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## **ARTICLE TYPE**

## Synthesis and anti-cancer screening of novel heterocyclic-(2*H*)-1,2,3triazoles as potential anti-cancer agents

Narsimha Reddy Penthala,<sup>a</sup> Leena Madhukuri,<sup>b</sup> Shraddha Thakkar,<sup>a</sup> Nikhil Reddy Madadi,<sup>a</sup> Gauri Lamture,<sup>a</sup>

Robert L. Eoff<sup>b</sup> and Peter A. Crooks<sup>a</sup>\*

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*trans*-Cyanocombretastatin A-4 (*trans*-CA-4) analogues have been structurally modified to afford their more stable CA-4-(2H)-1,2,3-triazole analogues. Fifteen novel, stable 4-heteroaryl-5-aryl-(2H)-1,2,3-triazole CA-4 analogues (8a-i, 9 and 11a-e) were evaluated for anti-cancer activity against a panel of 60 human cancer cell lines. These analogues displayed potent cytotoxic activity against both

<sup>10</sup> hematological and solid tumor cell lines with  $GI_{50}$  values in the low nanomolar range. The most potent compound, **8a**, was a benzothiophen-2-yl analogue that incorporated a 3,4,5-trimethoxyphenyl moiety connected to the (2*H*)-1,2,3-triazole ring system. Compound **8a** exhibited  $GI_{50}$  values of <10 nM against 80% of the cancer cell lines in panel. Three triazole analogues, **8a**, **8b** and **8g**, showed particularly potent growth inhibition against the triple negative Hs578T breast cancer cell line with  $GI_{50}$  values of 10.3 nM, 66.5 nM and 20.3 nM, respectively. Molecular docking studies suggest that these compounds bind to the same hydrophobic pocket at the

15 interface of  $\alpha$ - and  $\beta$ -tubulin that is occupied by colchicine and *cis*-CA-4, and are stabilized by Van der Waals' interactions with surrounding amino acid residues. Compound **8a** was found to inhibit tubulin polymerization *in vitro* with an IC<sub>50</sub> value of 1.7  $\mu$ M. The potent cytotoxicity of these novel compounds and their inhibition of tubulin dynamics make these triazole analogues promising candidates for development as anti-cancer drugs.

#### 20 Introduction

The combretastatins (Fig. 1, 1 and 2) were first isolated from the South African bush willow tree *Combretum caffrum*, of which *cis*-combretastatin A-4 (*cis*-CA-4, 1) is the major component.<sup>1</sup> Three analogues of combretastatin *cis*-CA-4: CA-<sup>25</sup> 4P (3), Oxi4503 (4) and AVE8062 (5) (Fig. 1) are currently in clinical trials for treatment of a variety of cancers, and have improved water solubility, cytotoxicity, efficacy and a better safety profile compared to *cis*-CA-4.<sup>2-5</sup> These combretastatin analogues are also known to be vascular disrupting agents <sup>30</sup> (VDAs) and have proven to be advantageous cytotoxic agents, since they target the tumor vasculature causing hemorrhagic necrosis and cell death.<sup>6</sup>

The CA-4 analogues are generally classified as tubulin <sup>35</sup> polymerization inhibitors with an anti-mitotic mechanism of action, due to their ability to disrupt microtubule dynamics and hence mitotic spindle formation by binding to the same site as colchicine on tubulin dimers.<sup>7, 8</sup> These small molecule VDAs also induce hypoxia-driven necrosis of solid tumors, and are <sup>40</sup> able to induce apoptosis by cell-cycle arrest.<sup>9, 10</sup>

<sup>a</sup>Department of Pharmaceutical Sciences, College of Pharmacy, University of Arkansas for Medical Sciences, Little Rock, AR 72205,

45 USA<sup>c</sup> Department of Biochemistry and Molecular Biology, University of Arkansas for Medical Sciences, Little Rock, AR 72205-7199, U.S.A. \* Corresponding author: Tel.: +1-501-686-6495; fax: +1-501-686-6057; e-mail: pacrooks@uams.edu



Fig 1. Chemical structures of *cis*-CA-4 (1) and its derivatives (2-6).

