiliated to Istituto Pasteur Italia e Fondazione Cenci Bolognetti, Piazzale Aldo Moro 5, I-00185 Roma, Italy
b Screening Technologies Branch, Developmental Therapeutics Program, Division of Cancer Treatment and Diagnosis, Frederick National Laboratory for Cancer Research, National Cancer Institute, National Institutes of Health, Frederick, MD 21702, United States
c Institute of Molecular Biology and Pathology (IBPM), CNR Consiglio Nazionale Delle Ricerche, Via degli Apuli 4, I-00185 Roma, Italy
d Laboratorio di Ricerca Pre-Clinica e Traslazionale, Istituto di Ricovero e Cura a Carattere Scientifico (IRCCS), Centro di Riferimento Oncologico Della Basilicata, Via Padre Pio 1, I-85028, Rionero in Vulture, Italy
e Department of Pharmacy, University of Pisa, Via Bonanno Pisano 6, I-56126 Pisa, Italy
f Experimental Therapeutics IFOM-the FIRC Institute of Molecular Oncology Foundation, Via Adamello 16, 20139 Milan, Italy

Abstract
We designed new 3-arythio- and 3-aryloyl-1H-indole derivatives 3e22 bearing a heterocyclic ring at position 5, 6 or 7 of the indole nucleus. The 6- and 7-heterocyclyl-1H-indoles showed potent inhibition of tubulin polymerization, binding of colchicine to tubulin and growth of MCF-7 cancer cells. Compounds 13 and 19 inhibited a panel of cancer cells and the NCI/ADR-RES multidrug resistant cell line at low nanomolar concentrations. Compound 13 at 50 nM induced 77% G2/M arrest in HeLa cells, and at 20 nM caused 50% stable arrest of mitosis. As an inhibitor of HepG2 cells (IC50 ¼ 20 nM), 13 was 4-fold superior to 19. Compound 13 was a potent inhibitor of the human U87MG glioblastoma cells at nanomolar concentrations, being nearly one order of magnitude superior to previously reported arylthioindoles. The present results highlight 13 as a robust scaffold for the design of new anticancer agents.

1. Introduction
Microtubules (MTs) are hollow, cylindrical, filamentous structures made of dimerized α- and β-tubulin subunits and are characterized by a highly dynamic equilibrium involving polymerization, where tubulin dimers bind non-covalently to αMT, and depolymerization to tubulin dimers [1]. MTs regulate key cellular functions, such as cell growth and division, intracellular trafficking, preservation of the architecture of the cell, and motility. Due to these key roles, disrupting the dynamic equilibrium of MTs at either the tubulin assembly or MT disassembly level, results in a fatal cellular event. Cancer cells are characterized by a high rate of cell division: hence, the strategy of inducing cell death through an interference with the dynamics of MTs has proved successful for the design of effective antitumor drugs [2, 6].

MT binding agents fall into two main groups: (i) inhibitors of tubulin polymerization, including colchicine [7, 8], combretastatin A-4 (CSA4) [9] (Chart 1), vincristine (VCR), vinorelbine (VRB) and vinblastine (VBL); and (ii) MT stabilizers, including taxoids and epothilones. The stabilizers stimulate MT polymerization and stabilization at high concentrations, whereas at lower concentrations paclitaxel (PTX) inhibits MT dynamics with little effect on the proportion of tubulin in polymer [10]. Taxoids and epothilones bind at a luminal site on the b-subunit [11, 12] following entry into the MT through pores in its wall [13] that are shaped by various tubulin subunits on the MT surface. Some evidence indicates a transient binding of MT stabilizers at a specific pore site.

Despite considerable clinical successes [14, 15], the anticancer therapies based on tubulin binding agents still have limitations perhaps due to multi-drug resistance (MDR), toxicity and unwanted side effects [16, 17]. Therefore, there is a quest for new effective MT inhibitors with fewer side effects to become components of improved anticancer treatments [18, 19]. Currently, there are no US Food and Drug Administration approved anticancer drugs on the market binding at the colchicine site [20]. Compared to other tubulin agents, these compounds (i) have shown less transporter mediated drug resistance in preclinical studies [21], (ii) are minimally affected by the overexpression of bIII-tubulin [22], and (iii) have generally better water solubility [9]. Improvements of activity of colchicine site binding agents by replacement or introduction of heterocyclic ring(s) to the parent scaffold have been reported [16, 23, 24]. Accordingly, as tubulin polymerization inhibitors, 2-heterocyclyl-3-arythio-1H-indole (2-HATI) derivatives were more effective than the corresponding 2-ary-3-arythio-1H-indole counterparts in both the sulfur and aroyl series [25]. For example, 2-(thiophen-2-yl)-3-[(30,40,50-trimethoxyphenyl)thio]-1H-indole (1) inhibited tubulin assembly with an IC50 of 0.74 mM and MCF-7 cancer cell growth with an IC50 of 39 nM [26]. Compound 1 was superior to VRB, VBL, and PTX as an inhibitor of the P-glycoprotein (Pgp) overexpressing NCI/ adriamycin-resistant (ADR-RES) MDR cell line and showed satisfactory metabolic stability. Recently, introduction of halogen or methoxy substituent(s) at positions 4e7 of 3-arythio-1H-indoles (ATIs) bearing the phenyl at position 2 of the indole provided potent tubulin polymerization inhibitors (for example 2) [27].
Herein we report the synthesis of new indole derivatives 3e22 bearing a five-membered heterocyclic ring at position 5, 6 or 7 of the indole ring (Chart 1 and Table 1). Our findings demonstrate that the new derivatives are potent anticancer agents, superior to the corresponding reference derivative 1 [26]. Compounds 13 and 19 inhibited the HT29, HCT116, HepG2, T98G, U87MG, U343G and NC/ADR-RES cell lines with IC50 values in the nanomolar range. Compound 13 was superior to the previously reported compounds [26] as an inhibitor of HepG2 and glioblastoma cells.

2. Chemistry

2.1. Synthetic procedures

 Arylthiindoles 3e7, 9, 11, 13, 15, 19 and 37 were prepared according to our previously reported venting-while-heating microwave (MW)-assisted procedure [28] by treating the appropriate indole with bis(3,4,5-trimethoxyphenyl)disulfide [29] in the presence of sodium hydride in anhydrous N,N-dimethylformamide (DMF) at 130 °C for 2 min (120 W) (Scheme 1). Reaction of the indole with 3,4,5-trimethoxybenzoyl chloride in the presence of diethylalumimium chloride in dichloromethane at 10 °C for 2 h furnished methanones 8, 10, 12 and 14. Derivatives 16e18, 20e22, 30e36, 35 and 36 were synthesized by a coupling reaction of the appropriate indole with the boronic derivative in the presence of tris(dibenzylideneacetone)dpalladium(0) (Pd2dba3), 2-dicyclohexylphosphino-20,60-dimethoxybiphenyl (SPhos) and potassium phosphate tribasic in 1-butanol at 100 °C for 15 h. A palladium(II) acetate (Pd(OAc)2)-catalyzed reaction of 5-iodo-1H-indole (24) with 2-furanboronic or 2-thienylboronic acid in the presence of tri(o-tolyl)phosphate (P[o-tol]3) and potassium phosphate tribasic in ethanol/toluene at 80 °C for 2 h gave the corresponding heteroaryl indoles 27 and 29, respectively. MW-assisted treatment of indole 24 with 3-furanboronic acid pinacol ester in the presence of Pd(OAc)2 and potassium carbonate in methylpyrrolidone/water at 110 °C for 15 min (200W) furnished 5-(furan-2-yl)-1H-indole (28). Compound 38 was prepared by MW-assisted Friedel-Crafts reaction of 7-ido-1H-indole (26) with 3,4,5-trimethoxybenzoyl chloride in the presence of anhydrous aluminum chloride in 1,2-dichloroethane at 110 °C for 2 min (150 W).

