Contents lists available at ScienceDirect

European Journal of Medicinal Chemistry

journal homepage: http://www.elsevier.com/locate/ejmech

Original article

Synthesis, anticancer activity and molecular docking studies on a series of heterocyclic *trans*-cyanocombretastatin analogues as antitubulin agents

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A R T I C L E I N F O

Article history: Received 16 August 2014 Received in revised form 21 November 2014 Accepted 28 December 2014 Available online 29 December 2014

Keywords: Trans-cyanocombretastatin analogues Anti-cancer activity Leukemia cell lines Molecular docking Tubulin binding

ABSTRACT

A series of heterocyclic combretastatin analogues have been synthesized and evaluated for their anticancer activity against a panel of 60 human cancer cell lines. The most potent compounds were two 3,4,5-trimethoxy phenyl analogues containing either an (Z)-indol-2-yl (8) or (Z)-benzo[b]furan-2-yl (12) moiety; these compounds exhibited GI₅₀ values of <10 nM against 74% and 70%, respectively, of the human cancer cell lines in the 60-cell panel. Compounds 8, and 12 and two previously reported compounds in the same structural class, i.e. 29 and 31, also showed potent anti-leukemic activity against leukemia MV4-11 cell lines with LD_{50} values = 44 nM, 47 nM, 18 nM, and 180 nM, respectively. From the NCI anti-cancer screening results and the data from the in vitro toxicity screening on cultured AML cells, seven compounds: 8, 12, 21, 23, 25, 29 and 31 were screened for their in vitro inhibitory activity on tubulin polymerization in MV4-11 AML cells; at 50 nM, 8 and 29 inhibited polymerization of tubulin by >50%. The binding modes of the three most active compounds (8, 12 and 29) to tubulin were also investigated utilizing molecular docking studies. All three molecules were observed to bind in the same hydrophobic pocket at the interface of α - and β -tubulin that is occupied by colchicine, and were stabilized by van der Waals' interactions with surrounding tubulin residues. The results from the tubulin polymerization and molecular docking studies indicate that compounds 8 and 29 are the most potent anti-leukemic compounds in this structural class, and are considered lead compounds for further development as anti-leukemic drugs.

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1. Introduction

A variety of natural products has been isolated from the bark of the South African tree *Combretum caffrum* which includes the combretastatins [1,2]. Combretastatins (Fig. 1, **I** and **II**) have been shown to be cytotoxic, with combretastatin A-4 (CA-4, Fig. 1, **I**) being the most potent [3,4]. CA-4 inhibits tubulin polymerization, and competitively inhibits the binding of radiolabeled colchicines to tubulin. CA-4 also exhibits potent cytotoxicity against a variety of cancer cell lines, and has been shown to be active against multidrug resistant (MDR) cancer cell lines [5–7].

Structurally related cyanocombretastatin analogues [8,9] (Fig. 1, **III**) and similar analogues that incorporate different heterocyclic moieties, such as indole, benzothiophene, quinoline and quinazoline have also been reported as cytotoxic compounds, and are potent inhibitors of tubulin polymerization potencies comparable to that of CA-4 [10–13]. In this respect, the 3,3-diarylacrylonitrile analog CC-5079 has been reported as a novel synthetic tubulin polymerization inhibitor with potential use in cancer chemo-therapy [14] and 2,3-diarylacrylonitriles have emerged as important synthons for the synthesis of a wide spectrum of biologically active molecules [15]. Such compounds have been shown to possess spasmolytic, estrogenic, hypotensive, antioxidant, tuberculostatic, antitrichomonal, and insecticidal properties [16].