CA-4 analogues are highly cytotoxic against human cancer <sup>70</sup> cell lines. CA4-P (**3**) exhibits potent cytotoxicity against anaplastic thyroid cancer cells lines,<sup>11</sup> and Oxi-4503 (**4**) is a potent anti-leukemic compound.<sup>12</sup> AVE8062 (**5**) is an effective cytotoxic agent against head and neck squamous cell carcinoma,<sup>13</sup> including multidrug resistant forms of this cell <sup>75</sup> line.<sup>14</sup> Tron *et al.* have reported that the *cis*-configured double bond of *cis*-CA-4, with a 3,4,5-trimethoxyphenyl ring (Ring A) on one side and a 4-methoxysubstituted phenyl ring on the other side (Ring B), is an essential structural moiety for optimal cytotoxic activity in *cis*-CA-4 analogues.<sup>15-17</sup> Recently, we have investigated the anticancer activities of a series of *trans*-cyano-CA-4 analogs in which replacement of ring B with <sup>5</sup> different heterocycles, such as quinoline (6), benzothiophene (7a-f and 10a-c), benzofuran (7g-h) and indole (7i and 10d-e),

has occurred; we have determined the effect of these structural changes on anticancer activity.<sup>18, 19</sup> We have also evaluated the effects of structurally modifying ring A by introducing <sup>10</sup> differently substituted methoxy and hydroxy moieties into this phenyl ring.<sup>18</sup>

The isomerization of the *cis*-configured CA-4 analogues to their *trans*-forms in solution is one of the major disadvantages <sup>15</sup> of these molecules. In this respect, we have shown that isomerization of *trans*-cyano CA-4 analogues to their *cis*-cyano CA-4 forms is also observed in solution.<sup>18</sup> To restrict *cis*-*trans* isomerization in these CA-4 analogues, different heterocylic ring systems, such as doxolane,<sup>20</sup> thiazole<sup>21</sup> 1,5-disubstituted <sup>20</sup> 1,2,3-triazoles,<sup>22-26</sup> 1,2,4-triazoles<sup>27, 28</sup> and imidazole,<sup>29</sup> have

been incorporated into the stilbene double bond.

In the present report, we have prepared a series of stable analogues of *cis*-cyano CA-4 by constructing a triazole ring <sup>25</sup> system across the olefinic bond between aromatic rings A and B. This (2*H*)-1,2,3-triazole heterocyclic bridge unit was established by chemical modification of existing *trans*-cyano CA-4 analogues via [3+2] cycloaddition of azide ion across the double bond. This novel and simple structural modification led

- <sup>30</sup> to an improvement in the stability and biological properties of these new CA-4 analogues. We now report on the synthesis and *in vitro* anti-cancer activities of a variety of 4-heteroaryl-5-aryl-(2*H*)-1,2,3-triazole CA-4 analogues, and their ability to inhibit tubulin polymerization *in vitro*. Some of the analogues were
- as also tested for cytotoxicity against Hs578T breast cancer cell lines. In addition, molecular modeling studies of the most active inhibitors indicate that they interact with the colchicine binding site on  $\alpha$ , $\beta$ -tubulin, and this has been confirmed in *in vitro* tubulin polymerization inhibition studies.

#### Chemistry

The formation of the 1,2,3-triazole ring system is well known in the literature as a "click chemistry" product from a <sup>45</sup> thermally induced Cu(I)-catalyzed (CuAAC) Huisgen [3+2]cycloaddition azide-alkyne reaction.<sup>30</sup> The click chemistry approach is widely used in the synthesis and regioisomeric formation of 1,4-disubstituted-1,2,3-triazoles in the presence of CuAAC in excellent yields.<sup>31-34</sup> The

<sup>50</sup> regioisomeric synthesis of 1,5-disubstituted-1,2,3-triazoles has also been reported with high selectivity utilizing different metal ions in magnesium-, cerium- and ruthenium-mediated reactions.<sup>35-37</sup>

<sup>55</sup> We have recently reported the synthesis of a wide variety of 4,5-disubstitued-1,2,3-(1*H*)-triazole analogues.<sup>38, 39</sup> We used a two stage synthesis, i.e. the preparation of the precursor *trans*-cyano-CA-4 analogue, which utilizes our previously published

procedure,<sup>18</sup> followed by synthesis of the target (2*H*)-1,2,3-<sup>60</sup> triazole CA-4 compounds **8a-8i** and **11a-11e** by reaction of the *trans*-cyano CA-4 analogue with sodium azide in DMSO at 100 °C over 1-6 hrs utilizing L-proline as a Lewis base. The desired (2*H*)-1,2,3-triazole products were obtained in 75-96% yield<sup>38</sup> (Schemes 1 and 2) (Note: the use of *cis*-cyano CA-4