3. Results and discussion

3.1. Inhibition of tubulin polymerization

We synthesized arylthiindole and arylindole derivatives 3e22 to explore the effects of five-membered heterocyclic rings at position 5, 6 or 7 of the indole nucleus (Table 1). Ten new derivatives (3, 7, 8, 10e15, and 19) inhibited tubulin polymerization with IC50 values at submicromolar concentrations, six compounds (4e6, 16, 17 and 21) were in the 1.0e2.0 mM range, as compared with colchicine (IC50 ¼ 3.2 mM) and CSA4 (IC50 ¼ 1.0 mM). With the exception of 9, the most potent tubulin polymerization inhibitors (7, 8 and 10e14) were characterized by the presence of the heterocyclic ring at position 6 of the indole; among others, compound 3, with the heterocyclic ring at position 7, showed IC50 values < 1.0 mM. As tubulin assembly inhibitors, the compounds bearing the furan-2-yl or the thiophen-2-yl ring were generally superior to the corresponding furan-3-yl/thiophen-3-yl derivatives (compare 3 with 4, 5 with 6, 7 with 9, 15 with 17, and 19 with 21). Replacement of the sulfur bridging atom with the carbonyl group also provided potent tubulin assembly inhibitors. The presence of the heterocycle at position 7 of the indole conferred significant differences of inhibition of tubulin polymerization among arylthiindole/arylindole derivatives (compare 15 with 16, 17 with 18, 19 with 20, and 21 with 22), while compounds 7e14 with the heterocycle 6 were almost equivalent, except 9 and 10, with 12 (IC50 ¼ 0.38 mM) being the most potent tubulin polymerization inhibitor among the new compounds.

3.2. MCF-7 breast cancer cell growth

Several new indole derivatives inhibited the growth of human MCF-7 nonmetastatic breast cancer epithelial cells with IC50 values at nanomolar concentrations (Table 1). Structure-activity relationship (SAR) analysis show three distinct groups: (i) introduction of the heterocyclic ring at position 5 of the indole had a relatively weak effect on inhibition of MCF-7 cell growth, with IC50 values ranging from 70 (3) to 200 nM (4, 6); (ii) compounds 7e14, bearing the heterocyclic ring at position 6 of the indole, all were highly potent MCF-7 cell growth inhibitors, with IC50 values ranging from 4.3 (10) to 18 (14) nM, with five compounds (8 and 10e13) having single digit nanomolar IC50 values; (iii) the indoles with the heterocycle at position 7 showed superior activity: the 7-heterocyclyl-3-arythio-1H-indoles 15, 17, 19 and 21 (IC50s ranging from 15 to 38 nM) were more than one order of magnitude superior to the corresponding aroyl derivatives 16, 18, 20 and 22.

A SAR summary of tubulin polymerization inhibition and inhibition of MCF-7 cell growth of compounds 3e22 is depicted in Chart 2. It should be noted that the SARs differ from those of the 2-phenyl-1H-indoles [27]. Compounds 3e6 with the heterocyclic ring at position 5 showed moderate inhibition of MCF-7 cell growth; compounds 7e14, bearing the heterocycle at position 6 of the indole, all were potent MCF-7 cell growth inhibitors and showed comparable IC50 values; derivatives 15, 17, 19 and 22, with the heterocyclic ring at position 7 and the sulfur atom bridging the two aromatic ring systems, were substantially superior to the corresponding aroyl compounds (16, 18, 20 and 22).

3.3. Inhibition of the binding of [3H]colchicine to tubulin

Compounds 3e22 were also examined for potential inhibition of the binding of [3H]colchicine to tubulin (Table 1). Thirteen compounds, 3, 5, 7e17, and 19 yielded >75% inhibition of the binding reaction. Interestingly, all the indoles with the heterocycle at position 7 were strong inhibitors of [3H]colchicine binding, with 13 (97% inhibition) nearly as
potent as CSA4 (98% inhibition). In general, good correlation between [3H]colchicine binding and MCF-7 cell growth inhibition was observed: compounds which inhibited [3H]colchicine binding >90%, inhibited the growth of the MCF-7 cancer cell with IC50s _ 15 nM; compounds with 80e90% inhibition of [3H]colchicine binding inhibited the growth of the MCF-7 cancer cell line with IC50s in the range of 24e70 nM; except for 21, compounds with 70e80% inhibition of [3H]colchicine binding inhibited the growth of the MCF-7 cancer cells with IC50s in the range of 150e350 nM.

3.4. Molecular modeling studies

Compounds 6, 13 and 19, representative members of the heterocyclic substitutions at position 5-, 6- or 7- of the indole nucleus, were docked into the tubulin-DAMA-colchicine complex (PDB code 1SAA) [8] by following our previously reported procedure [30]. The docking studies were also performed for five newly available tubulin crystal structures [31]. The docking results in the different tubulin structures revealed a consistent binding mode for compounds bearing the heterocyclic ring at position 5-, 6- or 7- of the indole: (i) the 3,4,5-trimethoxyphenyl moiety formed an H-bond with the Cys241b side chain and hydrophobic contacts with Leu248b and Leu255b; (ii) the indole established hydrophobic contacts with Asn258b and Met259b; (iii) the heterocyclic ring of 6, 13 and 19 at position 5, 6 or 7 of the indole, respectively, arranged hydrophobic interactions with Met259b, Lys353b, Ala180a and Val181 in the same cleft of the colchicine site (Figs. 1 and 2).

3.5. Inhibition of growth of glioblastoma T98G, U87MG and U343MG cell lines

Treatments of T98G (human glioblastoma), U87MG (human glioblastoma-astrocytoma) and U343MG (human glioblastoma/astrocytoma) cells with increasing concentrations of 13 and 19 significantly inhibited cell growth in a dose-dependent manner (Table 2 and Fig. 1S-D3D, supplementary data). The IC50 values were calculated taking into account the relative cellular doubling times [32,33] of 48 h for the T98G and [U87MG cells and of 72 h for the U343MG cells. Compound 13 potently inhibited the glioblastoma cells at nanomolar concentrations. Arylthioindoles with the imidazole-1-y1 or pyridin-4-y1 heterocyclic ring at position 2 of the indole were weaker inhibitors of glioblastoma U87MG cells than were 13 and 19, even though such compounds exhibited potent inhibition of the MCF-7 cancer cell growth [26].

3.6. Inhibition of HT29, HCT116 and HepG2 cancer cell growth

Compounds 13 and 19 were evaluated as growth inhibitors of the human cell lines HT29 (colon adenocarcinoma), HCT116 (colon carcinoma), and HepG2 (hepatocellular carcinoma) using PTX as reference compound (Table 2). Compounds 13 and 19 showed strong inhibition of the HT29 and HCT116 cell lines. As inhibitors of the HepG2 cells, 13 and 19 were more effective than PTX (IC50 ¼ 2660 nM).

3.7. Inhibition of MV4-11, THP-1, A-549 and PC3 cell lines

Compounds 13 and 19 were evaluated as inhibitors of MV4-11 (leukemia, acute myeloid), THP1 (leukemia, acute monocytic), A549 (lung carcinoma) and PC3 (prostate carcinoma) cancer cells. Compound 13 inhibited MV4-11, THP-1, A-549 and PC3 cells at single digit nanomolar concentration, while 19 was consistently less active (Table 2).

3.8. MDR cell lines

Compounds 13 and 19, representative of the 6- and 7- heterocyclicyl series, respectively, were compared with CSA4, VRB, VLB and PTX in the ovarian carcinoma cell lines OVCAR-8 and its cognate Pgp overexpressing line NCI/ADR-RES (Table 3). In contrast to CSA4, the standard agents VRB, VLB and PTX showed weak inhibition of the MDR line NCI/ADR-RES. The IC50 values of 13 and 19 for the MCF-7 cells, repeated in this study, resembled those obtained in the studies summarized in Table 1.