Many natural medicinal agents, such as the combretastatins, the colchicines (Fig. 1, **IV**), and the podophyllotoxins, possess a





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Fig. 1. Chemical structures of potent antitubulin agents.

trimethoxyphenyl moiety in their structure. The cytotoxic properties of such compounds are believed to be related to their antitubulin properties [17–21]. A large number of CA-4 derivatives have been synthesized and evaluated for anti-tubulin activity [22–29]. SAR studies have revealed that a 3',4',5'-trimethoxyphenyl moiety and a cis-configuration of the olefinic bond in these compounds are essential structural elements for biological activity [30,31]. The presence of the *cis*-ethylene bridge that connects the two aryl rings (two planar rings tilted at 50-60⁰ to each other [1,31]) is believed to be the key structural factor that holds these structural moieties at an appropriate distance apart, to maintain the correct dihedral angle that maximizes interaction with the colchicine binding site on tubulin protein [32]. We have recently reported on the synthesis and anti-cancer activities of a series of (Z)-quinolinylcyano- (\mathbf{V}) and (E)- and (Z)-benzothiophene cyanocombretastatin analogues (Fig. 1, VI) [11,12], and have determined that such analogues (e.g. compound 29, Fig. 2) appear to overcome cell-associated P-glycoprotein (P-gp)-mediated resistance in tumor cells, since they are equipotent in inhibiting both OVCAR8 and NCI/ ADR-RES cell growth [11].

Previous studies on combretastatin analogues have shown that replacement of the 3-hydroxy-4-methoxyphenyl moiety of CA-4 with heterocyclic groups such as quinoline, quinazoline and benzothiophene results in improved anti-cancer activity [11,12,33]. In the present work, we have synthesized a variety of heterocyclic (*Z*)cyanocombretastatin analogues that incorporate 2- and 3-indolyl, 2- and 3-benzofuranyl, 2-benzothiophenyl, and 2-benzothiazolyl moieties as replacements for the 3-fluoro-4-methoxyphenyl group in (*Z*)-cyano CA-4 [9] (Fig. 1, **III**).

2. Results and discussion

2.1. Drug synthesis

(*Z*)-Indol-2-yl cyanocombretastatin analogues (**8**–**11**), (*Z*)-benzo [*b*]furan-2-yl cyanocombretastatin analogues (**12**–**14**) and (*Z*)-benzo[*d*]thiazol-2-yl cyanocombretastatin analogues (**15**–**17**) were synthesized by refluxing the appropriate indole-2-carbaldehyde (**1**), 4-cyanobenzyl substituted indole-2-carbaldehyde (**2**), benzo-[*b*]furan-2-carbaldehyde (**3**) or benzo[*d*]thiazole-2-carbaldehyde (**4**) with a variety of phenylacetonitriles, i.e., 3,4,5-trimethoxyphenylacetonitrile (**5**), 3,4-dimethoxyphenyl acetonitrile (**6**), and 3,5-dimethoxyphenyl acetonitrile (**7**), in 5% sodium methoxide/methanol (Scheme 1). Confirmation of the structure and purity of these analogues was obtained from ¹H NMR, ¹³C NMR and high resolution mass spectroscopic analysis. The geometry of the double bond (*E*- or *Z*-configuration) in these molecules was established as the (*Z*)-isomer from single crystal X-ray crystallographic data [34,35].

A second series of indole-3-yl (21-24), and (Z)-benzo[b]furan-



Fig. 2. Z-Benzo[b]thiophen-2-yl cyanocombretastatin analogues (29-31) [11].



1:	X=NH, Y=CH	; 2: X=N-CH ₂ -4-CN-C ₆	₅ H ₄ Y=CH
3:	X=O, Y=CH:	4: X=S, Y= N	

Compound	Х	Y	R^1	\mathbb{R}^2
8	NH	CH	-OCH ₃	-OCH ₃
9	NH	CH	Н	-OCH ₃
10	NH	CH	-OCH ₃	Н
11	-N-CH ₂ -C ₆ H ₄ - <i>p</i> -CN	СН	-OCH ₃	-OCH ₃
12	0	CH	-OCH ₃	-OCH ₃
13	0	CH	Н	-OCH ₃
14	0	CH	-OCH ₃	Н
15	S	Ν	-OCH ₃	-OCH ₃
16	S	Ν	Н	-OCH ₃
17	S	Ν	-OCH ₃	Н

Scheme 1. Synthesis of (*Z*)-indol-2-yl, (*Z*)-benzo[*b*]furan-2-yl, and (*Z*)-benzo[*d*] thia-zol-2-yl acrylonitriles (**8**–**17**).