<sup>65</sup> analogues in the above reaction led to lower yields of the triazole product<sup>39</sup>). The *N*-2 methylated analogue **9** was prepared from the reaction of **8a** with methyl iodide/K<sub>2</sub>CO<sub>3</sub> in acetone (Scheme 2). Confirmation of the structure and purity of these analogues is reported elsewhere, and was obtained from <sup>70</sup> single crystal X-ray diffraction studies,<sup>40</sup> and <sup>1</sup>H-NMR, <sup>13</sup>C-

NMR and high resolution mass spectroscopic analysis.<sup>38</sup>



Scheme 1 Synthesis of substituted 4-heteroaryl-2-yl-5-phenyl-2*H*-1,2,3-triazoles (8a-i).



Scheme 2 Synthesis of substituted 4-heteroaryl-3-yl-5-phenyl-2*H*-1,2,3-triazoles (11a-e).



Scheme 3 Synthesis of 4-(benzo [*b*]thiophen-2-yl)-2-methyl-5-(3,4,5-trimethoxyphenyl)-2*H*-1,2,3-(2*H*)-triazole (9).

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#### **Biological Evaluation** A. *In vitro* growth inhibition and cytotoxicity

The above triazole analogues were evaluated for their <sup>5</sup> anticancer activities against a panel of 60 human cancer cell lines, which incorporates different subpanels representing leukemia, non-small cell lung, colon, central nervous system, melanoma, ovary, renal, prostate, and breast cancer cell lines, at a concentration of  $10^{-5}$  M utilizing the procedure described

<sup>10</sup> by Rubinstein et al.<sup>41</sup> Compounds that showed 60% growth inhibition in at least eight of the 60 cell lines screened were further evaluated in a five-dose screen.

From the preliminary screen, compounds **8a**, **11b**, **11d**, **11e** <sup>15</sup> and **9** were selected for five-dose studies and their GI<sub>50</sub> values determined (Table 1). GI<sub>50</sub> values represent the molar drug concentration required to cause 50% growth inhibition. In the five-dose studies, 5 different concentrations at 10-fold dilutions (10<sup>-4</sup> M, 10<sup>-5</sup> M, 10<sup>-6</sup> M, 10<sup>-7</sup> M and 10<sup>-8</sup> M) were utilized and <sup>20</sup> incubations were carried out over 48 h exposure to drug. Compounds **8a**, **11b**, **11d**, **11e** and **9** exhibited GI<sub>50</sub> values in the low nanomolar range against all 60 human cancer cell lines in the panel. The results are presented in Table 1.

<sup>25</sup> Compound **8a** (4-(benzo[*b*]thiophen-2-yl)-5-(3,4,5-trimethoxyphenyl)-2*H*-1,2,3-triazole) exhibited GI<sub>50</sub> values <0.050  $\mu$ M in all the 60 human cancer cell lines in the panel, and had GI<sub>50</sub> <0.01  $\mu$ M in 80% of the cancer cells in the panel (Table 1). Compound **8a** exhibited particularly potent growth <sup>30</sup> inhibition properties against all six cell lines in the breast cancer subpanel, with GI<sub>50</sub> values of <0.01  $\mu$ M to 0.014  $\mu$ M.

Compound **9**, obtained by *N*-methylation of **8a**, exhibited  $GI_{50}$  values ranging from 0.023  $\mu$ M to 0.744  $\mu$ M in 84% of the <sup>35</sup> cancer cell lines screened and showed moderate growth inhibition properties in all six breast cancer cell lines, with  $GI_{50}$  values in the range of 0.040  $\mu$ M to 1.6  $\mu$ M (except for the T-47D cell line). This compound also exhibited potent growth inhibition against NCI-H522 lung cancer ( $GI_{50} = 0.023 \ \mu$ M) <sup>40</sup> and MDA-MB-435 melanoma ( $GI_{50} = 0.025 \ \mu$ M) cell lines (Table 1).