3.9. Compounds 13 and 19 induce cell cycle arrest in the G2/M phase in HeLa cells

We compared cell cycle progression in HeLa cells exposed to 20, 50 and 100 nM 13 and 19, using 20 nM VBL as a reference compound (a concentration that effectively arrests cells in mitosis [27]) and DMSO as a control. Cells were treated for 24 h to cover the average duration of an entire cell cycle, then harvested, incubated with propidium iodide (PI) and analyzed for their genomic content by FACS analysis. Representative cell cycle profiles are shown in Fig. 3A, left panel; data from three experiments are quantified in the graph in Fig. 3A, right panel. Both molecules inhibited cell cycle progression in a dose-dependent manner but with different effectiveness: treatment with 100 nM 19 yielded a substantial fraction (over 71%) of the cell population in the G2/M phase, whereas lower doses had no significant effect; compound 13 was already partly effective at 20 nM and induced substantial G2/M arrest at 50 nM (77% arrest). Immunofluorescence (IF) analysis of 13 and 19-treated HeLa cell cultures confirmed that treatment with 20 nM 13 (24 h) induced arrest in mitosis in over 50% of all cells, similar to the effect of VBL. The mitotic arrest was stable, and only 15% of the cell population included multinucleated cells, resulting from mitotic “slippage” and interphase re-entry with unsegregated, or randomly segregating, chromosomes. Compound 19 induced mitotic arrest with lower effectiveness than 13: (i) 100 nM 19 arrested 50% of the cell population in mitosis, compared with 20 nM 13 or VBL; (ii) the induction of mitotic arrest by 19 was not fully sustained. The higher accumulation of cells in mitosis observed with 100 nM 19 was accompanied by induction of a relevant fraction (around 32%) of multinucleated cells: this suggests that cells that reached mitosis arrested only transiently, and eventually resumed mitotic progression with an inefficient mitotic apparatus, generating multinucleated cellular offspring (Fig. 3B). We wondered whether cell cycle arrest by 13 and 19 was paralleled by cell death induction, as is the case with classical MTtargeting drugs. Cultures treated for 24 h were
incubated with annexin V to identify cells with damaged plasma membranes committed to death. Both compounds activated cell death parallel to arresting cell cycle progression; again, 19 was effective when used at 100 nM, whereas 20 nM 13 was already partly effective and reached the same death-inducing capacity as VBL at the 50 nM dose (Fig. 3C). The tubulin inhibitory effects of 13 and 19 at the cytological level were evaluated by IF analysis (Fig. 4). These assays provided a visual demonstration of the differential effectiveness of the two compounds. Compounds at 20 or 50 nM induced a mixture of apparently normal, or fragmented or multipolar spindles in variable proportions (examples are shown in Fig. 4A, phenotypes are quantified in Fig. 4C); 100 nM 19 largely induced small formations of radially arranged short microtubules that failed to elongate (labeled as “MT asters”), Fig. 4A, central row). Compound 13 had a more pronounced effect at all tested concentrations, and no cell showed a normal or only weakly affected spindle. Even with 13 at 20 nM, most mitotic cells displayed multipolar (Fig. 4B, top row) or fragmented spindles (Fig. 4B, central row); at higher concentration, 50 nM or above, 13 inhibited MT formation altogether, with sparse tubulin foci in virtually all treated cells (Fig. 4B, bottom row). Representative phenotypes are shown in Fig. 4A and B and quantified in Fig. 4C.

In summary, both 13 and 19 arrested cells in mitosis, with formation of a defective mitotic apparatus, yet the two compounds showed differential effectiveness. Compound 13 induced the inhibition of MT polymerization to a greater extent than 19, and it showed a full inhibitory effect on tubulin polymerization in cells at 50 nM, comparable to the effect of VBL. Compound 19 induced milder inhibition: it essentially affected the mitotic spindle structural organization but did not fully inhibit tubulin polymerization in cells below 100 nM. The parallel observation that 13 was a more effective inducer of cell death than 19 correlated with cellular data on microtubule disruption and with inhibition of [3H]colchicine binding.

3.10. Effects of 13 or 19 on viability of HepG2 cells
HepG2 cells were treated with different concentrations of 13 or 19 and analyzed by the MTT assay. Treatment with compound 19 for 48 h caused a dose-dependent decrease of cell viability: at 80 nM nearly 30% of cells displayed reduced viability as compared to untreated cells (Fig. 5D, panel A, supplementary data). The dose-dependent decrease of cell viability on treatment with 13 became significant at 20 nM, nearly 40% of cells were affected (Fig. 5D, panel B, supplementary data). These experiments indicated 80 nM 19 and 20 nM 13 are optimal concentrations to induce significant modifications in cell viability but, at the same time, preserving a sufficient proportion of the cell population to permit further analysis. At such concentrations, 13 and 19 caused a marked impairment of HepG2 cell growth after a 48 h treatment. The cell cycle blocker p21cip1/Waf1 was significantly up-regulated in HepG2 treated cells at these doses. After 80 nM 19 (Fig. 5D, panel A, supplementary data) or 20 nM 13 (Fig. 5D, panel B, supplementary data) treatments, the transcript levels of p21cip1/Waf1, assessed by qRT-PCR, were increased by 1.5- and 3.9-fold, respectively, as compared with untreated cells.

3.11. Evaluation of the drug-like properties of 13 and 19
Drug like properties of compounds 13 and 19 were predicted through the most common descriptors of drug-likeness (Table 4). Oral absorption according to Lipinski’s rule of five [34] and Veber’s rule [35] was estimated by FAF drug server [36]. We referred to the 3/75 rule (logP > 3 and topological PSA < 75 Å2) [37] to estimate compound toxicity. Derivatives 13 and 19 properly fitted with both Lipinsky and Veber’s rules, suggesting a potential good absorption after oral administration. Furthermore, a low likelihood of in vivo toxicological outcome was inferred by the 3/75 rule (Fig. 5).

4. Conclusions
We designed new 3-arylthio- and 3-aryloxy-1H-indole derivatives 3-22 bearing a heterocyclic ring at position 5, 6 or 7 of the indole nucleus. Ten new derivatives inhibited polymerization of purified tubulin with IC50 values at submicromolar concentrations. Seven of the new indole derivatives inhibited the growth of MCF-7 cells with IC50 values 10 nM. Inhibition of cell growth showed good correlation with inhibition of [3H]colchicine binding to tubulin. Two representative highly potent members of the 6- and 7-heterocyclyl-1H-indoles, 13 and 19, inhibited an extensive panel of cancer cell lines, including the Pgp overexpressing NCI/ADR-RES cell line, with nanomolar IC50s. In cell cycle analysis, compound 13 became effective at 20 nM and induced 77% G2/M arrest at 50 nM, whereas 19 accumulated 71% cells in the G2/M phase at 100 nM. Compound 13 extensively inhibited cellular tubulin polymerization at 50 nM, and its activity was comparable to that of VBL. Compound 19 at 100 nM affected mitotic spindle structural organization but did not fully inhibit cellular tubulin polymerization. As inhibitors of HepG2 cells, 13 and 19 (IC50 values of 20 nM and 80 nM, respectively) were distinctly superior to PTX. Compound 13 was also a potent inhibitor of the U87MG glioblastoma cell line at nanomolar concentrations [41] being almost one order of magnitude more active than the previously reported arylthiinones with the imidazole-1-yl or pyridin-4-yl ring at position 2 of the indole nucleus [26]. Analysis of the gene expression levels of p21cip1/Waf1, the cell cycle blocker, indicated that at sub-cytotoxic concentrations 13 caused a decrease in cell growth with strong up-regulation of p21cip1/Waf1 [42].

In conclusion, these new indoles are potent inhibitors of tubulin polymerization and cancer cell growth, including human liver carcinoma and glioblastoma cells. The introduction of the heterocyclic ring at position 5, 6 or 7 of the 3-(30,40,50-trimethoxyaryltio)-1H-indole and 3-(30,40,50-trimethoxyaroyl)-1H-indole scaffold was a productive strategy for the
5. Experimental protocols