3-yl cyanocombretastatin analogues (**25–27**) were synthesized by refluxing a variety of indole-3-carbaldehydes (**18–19**) or benzo[*b*] furan-3-carbaldehyde (**20**) with appropriate phenylacetonitriles (**5–7**) in 5% sodium methoxide/methanol (Scheme 2).

2.2. Biological evaluation

2.2.1. In vitro growth inhibition and cytotoxicity

All compounds were evaluated for their cytotoxic potency in a preliminary screen against a panel of 60 human cancer cell lines (NCI-60 panel) at a single analogue concentration (10^{-5} M) . The 60 cell line panel is organized into subpanels representing leukemia, non-small cell lung, colon, central nervous system, melanoma,



Scheme 2. Synthesis of (*Z*)-indole-3-yl, and (*Z*)-benzo[*b*]furan-3-yl cyanocombretastatins (**21–27**).

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Table 1

Table 2

Growth inhibition $(GI_{50}/\mu M)^a$ data for (*Z*)-indole-3-yl (**21–24**) and (*Z*)-benzo[*b*] furan-3-yl (**25**) cyanocombretastatin analogues.

μM μM μM μM μM μM μM Leakemia CCRF-CEM 0.017 0.052 0.136 0.024 0.777 0.055 na LE60(TB) <.001 0.039 0.039 0.010 0.467 0.054 0.33 MOLT-4 0.023 0.084 0.274 0.047 0.548 9.20 3.38 RMI-5226 <.001 0.151 0.199 0.019 1.32 0.237 na A549/ATCC <.001 0.114 0.076 <.001 0.520 0.379 na na HOP-92 <.001 0.399 1.45 <.001 3.15 3.25 2.84 NCI-H226 <.001 0.236 0.246 <.001 3.11 3.25 2.84 NCI-H226 <.001 0.033 <.001 0.472 0.333 2.97 8.39 NCI-H322 <.010 0.044 0.033 <.011 4.32 2.84 NCI-1522	Panel/cell line	8	9	11	12	13	15	17
Leukemia CCR-CEM 0.017 0.062 0.136 0.024 0.777 0.055 na HL-60(TB) <0.01 0.031 <0.01 0.481 0.066 0.345 RMI-8226 <0.01 0.039 0.039 <0.01 0.467 0.051 0.333 MOLT-4 0.023 0.084 0.274 0.047 0.548 9.20 3.33 MCI-122 <0.01 0.115 0.019 1.32 0.237 2.29 SR <0.01 0.042 na <0.01 0.470 0.107 0.402 Lung cancer <0.01 0.309 1.45 0.044 4.11 Nd.429 Nd.16 0.339 1.45 0.044 4.11 Nd.144 1.11 Nd.15 0.344 4.11 Nd.1423 <0.01 0.236 0.265 0.01 3.45 0.844 4.11 Nd.14232 <0.01 0.045 0.001 0.438 0.027 Ra Nd.142 0.333 0.01 4.423 <		μM	μM	μM	μM	μM	μM	μM
Leturema CCRR-CEM0.0170.0620.1360.0240.7770.055naHL-60(TB)<0.017	T							_
CCLPC-LEM0.0170.00240.1360.0240.770.0330.018HL-60(TR)0.0100.0330.0010.4610.0660.345K-552<0.01	Leukemia	0.017	0.002	0.120	0.024	0 777	0.055	