Compound **11b** exhibited GI<sub>50</sub> values ranging from 0.337  $\mu$ M to 13.3  $\mu$ M against the entire 60 cancer cell panel screened, <sup>45</sup> and showed moderate growth inhibition properties against all six breast cancer cell lines, with GI<sub>50</sub> values in the range of 0.63  $\mu$ M to 3.21  $\mu$ M. This compound exhibited potent growth inhibition against MDA-MB-435 melanoma cells (GI<sub>50</sub> = 0.337  $\mu$ M) (Table 1).

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Compound **11d** exhibited  $GI_{50}$  values ranging from 0.014  $\mu$ M to 0.44  $\mu$ M in 90% of the cancer cell lines screened and showed potent growth inhibition properties in all six breast cancer cell lines, with  $GI_{50}$  values in the range of 0.028  $\mu$ M to <sup>55</sup> 0.089  $\mu$ M. This compound exhibited potent growth inhibition

against MDA-MB-435 melanoma cells ( $GI_{50} = 0.014 \mu M$ ) (Table 1). The average  $GI_{50}$  value of this compound against all the cancer cell lines screened was 0.424  $\mu M$ .

Table 1.	Antitumor	activity	(GI <sub>50</sub>	in	$\mu M)^a$	data	for	compou	inds	8a,	9,
11b, 11d,	11e in the 1	NCI 60-c	ell lin	e s	creen						

	8a	9	11b	11d	11e
Panel/cell line	GI50 (µM)	GI50 (µM)	GI50 (µM)	GI50 (µM)	GI50 (µM)
Leukemia					
CCRF-CEM	0.028	0.22	3.29	0.048	0.037
HL-60(TB)	< 0.01	0.048	1.70	0.046	0.033
K-562 MOLT 4	< 0.01	0.042	2.05	0.053	0.031
RDMI 8226	0.042	0.353	3.03	0.10	0.063
SR	< 0.01	0.302 nd	3.04 nd	0.09	0.033
Lung Cancer	<0.01	nu	nu	0.07	0.047
A549/ATCC	0.023	0.167	4.52	0.09	0.088
EKVX	0.010	0.466	3.42	na	na
HOP-62	0.015	0.553	2.18	na	na
HOP-92	< 0.01	0.080	0.921	nd	0.029
NCI-H226	nd	>100	13.2	0.39	0.075
NCI-H23	< 0.01	0.392	3.94	0.27	0.124
NCI-H322M	< 0.01	1.26	7.79	nd	40.7
NCI-H460	< 0.01	0.069	3.56	0.042	0.038
NCI-H522	< 0.01	0.023	0.601	0.021	0.013
COLO 205	< 0.01	0.082	2 82	0.034	0.035
HCC-2998	0.01	0.407	4.21	0.034	0.055
HCT-116	< 0.01	0.073	2.84	0.041	0.031
HCT-15	< 0.01	0.058	1.67	0.035	0.031
HT29	0.020	0.055	2.78	0.033	0.031
KM12	0.022	0.054	2.28	0.045	0.045
SW-620	< 0.01	0.063	3.09	0.043	0.035
CNS Cancer					
SF-268	< 0.01	0.63	3.69	0.071	0.067
SF-295 SE 520	< 0.01	0.030	2.33	0.032	0.026
ST-339 SNB 10	< 0.01	0.149	1.64	0.051	0.024
SNB-75	< 0.01	0.2.34 nd	4.07 nd	0.001	0.030
U251	< 0.01	0.078	2.19	0.050	0.023
Melanoma	-0.01	0.070	2.17	0.007	0.001
LOX IMVI	< 0.01	0.168	3.04	0.071	0.055
MALME-3M	nd	16.9	13.3	nd	nd
M14	< 0.01	0.076	1.95	0.036	0.031
MDA-MB-435	< 0.01	0.025	0.337	0.014	0.010
SK-MEL-2	< 0.01	0.059	1.44	0.030	0.031
SK-MEL-28	< 0.01	13.0	4.63	0.112	Nd
SK-MEL-5	< 0.01	0.225 51.6	2.85	12.0	0.034
UACC-237	< 0.01	0.053	2 77	13.9	>100.0
Ovarian Cancer	<0.01	0.055	2.17	0.045	0.034
IGROV1	< 0.01	0.707	4.21	na	na
OVCAR-3	< 0.01	0.354	1.94	0.033	0.028
OVCAR-4	0.042	nd	6.33	1.16	2.67
OVCAR-5	0.042	2.36	7.91	0.381	0.416
OVCAR-8	< 0.01	0.352	4.24	1.16	0.070
NCI/ADR-RES	< 0.01	0.103	2.39	0.034	0.030
SK-OV-3	< 0.01	0.311	2.37	0.042	0.034
Renal Cancer	<0.01	0.212	2.26	0.025	0.020
A498	< 0.01	0.312	2.96	0.035	0.030
ACHN	< 0.01	0.294	3.94	0.043	0.035
CAKI-1	< 0.01	3.77	3.69	0.044	0.038
RXF 393	nd	0.286	2.11	0.164	0.052
SN12C	0.048	nd	8.80	1.39	0.069
TK-10	nd	0.578	2.98	1.39	0.083
UO-31	< 0.01	0.744	2.78	0.087	0.062
Prostate Cancer					
PC-3	0.034	0.077	3.08	0.044	0.038
DU-145	< 0.01	0.263	3.77	0.090	0.075
MCF7	<0.01	0.040	1 15	0.020	0.020
IVICI /	~0.01	0.040	1.43	0.029	0.029