5.1. Chemistry

All reagents and solvents were handled according to the material safety data sheet of the supplier and were used as purchased without further purification. 5-Chloro-1H-indole (23), 5-iodo-1Hindole (24), 6-bromo-1H-indole (25), 7-iodo-1H-indole (26), 6-(thiophen-2-yl)-1H-indole (33) and 6-(thiophen-3-yl)-1H-indole (34) were commercially available. MW-assisted reactions were performed on a CEM Discover SP single-mode reactor equipped with an Explorer 72 autosampler, controlling the instrument settings by PC-running CEM Synergy 1.60 software. Closed vessel experiments were carried out in capped MW-dedicated vials (10 mL) with a cylindrical stirring bar (length 8 mm, diameter 3 mm). Stirring, temperature, irradiation power, maximum pressure (Pmax), pressure set point, times at set point, delta pressure, PowerMAX (simultaneous cooling-while-heating), ActiVent (simultaneous venting-while-heating), and ramp and hold times were set as indicated. Reaction temperature was monitored by an external CEM fiber optic temperature sensor. After completion of the reaction, the mixture was cooled to 25 _C via air-jet cooling. Organic solutions were dried over anhydrous sodium sulfate. Evaporation of solvents was carried out on a Büchi Rotavapor R-210 equipped with a BÜCHI V-850 vacuum controller and a BÜCHI V-700 vacuum pump. Column chromatography was performed on columns packed with silica gel from Macherey-Nagel (70e230 mesh). Silica gel thin layer chromatography (TLC) cards from Macherey-Nagel (silica gel precoated aluminum cards with fluorescent indicator visualizable at 254 nm) were used for TLC. Developed plates were visualized with a Spectroline ENF 260C/FE UV apparatus. Melting points (mp) were determined on a Stuart Scientific SMP1 apparatus and are uncorrected. Infrared (IR) spectra were recorded on a PerkinElmer Spectrum 100 FT-IR spectrophotometer equipped with a universal attenuated total reflectance accessory and IR data acquired and processed by PerkinElmer Spectrum 10.03.00.0069 software. Band position and absorption ranges are given in cm^-1. Proton nuclear magnetic resonance (1H NMR) spectra were recorded with a Varian Mercury (300 MHz) or a Bruker Avance (400 MHz) spectrometer in the indicated solvent, and the corresponding fid files were processed by MestreLab Research SL MestreReNova 6.2.1e769 software. Chemical shifts are expressed in ppm (ppm) from tetramethylsilane. Mass spectra were recorded on a Bruker Daltonics MicroTOF LC/MS mass spectrometer equipped with a positive ion ESI source. Compound purity was checked by high pressure liquid chromatography (HPLC). Purity of tested compounds was found to be >95%. The HPLC system used (Thermo Fisher Scientific Inc. Dionex UltiMate 3000) consisted of an SR-3000 solvent rack, a LPG-3400SD quaternary analytical pump, a TCC-3000SD column compartment, a DAD-3000 diode array detector, and an analytical manual injection valve with a 20 ml loop. Samples were dissolved in acetonitrile (1 mg/mL). HPLC analysis was performed by using a Thermo Fisher Scientific Inc. Acclaim 120C18 column (5 mm, 4.6 mm, 250 mm) at 25 ± 1 _C with an appropriate solvent gradient (acetonitrile/water), flow rate of 1.0 mL/min and signal detector at 206, 230, 254 and 365 nm. Chromatographic data were acquired and processed by Thermo Fisher Scientific Inc. Chromelon 6.80 SR15 Build 4656 software.

5.1.1. General procedure for the synthesis of derivatives 3e-7, 9, 11, 13, 15, 19 and 37

Example. 5-(Furan-2-yl)-3-((3,4,5-trimethoxyphenyl)thio)-1H-indole (3)

A mixture of indole 27 (183 mg, 1 mmol), bis(3,4,5-trimethoxyphenyl)disulfide [29] (438 mg, 1.1 mmol), and sodium hydride (88 mg, 2.2 mmol; 60% in mineral oil) in anhydrous DMF (3 mL) was placed into the MW cavity (closed vessel mode, Pmax ¼ 250 psi). Starting MW irradiation of 120W was used, the temperature being ramped from 25 to 130 _C, while rapidly stirring and venting (pressure set point: 100 psi; times at set point: 100; delta pressure: 20 psi). Once 130 _C was reached, taking about 1 min, the reaction mixture was held at this temperature for 2 min. The mixture was diluted with water and extracted with ethyl acetate. Evaporation of solvents was carried out on a Büchi Rotavapor R-210 equipped with a BÜCHI V-850 vacuum controller and a BÜCHI V-700 vacuum pump. Column chromatography was performed on columns packed with silica gel from Macherey-Nagel (70e230 mesh). Silica gel thin layer chromatography (TLC) cards from Macherey-Nagel (silica gel precoated aluminum cards with fluorescent indicator visualizable at 254 nm) were used for TLC. Developed plates were visualized with a Spectroline ENF 260C/FE UV apparatus. Melting points (mp) were determined on a Stuart Scientific SMP1 apparatus and are uncorrected. Infrared (IR) spectra were recorded on a PerkinElmer Spectrum 100 FT-IR spectrophotometer equipped with a universal attenuated total reflectance accessory and IR data acquired and processed by PerkinElmer Spectrum 10.03.00.0069 software. Band position and absorption ranges are given in cm^-1. Proton nuclear magnetic resonance (1H NMR) spectra were recorded with a Varian Mercury (300 MHz) or a Bruker Avance (400 MHz) spectrometer in the indicated solvent, and the corresponding fid files were processed by MestreLab Research SL MestreReNova 6.2.1e769 software. Chemical shifts are expressed in ppm (ppm) from tetramethylsilane. Mass spectra were recorded on a Bruker Daltonics MicroTOF LC/MS mass spectrometer equipped with a positive ion ESI source. Compound purity was checked by high pressure liquid chromatography (HPLC). Purity of tested compounds was found to be >95%. The HPLC system used (Thermo Fisher Scientific Inc. Dionex UltiMate 3000) consisted of an SR-3000 solvent rack, a LPG-3400SD quaternary analytical pump, a TCC-3000SD column compartment, a DAD-3000 diode array detector, and an analytical manual injection valve with a 20 ml loop. Samples were dissolved in acetonitrile (1 mg/mL). HPLC analysis was performed by using a Thermo Fisher Scientific Inc. Acclaim 120C18 column (5 mm, 4.6 mm, 250 mm) at 25 ± 1 _C with an appropriate solvent gradient (acetonitrile/water), flow rate of 1.0 mL/min and signal detector at 206, 230, 254 and 365 nm. Chromatographic data were acquired and processed by Thermo Fisher Scientific Inc. Chromelon 6.80 SR15 Build 4656 software.
5.1.4. 5-(Thiophen-3-yl)-3-((3,4,5-trimethoxyphenyl)thio)-1H-indole (6)
Synthesized as 3 starting from 30. Yield 40%, mp 1161612_C (from ethanol). 1H NMR (DMSO-d6, 400 MHz): d 3.57 (s, 3H), 3.58 (s, 6H), 6.44 (s, 2H), 7.49e7.51 (m, 1H), 7.53 (s, 1H), 7.55 (d, J ¥ 1.3 Hz, 1H), 7.59e7.62 (m, 1H), 7.72e7.74 (m, 2H), 7.79 (d, J ¥ 2.4 Hz, 1H), 11.71 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3448 cm_1. MS (ESI): 398.5 (M+H). C21H19NO3S2 (397.51).

5.1.5. 6-(Furan-2-yl)-3-((3,4,5-trimethoxyphenyl)thio)-1H-indole (7)
Synthesized as 3 starting from 11. Yield 4%, slurry. 1H NMR (DMSO-d6, 400 MHz): d 3.57 (s, 3H), 3.85 (s, 6H), 6.41 (s, 2H), 7.10e7.13 (m, 1H), 7.41e7.43 (m, 1H), 7.44e7.48 (m, 3H), 7.70e7.72 (m, 1H), 7.82 (s, 1H), 11.74 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3379 cm_1. MS (ESI): 382.2 (M+H). C21H19NO3S2 (381.45).

5.1.6. 6-(Furan-3-yl)-3-((3,4,5-trimethoxyphenyl)thio)-1H-indole (9)
Synthesized as 3 starting from 32. Yield 4%, slurry. 1H NMR (DMSO-d6, 400 MHz): d 3.57 (s, 3H), 3.85 (s, 6H), 6.40 (s, 2H), 7.45e7.47 (m, 2H), 7.52e7.57 (m, 1H), 7.60e7.64 (m, 1H), 7.76 (s, 1H), 7.79 (s, 1H), 7.77e7.81 (m, 1H), 11.72 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3376 cm_1. MS (ESI): 399.5 (M+H). C21H19NO3S2 (397.51).
5.1.16. General procedure for the synthesis of derivatives 16e18,20e22, 30e32, 35 and 36. (7-(Furan-2-yl)-1H-indol-3-yl) (3,4,5-trimethoxyphenyl) methanone (16)
A mixture of derivative 38 (437 mg, 1 mmol), tris(dibenzylideneacetone) dipalladium(0) (20 mg, 0.022 mmol), 2-dicyclohexylphosphino-20,60-dimethoxybiphenyl (3.2 mg, 0.008 mmol) and potassium phosphate tribasic (435 mg, 2.05 mmol) was degassed for 20 min. 1-Butanol (2.2 mL) and a solution of 2-furanylboronic acid MIDA ester (2-(furan-2-yl)-6-methyl-1,3,6,2-dioxazaborocane-4-dione) (334 mg, 1.5 mmol) in the same solvent (4.4 mL) were added. The reaction mixture was heated at 100 °C for 15 h. After cooling, the mixture was treated dropwise with 1 N HCl and extracted with ethyl acetate. The organic layer was washed with brine, dried and filtered. Evaporation of the solvent gave a residue that was purified by column chromatography (silica gel, n-hexane:acetone ¼ 2:1 as eluent) to give 16 (124 mg, yield 33%), mp 212e224 °C (from ethanol). 1H NMR (DMSO-d6, 400 MHz): d 3.76 (s, 3H), 3.85 (s, 6H), 7.00e7.05 (m, 1H), 7.11 (s, 2H), 7.29 (t, J = 7.9 Hz, 1H), 7.42 (dd, J = 1.1 and 7.4 Hz, 1H), 7.86 (t, J = 1.8 Hz, 1H), 7.97 (s, 1H), 8.21 (dd, J = 1.2 and 8.0 Hz, 1H), 8.25e8.30 (m, 30, 1H), 11.76 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3318 cm_1. MS (ESI): 382.4 (MHþ). C27H20NO6S (381.45).