Introd(1b) Could Could <thcould< th=""> Could Could</thcould<>		<0.017	0.062	0.130	0.024	0.777	0.055	11d 0.245
MOLT-4 0.023 0.024 0.047 0.548 9.20 3.35 RPMI-S226 <0.01	K-562	<0.01	0.034	0.031	<0.01	0.467	0.000	0.343
RPMI-8226 (0.01) 0.151 0.199 0.019 1.32 0.237 2.29 SR (0.01) 0.042 na (0.01) 0.170 0.402 MS49/ATCC (0.01) 0.14 0.076 (0.01) 0.132 0.379 na REVX 1.16 0.307 na 0.092 (0.01) 2.29 1.93 3.35 HOP-62 (0.01) 0.33 0.077 (0.01) 1.43 nd 0.534 NCI-H226 (0.01) 0.033 0.01 0.438 0.229 1.00 NCI-H420 (0.01 0.033 (0.01 0.445 0.01 0.445 NCI-H522 (0.01 0.044 0.033 (0.01 0.448 0.022 0.444 HCC-1998 0.016 0.232 0.097 (0.01 0.448 0.033 0.33 SW-620 (0.01 0.048 0.497 0.035 0.445 0.414 HT29 (0.01 0.404	MOLT-4	0.023	0.033	0.055	0.047	0.407	9.20	3 38
SR COID 0.040 na COID 0.470 0.107 0.402 Lung cancer A549/ATCC <0.01	RPMI-8226	< 0.023	0.001	0 199	0.019	1 32	0.237	2.29
Lung cancer	SR	< 0.01	0.042	na	< 0.01	0.470	0.107	0.402
A>640/ATCC.0.010.1140.076.0.01.0.200.379naEKVX1.160.307na0.059na0.03na0.534HOP-62.0.010.2380.077.0.011.13nd0.534HOL-1423.0.010.2360.246.0.013.113.252.84NCI-H322nd0.7350.465.0.010.3360.3621.0332.28NCI-H420.0.010.0330.003.0.010.4720.3362.85NCI-H522.0.010.0490.033.0.010.4380.022naColo cancer0.010.4380.022naColo 205.0.0180.3660.2650.010.3450.3741.414HCT-116.0.010.0480.049.0.010.3160.2782.01SW-620.0.010.0480.047.0.010.3580.3332.33CNS Cancer	Lung cancer							
ICVX1.160.307na0.0020.0050.011.31nan.3HOP-92-0.010.0580.077-0.011.310.354NCI-H226-0.010.3281.45-0.010.3113.252.44NCI-H226-0.010.0330.033-0.010.4180.2252.010NCI-H322Mnd0.7350.465-0.010.4720.3362.36NCI-H420-0.010.0430.033-0.010.4380.2278.39HCI-1460-0.010.0440.047-0.010.4380.2798.39HCI-116-0.010.0440.047-0.010.3380.2978.39HCI-115-0.010.0440.047-0.010.3560.2782.01SW-620-0.010.0460.047-0.010.3580.0380.3130.33SW-620-0.010.0470.010.3740.3080.3430.343SF-295-0.010.0370.010.7080.582.465.539-0.010.3380.0111.560.99SNB-75-0.010.0360.039-0.011.780.1452.131.21M26A-0.010.0270.044-0.010.7030.731.48SNB-19-0.010.0360.011.780.1451.31U251-0.010.0280.011.780.1451.31U251-0.010.028	A549/ATCC	< 0.01	0.114	0.076	< 0.01	0.520	0.379	na
HOP-62<0.01na0.092<0.012.291.933.35HOP-92<0.01	EKVX	1.16	0.307	na	0.059	na	na	na
HOP-92<0.010.0580.077<0.011.13nd0.534NCI-H226<0.01	HOP-62	< 0.01	na	0.092	< 0.01	2.29	1.93	3.35