	0.01	0.002	0.05	0.0 .2	0.007
MDA-MB-468	< 0.01	0.082	0.63	0.042	0.037
T-47D	nd	>100	2.83	0.048	0.044
BT-549	< 0.01	0.422	2.17	0.028	0.017
HS 578T	< 0.01	0.270	2.25	0.042	0.031
MDA-MB-	0.014	1.60	3.21	0.089	0.076

na: not analyzed; nd: not determined;  ${}^{a}GI_{50}$ : 50% growth inhibition, concentration of drug resulting in a 50% reduction in net cell growth as compared to cell numbers on day 0.

<sup>5</sup> Compound **11e** exhibited GI<sub>50</sub> values ranging from 0.01 μM to 41.6 μM in 94% of the cancer cell lines screened and showed potent growth inhibition properties in all six breast cancer cell lines, with GI<sub>50</sub> values in the range of 0.017 μM to 0.076 μM. This compound exhibited potent growth inhibition against <sup>10</sup> MDA-MB-435 melanoma cells (GI<sub>50</sub> = 0.01 μM) and NCI-H522 (GI<sub>50</sub> = 0.013 μM) non-small cell lung cancer cell lines (Table 1). The average GI<sub>50</sub> value of this compound against all 60 cancer cell lines screened was 0.856 μM.

## 15 B. Cytotoxic effects of 8a, 8b, and 8g on Hs578T breast cancer cell lines

Compound 8a and its analogues (8b and 8g) were tested for cytotoxicity against the triple negative breast cancer (TNBC)-<sup>20</sup> derived Hs578T cell line. The GI<sub>50</sub> estimates for compounds 8a, 8b, and 8g were all below 100 nM (Fig. 2). However, the potency of compound 8a remained two- to six-fold greater than either 8b or 8g. Comparing the results for 8a, 8b and 8g revealed that replacing the 3,4,5-trimethoxyphenyl moiety of <sup>25</sup> 8a with a 3,5-dimethoxyphenyl group to afford 8b, results in a more pronounced effect on cytotoxicity than replacing the

benzothiophen-2-yl moiety with a benzofuran-2-yl group (8g).



Fig. 2 Hs578T TNBC cell viability is sensitive to compounds 8a, 8b, and 8g. Dose-response curves for Hs578T cell viability in the presence of varying concentrations of compounds 8a (closed circles), 8b (closed squares), and 8g (open circles) are

(costed energy, 60 (costed squares), and 6**g** (open energy) are 45 shown. The values shown represent the mean  $\pm$  s.d. (n=3). Fitting the data to a variable slope log (inhibitor) vs. response equation yields the following GI<sub>50</sub> estimates (with 95% confidence intervals): **8a**: GI<sub>50</sub> = 10 (9.8-11) nM, **8b**: GI<sub>50</sub> = 65 (52-82) nM, **8g**: GI<sub>50</sub> = 20 (16- 26) nM.