5.1.17. 7-(Furan-3-yl)-3-(3,4,5-trimethoxyphenyl)thio)-1H-indole (17)
Synthesized as 16 starting from 37 and furan-3-boronic acid pinacol ester. Yield 46%, mp 215e220 °C (from ethanol). 1H NMR (DMSO-d6, 400 MHz): d 3.37 (s, 3H), 3.85 (s, 6H), 7.04 (m, 1H), 7.15 (t, J = 7.7 Hz, 1H), 7.40 (m, 2H), 7.78 (s, 1H), 7.85 (t, J = 1.8 Hz, 1H), 9.41 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3318 cm_1. MS (ESI): 382.4 (MHþ). C27H20NO6S (381.45).

5.1.18. 7-(Furan-3-yl)-1H-indol-3-yl) (3,4,5-trimethoxyphenyl) methanone (18)
Synthesized as 16 starting from 38 and furan-3-boronic acid pinacol ester. Yield 46%, mp 215e220 °C (from ethanol). 1H NMR (DMSO-d6, 400 MHz): d 3.76 (s, 3H), 3.85 (s, 6H), 7.00e7.05 (m, 1H), 7.11 (s, 2H), 7.29 (t, J = 7.9 Hz, 1H), 7.42 (dd, J = 1.1 and 7.4 Hz, 1H), 7.86 (t, J = 1.8 Hz, 1H), 7.97 (s, 1H), 8.21 (dd, J = 1.2 and 8.0 Hz, 1H), 8.25e8.30 (m, 30, 1H), 11.76 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 1568 and 3285 cm_1. MS (ESI): 378.4 (MHþ). C27H20NO6S (377.40).

5.1.19. 7-(Thiophen-2-yl)-1H-indol-3-yl) (3,4,5-trimethoxyphenyl) methanone (20)
Synthesized as 16 starting from 38 and thiophen-3-boric acid pinacol ester. Yield 6%, mp 209e211 °C (from ethanol). 1H NMR (DMSO-d6, 400 MHz): d 3.76 (s, 3H), 3.85 (s, 6H), 7.11 (s, 2H), 7.28e7.33 (m, 2H), 7.38 (dd, J = 1.2 and 7.4 Hz, 1H), 7.56 (dd, J = 1.2 and 3.6 Hz, 1H), 7.68 (dd, J = 1.1 and 5.1 Hz, 1H), 7.95 (s, 1H), 8.25 (dd, J = 1.2 and 7.9 Hz, 1H), 11.94 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3327 cm_1. MS (ESI): 394.4 (MHþ). C27H20NO6S (393.46).

5.1.20. 7-(Thiophen-3-yl)-3-(3,4,5-trimethoxyphenyl)thio)-1Hindole (21)
Synthesized as 16 starting from 38 and thiophen-3-boric acid. Yield 5%, slurry. 1H NMR (DMSO-d6, 400 MHz): d 3.76 (s, 3H), 3.84 (s, 6H), 7.10 (s, 2H), 7.30 (t, J = 7.8 Hz, 1H), 7.40 (dd, J = 1.1 and 7.4 Hz, 1H), 7.52 (dd, J = 1.3 and 5.0 Hz, 1H), 7.70e7.75 (m, 1H), 7.80e7.86 (m, 1H), 7.93 (s, 1H), 8.23 (dd, J = 1.1 and 7.8 Hz, 1H), 11.86 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 1580 and 2925 cm_1. MS (ESI): 394.4 (MHþ). C27H19NO4S (393.46).

5.1.22. 5-(Thiophen-3-yl)-1H-indole (30)
Synthesized as 16 starting from 23 and thiophen-3-boric acid. Yield 40%, mp 80e83 °C (from n-hexane), lit. 80e82 °C [43].

5.1.23. 6-(Furan-2-yl)-1H-indole (31)
Synthesized as 16 starting from 25 and furan-2-boronic acid MIDA ester. Yield 19%, mp 90e96 °C (from n-hexane). 1H NMR (DMSO-d6, 400 MHz): d 6.38e6.42 (m, 1H), 6.50e6.56 (m, 1H), 6.81 (dd, J = 0.8 and 3.4 Hz, 1H), 7.32e7.36 (m, 2H), 7.55 (d, J = 8.3 Hz, 1H), 7.69e7.71 (m, 2H), 11.17 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3388 cm_1.

5.1.24. 6-(Furan-3-yl)-1H-indole (32)
Synthesized as 16 starting from 25 and furan-3-boronic acid MIDA ester. Yield 7%, mp 155e158 °C (from n-hexane). 1H NMR (DMSO-d6, 400 MHz): d 6.37e6.40 (m, 1H), 6.90e6.95 (m, 1H), 7.25 (dd, J = 1.6 and 8.2 Hz, 1H), 7.30e7.32 (m, 1H), 7.50e7.53 (m, 2H), 7.77 (t, J = 1.8 Hz, 1H), 8.08e8.12 (m, 1H), 11.07 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 3384 cm_1.

5.1.25. 7-(Furan-2-yl)-1H-indole (35)
Synthesized as 16 starting from 26 and 2-furanylboronic acid MIDA ester. Yield 21% as an oil [44].

5.1.26. 7-(Thiophen-2-yl)-1H-indole (36)
Synthesized as 16 starting from 26 and 2-thienylboronic acid. Yield 67% as an oil [45].

5.1.27. General procedure for the synthesis of derivatives 27 and 29.
Example. 5-(Furan-2-yl)-1H-indole (27)
A mixture of 24 (243 mg, 1 mmol), 2-furanboronic acid (223 mg, 2 mmol), palladium(ii) acetate (11 mg, 0.05 mmol), tri(o-tolyl) phosphine (30 mg g, 0.1 mmol), potassium phosphate tribasic (745 mg, 3.5 mmol) in ethanol (10 mL) and toluene (5 mL) was heated at 80 °C for 2 h. After cooling, the reaction mixture was diluted with a saturated aqueous solution of sodium hydrogen carbonate and extracted with ethyl acetate. The organic layer was washed with brine, dried and filtered. Evaporation of the solvent gave a residue that was purified by column chromatography (silica gel, n-hexane:ethyl acetate 5:1 as eluent) to give 27 (159 mg, yield 87%), mp 108e110 _C (from ethanol), lit. 110e111 _C.

5.1.28. 5-(Thiophen-2-yl)-1H-indole (29)
Synthesized as 27 starting from 2-thienylboronic acid. Yield 73%, mp 50e51 _C (from n-hexane), lit. 53e54 _C [46].

5.1.29. Synthesis of 5-(furan-2-yl)-1H-indole (28)
A mixture of 24 (243 mg, 1 mmol), 3-furanboronic acid pinacol ester (253 mg, 1.3 mmol), potassium carbonate (179 mg, 1.3 mmol) in methylpyrrolidone (2 mL) and water (0.15 mL) was degassed for 15 min. Palladium(ii) acetate (30 mg, 0.13 mmol) was added, and the reaction mixture was placed into the MW cavity (closed vessel mode, Pmax ¼ 250 psi). Starting MW irradiation of 200W was used, the temperature being ramped from 25 to 110 _C. Once this was reached, taking around 2 min, the mixture was held at this temperature for 15 min with rapid stirring. The reaction mixture was diluted with water and extracted with ethyl acetate. The organic layer was washed with brine, dried and filtered. Evaporation of the solvent gave a residue that was purified by column chromatography (silica gel, n-hexane:ethyl acetate 5:1 as eluent) to give 28 (82 mg, yield 45%) mp 89e90 _C (from n-hexane), lit. 89e91 _C [47].