NCI-H226 <0.01 0.399 14.5 <0.01 3.4.5 0.8.4 4.11 NCI-H23 <0.01	HOP-92	<0.01	0.058	0.077	<0.01	1.13	nd	0.534
NCI-H23 <0.01 0.236 0.246 <0.01 5.18 2.29 >10.0 NCI-H420 <0.01	NCI-H226	<0.01	0.399	14.5	<0.01	34.5	0.844	4.11
NCI-H322M nd 0.735 0.465 <0.01 0.512 >510.0 NCI-H522 <0.01	NCI-H23	<0.01	0.236	0.246	< 0.01	3.11	3.25	2.84
NC1-H400 <0.01 0.053 0.039 <0.01 0.472 0.336 2.86 NC1-H522 <0.01	NCI-H322M	nd	0.735	0.465	< 0.01	6.18	2.29	>10.0
NC1-H522 c001 0.043 c003 c001 0.438 0.022 na COLO 205 0.016 0.232 0.097 -0.01 0.404 0.036 0.574 HCC-29298 0.018 0.056 0.011 0.248 0.049 0.0148 0.045 0.414 HT29 -0.01 0.048 0.089 -0.01 0.356 0.278 2.01 SW-620 -0.01 0.042 0.048 -0.01 0.356 0.278 2.01 SV-620 -0.01 0.042 0.048 -0.01 0.388 0.388 0.343 CNS Cancer - - - 0.01 0.76 0.447 -0.01 0.708 5.55 2.49 >1.00 SNB-75 -0.01 0.036 0.039 -0.01 1.78 0.145 2.13 U251 -0.01 0.056 -0.01 0.733 1.84 MALME-3M 8.28 nd nd 59.9 0.629 0.515	NCI-H460	<0.01	0.053	0.093	< 0.01	0.472	0.336	2.86
COLO 205 0.016 0.232 0.097 <0.011 0.404 0.036 0.574 HCC-2998 0.018 0.366 0.265 0.011 2.33 0.297 8.39 HCT-116 <0.01	NCI-H522 Colon cancor	<0.01	0.049	0.033	<0.01	0.438	0.022	na
COLD 205 0.018 0.366 0.265 0.017 0.404 0.036 0.33 0.297 8.39 HCT-116 0.01 0.044 0.047 0.01 0.448 0.045 0.414 HT29 <0.01	COLO 205	0.016	0 222	0.007	-0.01	0.404	0.026	0 574
Incc-106 O.010 O.044 O.020 O.011 D.13 O.023 O.014 O.024 O.011 O.148 O.023 O.037 O.038 O.038 O.038 O.038 O.338 O.338 O.338 O.338 O.338 O.338 O.338 O.339 O.31 I.48 O.856 O.247 O.01 D.788 O.588 Z.46 Sr-539 O.01 O.189 O.030 O.011 I.78 O.145 I.38 SNB-75 <0.01	HCC-2008	0.010	0.252	0.097	<0.01	0.404	0.030	0.574 830
Int Indication Count	HCT-116	<pre>0.018</pre>	0.300	0.203	<0.01	2.55	0.257	0.39
HT29 COII O.048 COUI O.037 O.0373 O.0374 O.011 SW-620 <0.01 O.042 O.044 <0.01 O.358 O.037 O.01 O.358 O.038 O.038 O.037 O.01 O.038 O.039 C.011 D.55 2.49 >1.00 SF-539 <0.01	HCT-15	< 0.01	0.044	0.047	< 0.01	0.469	0.032	0.404
KM12 c0.01 0.076 0.047 c0.01 0.356 0.278 2.01 SW-620 <0.01	HT29	< 0.01	0.048	0.005	< 0.01	0.105	0.037	0.438
SW-620 <0.01 0.042 0.048 <0.01 0.388 0.038 0.343 CNS Cancer SF-268 0.015 1.48 0.086 0.247 2.05 1.03 3.09 SF-255 <0.01	KM12	< 0.01	0.076	0.047	< 0.01	0.356	0.278	2.01