#### C. Molecular modeling studies

Molecular modeling studies were performed for four active (2*H*)-1,2,3-triazoles analogues (8a, 8g, 11d, and 11e) using 55 SYBYL-X 2.1. Binding interactions of these triazole analogues

were studied by docking them at the colchicine-binding site on tubulin. 3-D coordinates for the tubulin-colchicine complex were obtained from the RSCB protein data bank (pdb id: 4O2B). For preparation of tubulin to perform the docking 60 calculations, the geometry of the protein molecule was optimized using energy minimization techniques after the

- addition of hydrogen atoms. Since no structured water molecule was present in the binding pocket, all the water molecules in the protein were deleted during the preparation of
- 65 structure for docking calculations. Amino acid side chain bumps were fixed and terminal amino acid were charged to mimic the biological environment. Hydrogen atoms were added to the structure. The Kollman force field was applied and the protein was minimized using the Powell method and Pullman 70 charges. The triazole analogue structures were initially
- generated in 2-D format in Chem-Draw. For generating energyminimized structures, the 2-D structures were converted to 3-D using ChemDraw3D and all the structures were saved in Mol2 format. The structures were imported to Sybyl in the Mol2
- <sup>75</sup> format. In SYBYL, TRIPOS force field was applied and 3-D structure coordinates were taken through a series of minimization processes. Firstly, all the structures we minimized using the BFGS method. Then, MOPAC charges were calculated for each molecule followed by another <sup>80</sup> minimization step. Each molecule was then checked for
- charges, bond angle, torsion angles and geometry. All the molecules were saved in one directory and prepared for docking via the ligand preparation tool. For the docking analysis using the Surflex program, a protomol was generated
- at the colchicine-binding site in tubulin. This protomol is the representation of the binding site that simulates the binding environment experienced by the ligand. The docking analysis is performed using the Surflex Dock-Gemox module which lists the C-Scores (consolidated scores) for each molecule. The C-
- <sup>90</sup> Score is a measure of the goodness of fit. The C-Score function combines the binding score obtained from five different scoring algorithms; namely: Total Score, G Score, PMF Score, D Score and Chem Score. In these docking calculations, the flexibilities of the ligands are accounted for by considering 20 different <sup>95</sup> conformational states and scoring each of them.

Docking analysis at the colchicine binding site determined the binding modes of the four triazole analogues. Compound 8a exhibited the strongest binding interactions at the colchicine 100 binding site in tubulin compared to the other three analogues. 8a participates in strong hydrogen bonding with ASN 258 and THR 353 (Fig. 3) and makes a weak hydrogen bond with ASN 258. Out of 20 different tested conformations of 8a, four conformations showed the highest C-Score of 4, which was 105 higher than any of the C-Scores for the other three triazole analogues. Compound 8g is participates in hydrogen bonding with ASN 258 and ALA 317, with one conformation exhibiting the highest C-Score of 4. Compounds 11d and 11e each participate in a hydrogen bonding interaction with ASN 258, 110 while **11d** participates in another hydrogen bonding interaction with THR 356, and 11e makes hydrogen bond with ALA 317 (Fig. 3). ASN 258 hydrogen bond interactions are common in all four docked triazoles.



Fig. 3 Binding modes of compounds 8a, 8g, 11d and 11e at the colchicine binding site on tubulin

In summary, **8a** exhibited the strongest binding interactions with tubulin at the colchicine-binding site, which is consistent with the anticancer screening data (Table 1). Thus, the above <sup>30</sup> molecular modeling data predicts that stable, more drug-like triazolo derivatives of the *cis*-cyano CA-4 analogues formed by constructing a heterocyclic ring system across the olefinic bond

between aromatic rings A and B will retain the tubulin targeting properties of the parent compounds.

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#### D. Inhibition of tubulin polymerization

The ability of compounds **8a**, **11d**, **11e** and **9** to inhibit tubulin polymerization was determined using a commercially 40 available *in vitro* assay (Cytoskeleton Inc., Denver, CO). We examined the ability of three of the most active CA-4-(2*H*)-1,2,3-triazoles, **8a**, **11d**, and **11e**, to inhibit tubulin polymerization using an *in vitro* assay. Compound **8a**, which exhibited the greatest overall potency against the NCI 60 45 human cancer cell line panel, was found to inhibit tubulin

- polymerization with an IC<sub>50</sub> value of 1.7 ( $\pm$  0.4)  $\mu$ M (Fig. 4), where the reported values represent the mean  $\pm$  s.d. (n=3). Compounds **11d** and **11e** inhibited tubulin polymerization with IC<sub>50</sub> values of 18.5 ( $\pm$  9.8)  $\mu$ M and 13.5 ( $\pm$  3.5)  $\mu$ M,
- <sup>50</sup> respectively, in this assay. The *N*-methylated analogue **9** failed to inhibit tubulin polymerization even at concentrations as high as 200  $\mu$ M (data not shown). These results are consistent with the idea that compound **8a** exerts some of its cytotoxic effects on cells through inhibition of tubulin polymerization. The less
- <sup>55</sup> potent inhibition of tubulin polymerization by compounds **11d** and **11e** is consistent with the general trend towards decreasing potency (but still significant), as determined from  $GI_{50}$  values from the cancer cell screens, when compared to compound **8a**,

culture are not mediated by inhibition of tubulin polymerization, but are instead the result of an unknown mechanism of action. A E 0.20 DMSO

85 and is also consistent with the molecular docking studies.