5.1.30. Synthesis of 7-iodo-1H-indol-3-yl) (3,4,5-trimethoxyphenyl) methanone (38)
A mixture of 26 (243 mg, 1 mmol), 3,4,5-trimethoxybenzyl chloride (230 mg, 1 mmol) and anhydrous aluminum chloride (133 mg, 1 mmol) in 1,2-dichloroethane (2 mL) was placed into the MW cavity (closed vessel mode, Pmax ¼ 250 psi). A starting MW irradiation of 150W was used, the temperature being ramped from 25 to 110 _C with rapid stirring. Once 110 _C was reached, taking around 1 min, the reaction mixture was heated at this temperature for 2 min, diluted with water, treated with 1 N HCl and extracted with chloroform. The organic layer was washed with brine, dried and filtered. Evaporation of the solvent gave a residue that was purified by column chromatography (silica gel, n-hexane:ethyl acetate 1:1 as eluent) to give 38 (223 mg, yield 51%, mp 148e153 _C (from ethanol). 1H NMR (DMSO-d6, 400 MHz): δ 3.76 (s, 3H), 3.85 (s, 6H), 7.04 (t, J ¼ 7.6 Hz, 1H), 7.09 (s, 2H), 7.66 (dd, J ¼ 1.0 and 7.6 Hz, 1H), 7.98 (s, 1H), 8.23 (dd, J ¼ 1.0 and 7.96 Hz, 1H), 11.86 ppm (br s, disappeared after treatment with D2O, 1H). IR: n 1577 and 3112 cm^-1.

5.2. Molecular modeling studies
All molecular modeling studies were performed on a MacPro dual 2.66 GHz Xeon running Ubuntu 14LTS. The tubulin structures were downloaded from the PDB data bank (http://www.rcsb.org/). PDB code: 1SAO [30], 4O2A [48], 4O2B [48], 5CA0 [8], 5CB4 [8], 5LYJ [49] and 3HKC [50]. Ligand structures were prepared with Maestro [51]. Proteins were prepared by the protein preparation wizard [52] of Maestro. The docking simulations were performed using Plants [53] and Autodock [54]. Images shown in the manuscript were prepared with Pymol [55].

5.3. Biological assays
5.3.1. Tubulin assembly
The reaction mixtures contained 0.8M monosodium glutamate (pH 6.6 with HCl in a 2M stock solution), 10 mM tubulin, 4% (v/v) DMSO, and varying concentrations of drug. Following a 15 min preincubation at 30 C, samples were chilled on ice, GTP to 0.4mM was added, and turbidity development was followed at 350nm in a temperature-controlled recording spectrophotometer for 20 min at 30 C. The extent of reaction was measured. Full experimental details were previously reported [56].

5.3.2. [3H]Colchicine binding assay
The reaction mixtures contained 1.0 mM tubulin, 5.0 mM [3H] colchicine, and 5.0 mM inhibitor and were incubated for 10 min at 37 C. Complete details were described previously [57].

5.3.3. Cell cultures
Cell lines were obtained from the American Type Culture Collection (ATCC), unless otherwise specified. MCF-7 breast carcinoma, OVCAR-8, and NCI/ADR-RES cells were obtained from the National Cancer Institute drug screening laboratory. U87MG and U343MG cell lines were obtained from the National Institute for Cancer Research of Genoa (Italy) and Cell Lines Service GmbH (Germany), respectively. All cell lines, except as indicated, were grown in Dulbecco’s modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS), 20mMHEPES,100 U/mL penicillin, 100 mg/mL streptomycin, and 1% L-glutamine; specific requirements include the addition of glucose (4.5 g/L for HT29 and HCT116 cells; 1 g/L for HepG2). Cell lines were cultured at 37 C in 5% CO2/95% air in a humidified incubator. Treatments were initiated 24 h after cell seeding using compound 13 or 19 diluted in 0.1% DMSO, the indicated reference compound, or 0.1% DMSO vehicle, for 24e72 h as indicated. T98G and U87MG cells were cultured in RPMI medium and minimum essential medium Eagle, respectively, supplemented with 10% FBS, 2mM L-glutamine, 100 U/mL
penicillin, 100 mg/mL streptomycin and 1% non-essential amino acids at 37 C in 5% CO2. The U343MG cells were cultured in minimum essential medium Eagle with 2mM L-glutamine and Earle's BSS adjusted to contain 1.5 mg/mL sodium bicarbonate and supplemented with 10% FBS, 100 U/mL penicillin, 100 mg/mL streptomycin, 1% non-essential amino acids and 1.0mM sodium pyruvate at 37 C in 5% CO2.

5.3.4. Cell viability assays
The methodology for the evaluation of the growth of human MCF-7 breast carcinoma, OVCAR-8, and NCI/ADR-RES cells, obtained from the National Cancer Institute drug screening laboratory, was previously described, except that cells were grown for 96 h for IC50 determinations [58]. Cell viability of HT29, HCT116 and HepG2 cells was determined using the MTT colorimetric assay. HT29, HepG2 and HCT116 cells were seeded into 24-well plates to a density of 15 x 10^3/mL in each well. After 24 h of growth to allow attachment of cells to the wells, test compounds were added at 20-320 nM. After 48 h of growth and removal of the culture medium, 500 mL/well of PBS containing 500 mM MTT was added. Cell cultures were further incubated at 37 C for 2 h in the dark. The solutions were then gently aspirated from each well, and the formazan crystals within the cells were dissolved in propan-2-ol and 0.04 N HCl (200 mL). Optical densities were read at 550 nm using a Multiskan Spectrum Thermo Electron Corporation reader. The results were expressed as % relative to vehicle-treated control (0.1% DMSO), and IC50 values were calculated by nonlinear regression analysis (GraphPad Prism statistics software). The effect of the treatment with compounds 13 or 19 on the T98G, U87MG and U343MG cell lines was estimated using the colorimetric MTS conversion assay, as previously reported.27 After compound incubations, the MTS reagent was added, and the absorbance at 590 nm was measured by a microplate reader (Wallac, Victor 2, 1420 Multilabel Counter, PerkinElmer). The percentage of proliferating cells after compound exposure was calculated with respect to control cells (100%). The effect of treatment with compounds 13 or 19 on MV4-11, THP-1, A-S49 and PC3 cell was determined by CellTiter-Fluor™ (Promega cod. G6082). The cells were seeded in 96 multwell plates at the indicated densities (MV-4: 5000 cells/50 mL/ well; THP-1: 2000 cells/50 mL/well; A549: 3500 cells/50 mL/well; MDA-MB-231: 3500 cells/50 mL/well; PC-3: 2000 cells/50 mL/well). 24 h after plating, compound 13 or 19 was added to the cells, and the effect on cell proliferation was determined after 72 h by the CellTiter-Fluor™ assay. After adding the reagents to the cells plates, the cells were incubated for 90 min at 37 C, and the fluorescent signal was read using a TECAN reader. The IC50 and GI50 results were obtained by analysis with GraphPad Prism and Assay Explorer software.

5.3.5. Statistical analyses
Graph-Pad Prism 5 software (Graph-Pad Software Inc, San Diego, CA) was used for data analysis and graphic presentations. Statistical analysis was performed by non-linear regression fitting; sigmoidal-dose response curves were performed using the log(inhibitor) vs response analyses. The IC50 value and the maximal efficacy of compounds in inhibiting cell viability (E max) were derived.

5.3.6. Immunofluorescence and microscopy
After treatment, HeLa cells directly grown on sterile poly-L-lysine (Sigma P4832) coated coverslips were fixed in methanol for 6 min at 20 C and processed for IF using mouse a-tubulin antibody (Sigma clone B-5-1-2, 1:3000 dilution) followed by FITC-conjugated anti-mouse secondary antibody (Jackson Immunoresearch Laboratories). Slides were counterstained with 0.05 mg/mL DAPI (Sigma) and mounted in Vectashield (Vector). IF-processed cells were examined under an epifluorescence Nikon Eclipse 90i with a QICAM Fast 1394 (Qimaging) camera and using the NISElements AR 4.0 software (Nikon). Single cell images were routinely taken using immersion oil 100_ objectives with NA 1.3.

5.3.7. Flow cytometric analysis
Cell cycle phase distribution was analyzed after incubation with PI (Sigma P4170). All parameters (FS, SS and FL-3) were acquired in a linear amplification scale. Cell aggregates were gated out on the bi-parametric graph FL-3 lin/ratio. Cell death was analyzed using annexin V-FITC (Immunological Sciences, IK-11120). Cell samples were analyzed in a Coulter Epics XL cytometer (Beckman Coulter) equipped with EXPO 32 ADC software. Data from at least 10.000 cells per sample were acquired and processed using Win MDI software.

5.3.8. RNA extraction, reverse transcription and real-time polymerase chain reaction analysis
Total cellular RNA was processed, and 1 mg of total RNA was retro-transcribed. The cDNA was used to perform a real-time PCR using p21Cip1/Waf1 and GAPDH primers as indicated in Table 1S, Supporting Information. For p21 Cip1/Waf1 and GAPDH genes, the following protocols were used for the PCRs: initialization at 95 C for 3 min, followed by 40 cycles of 95 _C for 10 s, 60 _C for 10 s, 72 _C for 10 s. The melting program was 95 _C for 5 s, 65 _C for 1 min and 97 _C for 10 s. The rate of temperature increase was 1 _C/s, and fluorescence data were continuously acquired. The relative amounts of target genes were normalized to GAPDH expression by Light Cycler® 480 Software version 1.5 (Roche Diagnostics) using the 2DCt method.

Disclaimer
The content of this paper is solely the responsibility of the authors and does not necessarily reflect the official views of the National Institutes of Health.
Acknowledgments
This research was supported by grants PRIN 2015 n. 2015FCHJ8E (to R.S.), Finanziamenti Ateneo 2016 Sapienza Universit_a di Roma n. RG116154CF287B95 (to G.L.R.), AIRC (Italian Association for Cancer Research, grant n. IG10164 to P.L.), and Consiglio Nazionale delle Ricerche Flagship Project InterOmics (to P.L.).

Appendix A. Supplementary data
Supplementary data associated with this article can be found in the online version, at https://doi.org/10.1016/j.fsi.2016.10.010.
These data include MOL files and InChIKeys of the most important compounds described in this article.

References
Towards highly potent cancer agents by modulating the C-2 group of the arylthioindole class of tubulin polymerization inhibitors, J. Med. Chem. 56 (2013) 123e149.


Figures and Tables

![Chart 1](chart1.png)

**Chart 1.** Structures of compounds 1-22.
Table 1
Inhibition of tubulin polymerization, growth of MCF-7 human breast carcinoma cells, and colchicine binding by compounds 3-22.

<table>
<thead>
<tr>
<th>Compd</th>
<th>R</th>
<th>X</th>
<th>IC_{50} ± SD (µM) Tubulin</th>
<th>IC_{50} ± SD (nM) MCF-7</th>
<th>(%) ± SD Inh. Colch. Bind.</th>
</tr>
</thead>
<tbody>
<tr>
<td>3</td>
<td>5-</td>
<td>S</td>
<td>0.87 ± 0.1</td>
<td>70 ± 10</td>
<td>81 ± 1</td>
</tr>
<tr>
<td>4</td>
<td>5-</td>
<td>S</td>
<td>1.8 ± 0.1</td>
<td>200 ± 100</td>
<td>63 ± 1</td>
</tr>
<tr>
<td>5</td>
<td>5-</td>
<td>S</td>
<td>1.3 ± 0.09</td>
<td>150 ± 70</td>
<td>76 ± 2</td>
</tr>
<tr>
<td>6</td>
<td>5-</td>
<td>S</td>
<td>1.7 ± 0.03</td>
<td>200 ± 9</td>
<td>72 ± 3</td>
</tr>
<tr>
<td>7</td>
<td>6-</td>
<td>S</td>
<td>0.59 ± 0.1</td>
<td>10 ± 3</td>
<td>97 ± 1</td>
</tr>
<tr>
<td>8</td>
<td>6-</td>
<td>C=O</td>
<td>0.76 ± 0.04</td>
<td>47 ± 6</td>
<td>96 ± 0.1</td>
</tr>
<tr>
<td>9</td>
<td>6-</td>
<td>S</td>
<td>2.3 ± 0.04</td>
<td>10 ± 0</td>
<td>90 ± 2</td>
</tr>
<tr>
<td>10</td>
<td>6-</td>
<td>C=O</td>
<td>0.61 ± 0.06</td>
<td>43 ± 6</td>
<td>96 ± 0.1</td>
</tr>
<tr>
<td>11</td>
<td>6-</td>
<td>S</td>
<td>0.47 ± 0.05</td>
<td>60 ± 1</td>
<td>95 ± 1</td>
</tr>
<tr>
<td>12</td>
<td>6-</td>
<td>C=O</td>
<td>0.38 ± 0.1</td>
<td>96 ± 6</td>
<td>96 ± 0.8</td>
</tr>
<tr>
<td>13</td>
<td>6-</td>
<td>S</td>
<td>0.58 ± 0.06</td>
<td>45 ± 1</td>
<td>97 ± 0.3</td>
</tr>
<tr>
<td>14</td>
<td>6-</td>
<td>C=O</td>
<td>0.60 ± 0.04</td>
<td>18 ± 4</td>
<td>94 ± 0.6</td>
</tr>
<tr>
<td>15</td>
<td>7-</td>
<td>S</td>
<td>0.64 ± 0.06</td>
<td>15 ± 4</td>
<td>95 ± 0.6</td>
</tr>
<tr>
<td>16</td>
<td>7-</td>
<td>C=O</td>
<td>1.9 ± 0.03</td>
<td>320 ± 10</td>
<td>70 ± 1</td>
</tr>
<tr>
<td>17</td>
<td>7-</td>
<td>S</td>
<td>1.3 ± 0.1</td>
<td>24 ± 2</td>
<td>86 ± 1</td>
</tr>
<tr>
<td>18</td>
<td>7-</td>
<td>C=O</td>
<td>3.0 ± 0.3</td>
<td>1300 ± 0</td>
<td>29 ± 0.6</td>
</tr>
<tr>
<td>19</td>
<td>7-</td>
<td>S</td>
<td>0.57 ± 0.03</td>
<td>29 ± 1</td>
<td>89 ± 1</td>
</tr>
<tr>
<td>20</td>
<td>7-</td>
<td>C=O</td>
<td>2.6 ± 0.04</td>
<td>550 ± 70</td>
<td>48 ± 0.5</td>
</tr>
<tr>
<td>21</td>
<td>7-</td>
<td>S</td>
<td>1.3 ± 0.01</td>
<td>38 ± 10</td>
<td>77 ± 0.1</td>
</tr>
<tr>
<td>22</td>
<td>7-</td>
<td>C=O</td>
<td>6.2 ± 1</td>
<td>1400 ± 100</td>
<td>ND*</td>
</tr>
<tr>
<td>1</td>
<td></td>
<td></td>
<td>0.74 ± 0.05</td>
<td>39 ± 10</td>
<td>88 ± 2</td>
</tr>
</tbody>
</table>

(continued on next page)

Compd  R  X     IC_{50} ± SD (µM) Tubulin IC_{50} ± SD (nM) MCF-7 (%) ± SD Inh. Colch. Bind.
<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>GCH</td>
<td></td>
<td></td>
<td>3.2 ± 0.4</td>
<td>5 ± 1</td>
<td>-</td>
</tr>
<tr>
<td>Cl, Ac</td>
<td></td>
<td></td>
<td>1.0 ± 0.1</td>
<td>13 ± 3</td>
<td>98 ± 0.6</td>
</tr>
</tbody>
</table>

Table continued on next page.

* naïve results, ND, not done.

Experiments were performed in duplicate or triplicate.

\(^a\) Inhibition of tubulin polymerization. Tubulin was at 10 μM in the assembly assay.

\(^b\) Inhibition of growth of MCF-7 human breast carcinoma cells.

\(^c\) Inhibition of [\(\text{H}\)colchicine binding: tubulin, \([\text{H}\)colchicine, inhibitor at 1:5:5 µM.\]

\(^d\) ND, not done.
Scheme 1. Synthesis of Compounds 3e22 and 27e37. Reagents and reaction conditions: (a) (27, 29) boronic derivate, Pd(OAc)2, P(o-tol)3, K3PO4, EOH/PhMe, 80 _C, 2 h, 73e87%; (b) (28) 3-furanboronic acid pinacol ester, Pd(OAc)2, K2CO3, methylpyrrolidone/water, closed vessel, 200W, 110 _C, 15 min, 45%; (c) (3e7, 9, 11, 13, 15, 19 and 37) bis(3,4,5-trimethoxyphenyl)disulfide, NaH, anhydrous DMF, closed vessel, 120W, 130 _C, 2 min, 5e50%; (d) (8, 10, 12, 14) (i) diethylaluminum chloride, CH2Cl2, -78 _C, Ar stream; (ii) 3,4,5-trimethoxybenzoyl chloride, _10 _C/25 _C within 2 h, 14e69%; (e) (16e18, 20e22, 30e32, 35 and 36) boronic derivate, Pd2dba3, SPhos, K3PO4, BuOH, 100 _C, 15 h, 5e75%; (f) (38) 3,4,5-trimethoxybenzoyl chloride, CH2Cl2, anhydrous AlCl3, closed vessel, 150W, 110 _C, 2 min, 51%.