However, based on the in vitro assay results, it would appear

that the cytotoxic effects of compound 9 observed in cell



Fig. 4 Compound 8a inhibits tubulin polymerization. (A) An *in vitro* assay was used to measure inhibition of tubulin polymerization by compound 8a. (B) A dose-response curve for tubulin polymerization in <sup>95</sup> the presence of varying concentrations of compound 8a. The values shown represent the mean  $\pm$  s.d. (n=3).

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#### Conclusions

We have evaluated a small library of novel 4-heteroaryl-5aryl-(2H)-1,2,3-triazoles that are structurally related to trans-5 cyano CA-4 analogues that inhibit tubulin polymerization.

- These compounds have been evaluated in a panel of 60 human cancer cell lines for their anti-cancer activities, and in assays to determine their ability to inhibit tubulin polymerization. The most potent compound, 8a, and two closely related analogues,
- 10 8b and 8g, were also tested for cytotoxicity against triple negative Hs578T breast cancer cell lines. Compound 8a, a benzothiophen-2-yl derivative was shown to be a more potent anti-cancer agent than its isosteric indol-2-yl (8i) and benzofuran-2-yl (8g) congeners. Compound 8a was also found
- 15 to be the most effective inhibitor of tubulin polymerization. However, the benzothiophen-3-yl isostere of 8a, compound 11a, did not possess any significant anti-cancer activity, although indol-3-yl analogues that incorporate a 3,4,5trimethoxyphenyl (11d) or a 3,5-dimethoxyphenyl (11e)
- 20 moiety into their structure were potent anti-cancer agents. Interestingly, a significant decrease in anti-cancer activity was observed with the N-2-methylated analogue 9 when compared to the parent compound 8a.
- In summary, the anti-cancer screening data, inhibition of 25 tubulin polymerization, and molecular docking studies are all promising indicators that compound 8a warrants further study as a potential anti-cancer agent.

#### **30 EXPERIMENTAL PROCEDURES**

General synthetic procedure for the synthesis of 4-heteroaryl-5-(substituted phenyl)-2H-1,2,3-triazoles (8a-i and 11a-e):<sup>25</sup> Sodium azide (0.01 mol) in DMSO (1 ml) was added to the 35 appropriate heteroaryl cyanostilbene (0.01 mol) followed by addition of L-proline (0.002 mol). The resulting reaction mixture was heated to 100 °C and stirred for 1-6 h. After completion of the reaction, the reaction mixture was cooled to 20 °C, water (10 ml) was added, and the resulting slurry was 40 extracted with ethyl acetate (2x10 ml). The organic layers were separated and combined, dried over anhydrous sodium

- sulphate, concentrated on a rotovaporator and purified by silica gel column chromatography (ethylacetate:hexanes 1:4) to afford the pure product.
- Synthetic procedure for the synthesis of 4-(benzo/b) thiophen-2-yl)-2-methyl-5-(3,4,5-trimethoxyphenyl)-2H-1,2,3triazole (9):<sup>25</sup> A mixture of 4-(benzo[b]thiophen-2-yl)-5-(3,4,5-trimethoxy phenyl)-2H-1,2,3-triazole (6a) (1 mmol),
- 50 K<sub>2</sub>CO<sub>3</sub> (10 mmol) and MeI (2 mmol) in 10 volumes of acetone was refluxed for 8 h and the reaction monitored by TLC analysis. After completion, the reaction mixture was filtered to remove the K<sub>2</sub>CO<sub>3</sub>, washed with acetone and the filtrate evaporated to dryness on a rotovaporator. The resulting residue
- 55 was submitted to ethyl acetate/hexane flash column chromatography to yield 4-(benzo[b]thiophen-2-yl)-2-methyl-5-(3,4,5-tri methoxyphenyl)-2H-1,2,3-triazole (9) as a light yellow solid.

- NCI-60 cell line anti-cancer screening assay: All the 60 synthesized molecules were evaluated for their anti-cancer activity in a preliminary screen against a panel of 60 human cancer cell lines (NCI-60 panel) at a concentration of 10<sup>-5</sup> M utilizing the procedure described by Rubinstein et al.<sup>41</sup> In this cellular assay the growth inhibition of the test compounds is 65 measured by determining percentage cell growth (PG) inhibition. Optical density (OD) measurements of sulforhodamine B (SRB)-derived colour, just before exposing the cells to the test compound (ODt<sub>zero</sub>), and after 48 h exposure to the test compound (OD<sub>test</sub>) or the control vehicle 70 (OD<sub>ctrl</sub>) is recorded. The growth percentage compared to control is calculated utilizing the reported protocols.<sup>42</sup> From the preliminary screening potent compounds were further evaluated for five dose studies designed to determine GI<sub>50</sub> values, which represent the molar drug concentration required 75 for 50% cell growth inhibition. The compounds were dissolved in dimethyl sulfoxide (DMSO)/H2O and evaluated using five
- different concentrations at 10-fold dilutions (10<sup>-4</sup> M, 10<sup>-5</sup> M, 10<sup>-5</sup> <sup>6</sup> M, 10<sup>-7</sup> M and 10<sup>-8</sup> M) following 48 hr of incubation.
- 80 Screening against breast cancer cell lines (Hs578T) assay: Hs578T cells were seeded at 3,000 cells per well into 96-well plates and incubated overnight at 37 °C in DMEM containing 10% (v/v) FBS medium. The following day, various concentrations of triazole CA-4 analogues, or vehicle, were 85 added to each plate and the plates were then incubated for an additional 48 hr. Cells were washed twice with Dulbecco's phosphate-buffered saline (DPBS, pH 7.4) and incubated with 2 µM Calcein-AM for 30 min at 25 °C. Fluorescence was measured using a 490 nm excitation wavelength and 90 monitoring emission at 520 nm. The fluorescence intensity is proportional to the number of viable cells. The fluorescence intensity of control (untreated) cells was taken as 100% viability, and the relative cell viability compared to control was calculated.

Inhibition of tubulin polymerization assay: The tubulin polymerization was performed in 80 mM PIPES (pH 6.9) buffer containing 2 mM MgCl<sub>2</sub>, 0.5 mM EGTA, 100 mM GTP, and 10% (v/v) glycerol. The assay buffer was prepared on ice 100 immediately prior to performing the tubulin polymerization experiments in order to minimize GTP hydrolysis. Tubulin (4 mg) was reconstituted in 1 mL of cold assay buffer for a final concentration of 4 mg/mL. The reconstituted tubulin was placed on ice for a minimum of 3 minutes prior to starting the 105 experiment. Compounds were dissolved in 50% (v/v) DMSO/dH<sub>2</sub>O solution. The 96-well plate was pre-warmed to 37 °C for 30 minutes in a Synergy4 plate reader (BioTek, Winooski, VT). Compound (10x of final concentration) or DMSO control (4 µL) was added to the plate and incubated for 110 2 minutes at 37 °C. Cold tubulin (36 µL of 4 mg/mL) was added to each well and polymerization was initiated by placing the plate at 37 °C. Polymerization was measured over time by monitoring the change in absorbance at 340 nm. The initial portion of the reaction was fit to a linear equation and the slope

115 was used to estimate the rate of tubulin polymerization. The rate of polymerization in the presence of triazole analogue was

normalized against the control reaction and percent activity was plotted against the log of the inhibitor concentration. The  $IC_{50}$  values for test compounds were calculated using a variable slope dose-response curve.

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#### Graphical Abstract

## Synthesis and anti-cancer screening of novel heterocyclic-(2*H*)-1,2,3-triazoles as potential anticancer agents

Narsimha Reddy Penthala, Leena Madhukuri, Shraddha Takkar, Nikhil Reddy Madadi, Gauri Lamture,

Robert L. Eoff and Peter A. Crooks

Novel, stable combretastatin-A4 heterocyclic (2*H*)-1,2,3-triazole analogues displayed potent cytotoxic activity against both hematological and solid tumor cell lines with  $GI_{50}$  values in the low nanomolar range.