Chart 2. SAR summary for tubulin polymerization inhibition (TPI) and MCF-7 cell growth inhibition (CGI) by compounds 3-22.
Fig. 1. Plants proposed binding mode for derivatives 6 (yellow), 19 (pink) and 21 (green) into the 1SA0 tubulin structure. Residues involved in interactions are shown as stick diagrams. The tubulin polypeptide chains are shown as ribbon cartoons. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

Fig. 2. Plants proposed binding mode for derivatives 13 (left panel) and 19 (right panel) for the studied tubulin structures: 1SA0 (green), 3HKC (cyan), 4O2B (magenta), SCA0 (purple), 5CB4 (sky blue), 5LYI (pink). Residues involved in interactions are shown as stick diagrams. The tubulin polypeptide chains are shown as ribbon cartoons. Residues and cartoon are from 1SA0. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

Table 2
Growth Inhibition of HT29, HCT116, HepG2, T98G, U87MG, U343G, MV4-11, THP-1, A-549 and PC3 cells by 13 and 19.a,b

<table>
<thead>
<tr>
<th>Compd</th>
<th>HT29</th>
<th>HCT116</th>
<th>HepG2</th>
<th>T98G</th>
<th>U87MG</th>
<th>U343G</th>
<th>MV4-11</th>
<th>THP-1</th>
<th>A-549</th>
<th>PC3</th>
</tr>
</thead>
<tbody>
<tr>
<td>13</td>
<td>10 ± 1.4</td>
<td>21 ± 1.5</td>
<td>20 ± 1.1</td>
<td>26 ± 3.6</td>
<td>16 ± 2.0</td>
<td>31 ± 4.3</td>
<td>6 ± 2.0</td>
<td>2 ± 1</td>
<td>2 ± 0.1</td>
<td>35 ± 2</td>
</tr>
<tr>
<td>19</td>
<td>69 ± 1.4</td>
<td>70 ± 1.7</td>
<td>80 ± 1.2</td>
<td>211 ± 20</td>
<td>96 ± 14</td>
<td>154 ± 5.9</td>
<td>70 ± 16</td>
<td>20 ± 2</td>
<td>58 ± 1</td>
<td>39 ± 4</td>
</tr>
</tbody>
</table>

a Cytotoxic concentrations for the indicated cell lines; cytostatic concentrations for A-549 and PC3 cells.
b Incubation time was 48 h for U343G, MV4-11 and PC3 cells, incubation was 72 h.
Table 3

Growth Inhibition of the OVCA-8 and NCI/ADR-RES cells by compounds 13 and 19 and reference compounds CSA4, VRB, VLB and PTX.a

<table>
<thead>
<tr>
<th>Compd</th>
<th>IC_{50} ± SD (nM)</th>
<th>OVCA-8</th>
<th>NCI/ADR-RES</th>
<th>MCF-7</th>
</tr>
</thead>
<tbody>
<tr>
<td>13</td>
<td>9.3 ± 2</td>
<td>7.5 ± 2</td>
<td>6.7 ± 2</td>
<td></td>
</tr>
<tr>
<td>19</td>
<td>58 ± 8</td>
<td>34 ± 8</td>
<td>55 ± 7</td>
<td></td>
</tr>
<tr>
<td>CSA4</td>
<td>4.0 ± 0</td>
<td>3.5 ± 0.7</td>
<td>5.0 ± 1</td>
<td></td>
</tr>
<tr>
<td>PTX</td>
<td>4.0 ± 1</td>
<td>3100 ± 600</td>
<td>5.5 ± 0.7</td>
<td></td>
</tr>
<tr>
<td>VRB</td>
<td>300 ± 0</td>
<td>5000 ± 1000</td>
<td>Nd</td>
<td></td>
</tr>
<tr>
<td>VBL</td>
<td>15 ± 7</td>
<td>200 ± 0</td>
<td>Nd</td>
<td></td>
</tr>
</tbody>
</table>

a Inhibition of growth of the indicated cell lines.

Same value obtained in all experiments.

Nd, not done.

Fig. 3. FACS analysis of the cell cycle and cell death in HeLa cell populations treated with 13 or 19. Panel A left: distribution of cells after a 24 h treatment with 13 or 19 or VBL (DMSO as control). The DNA content indicates the cell cycle phase (2C: G1; 4C: G2/M; intermediate values between 2C and 4C: S phase). The panel shows a representative analysis of the cell cycle phase distribution, arranged according to the compound concentration (on the z axis). In the lower two diagrams, 20 nM, green curves; 50 nM, red curves; 100 nM; blue curves. The profiles illustrate that 19 only induces 4C cell accumulation when used at 100 nM, whereas 13 is fully effective at 50 nM. Panel A right: the histograms show the mean % values ± SD of 4C cells (G2/M phases) counted in 3 independent experiments. Panel B: the graph represents the frequency of mitotic figures (grey bars) and multinucleated cells (blue bars) in cell populations treated for 24 h under the indicated conditions, processed for immunofluorescence and scored by microscopy. Between 300 and 650 cells were counted for each condition in 2 independent experiments. Panel C: FACS analysis of cell populations
treated for 24 h under the indicated conditions and stained with annexin V. Mean ± SD values were calculated from three experiments. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

**Fig. 4.** IF analysis of mitotic cells in HeLa cultures treated with 13 or 19. Panel A: representative phenotypes induced by 19: all cells were arrested in prometaphase (compare the configuration of chromosomes in DAPI panels); the green channel shows the extent of inhibition of tubulin polymerization at increasing compound concentrations (indicated at the right of the panels in nM). Panel B: representative phenotypes induced by 13: the DAPI panels show the prometaphase configuration of chromosomes; the green channel shows the extent of inhibition of tubulin polymerization at the indicated concentrations. Panel C: Distribution of tubulin cytological phenotypes under the indicated conditions. The histograms represent the frequency of the indicated phenotypes (n, 100 to 250 prometaphase-arrested cells per condition). (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

**Table 4**
Physico-chemical profiles of compounds 13 and 19.

<table>
<thead>
<tr>
<th>Compd</th>
<th>Log P</th>
<th>MW</th>
<th>LogS</th>
<th>dPSA</th>
<th>HBA/HHBD</th>
<th>RotStab</th>
<th>Verbe</th>
<th>3/75</th>
</tr>
</thead>
<tbody>
<tr>
<td>13</td>
<td>5.34</td>
<td>397.51</td>
<td>-5.58</td>
<td>97.82</td>
<td>5</td>
<td>good</td>
<td>warning</td>
<td></td>
</tr>
<tr>
<td>19</td>
<td>5.37</td>
<td>397.51</td>
<td>-5.60</td>
<td>97.82</td>
<td>5</td>
<td>good</td>
<td>warning</td>
<td></td>
</tr>
</tbody>
</table>

- Log P: Logarithm of the partition coefficient between n-octanol and water computed by mLOGP3 method [38].
- MW: Molecular weight.
- LogS: Logarithm of compound water solubility computed by the ESOL method [39].
- dPSA: Molecular polar surface area, this parameter has been shown to correlate with human intestinal absorption (<1.40) [40].
- HBA/HHBD: Number H-bond acceptors and H-bond donors.
- RotStab: violation of the rule of five [36].
- Verbe: Verbe's rule matching [35].
- 3/75: 3/75 rule matching [36].
Fig. 5. Physico-chemical properties. Left. Compounds 13 and 19 (blue lines) are compared with the optimal light green area defined by Lipinski’s rule-of-five and Veber’s rule. The computations includes logP, molecular weight (MW), topological polar surface area (tPSA), rotatable bonds (RotB), H-bond acceptors and donors (HBA, HBD). The red area indicates low oral bioavailability. Estimated physico-chemical profiles of compounds 13 and 19 fall within the light green area. Right. Likelihood of in vivo toxicity according 3/75 rule [37]. tPSA is expressed as a function of logP. Compounds located in the red square are likely to cause toxicity. Estimated toxicities of compounds 13 and 19 (blues point) fall within the light green area. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)