CNS Cancer Similar	SW-620	< 0.01	0.042	0.048	< 0.01	0.388	0.038	0.343
SF-268 0.015 1.48 0.086 0.247 2.05 1.03 3.09 SF-295 <0.01	CNS Cancer							
SF-295<0.010.0360.037<0.010.7080.5082.46SF-539<0.01	SF-268	0.015	1.48	0.086	0.247	2.05	1.03	3.09
SF-539<0.010.0470.039<0.011.560.0971.68SNB-19<0.01	SF-295	< 0.01	0.036	0.037	< 0.01	0.708	0.508	2.46
SNB-19<0.010.1890.085<0.015.552.49>10.0SNB-75<0.01	SF-539	< 0.01	0.047	0.039	< 0.01	1.56	0.097	1.68
SNB-75 <0.01 0.036 0.039 <0.01 1.78 0.145 2.13 U251 <0.01	SNB-19	< 0.01	0.189	0.085	< 0.01	5.55	2.49	>10.0
U251 <0.01 0.075 0.044 <0.01 0.760 1.65 1.91 Melanoma LOX IMVI <0.01	SNB-75	< 0.01	0.036	0.039	< 0.01	1.78	0.145	2.13
MelanomaLOX IMVI<0.01	U251	<0.01	0.075	0.044	<0.01	0.760	1.65	1.91
LOX IMVI <0.01 0.064 0.090 <0.01 0.930 0.073 1.84 MALME-3M 8.28 nd nd 59.9 0.629 0.515 0.699 M14 <0.01	Melanoma							
MALME-3M 8.28 nd nd 59.9 0.629 0.515 0.699 M14 <0.01	LOX IMVI	<0.01	0.064	0.090	<0.01	0.930	0.073	1.84
M14 <0.01 0.039 0.056 <0.01 0.437 0.193 1.21 MDA-MB-435 <0.01	MALME-3M	8.28	nd	nd	59.9	0.629	0.515	0.699
MDA-MB-435 <0.01 0.021 0.024 <0.01 0.229 0.021 0.129 SK-MEL-2 <0.01	MDA MD 425	<0.01	0.039	0.056	<0.01	0.437	0.193	1.21
SK-MILL2 (0.01 0.312 0.322 (0.01 0.437 0.537 0.537 SK-MEL-28 na 1.94 0.161 <0.01	NIDA-INIB-435	<0.01	0.021	0.024	<0.01	0.229	0.021	0.199
SK-MEL-5 0.101 0.101 0.001 0.511 1.37 1.37 UACC-257 1.06 27.0 0.062 63.2 18.0 >100 na UACC-62 <0.01	SK-MEL-2	<0.01 no	1.0/	0.528	<0.01	2.51	1.207	1.09
JANCL 257 1.06 27.0 0.062 63.2 18.0 >100 na UACC-62 <0.01	SK-MEL-20	~0.01	0.142	0.101	<0.01	0.564	0.100	1.30
UACC-62 <0.01 0.038 0.058 <0.01 0.557 0.052 0.755 Ovarian cancer IGROV1 <0.01	LIACC-257	1.06	27.0	0.000	63.2	18.0	>100	na
Ovarian cancer Idea Idea <thidea< th=""> Idea Idea</thidea<>	UACC-62	< 0.01	0.038	0.058	< 0.01	0.557	0.052	0.755
IGROV1 <0.01 0.482 0.183 0.022 1.27 1.27 0.706 OVCAR-3 <0.01	Ovarian cancer							
OVCAR-3 <0.01 0.034 0.042 <0.01 0.279 0.031 0.285 OVCAR-4 15.2 0.735 0.672 0.039 7.19 3.86 >10.0 OVCAR-5 0.08 0.471 0.259 0.042 5.23 0.347 4.85 OVCAR-5 0.08 0.471 0.259 0.042 5.23 0.347 4.85 OVCAR-8 <0.01	IGROV1	< 0.01	0.482	0.183	0.022	1.27	1.27	0.706
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	OVCAR-3	< 0.01	0.034	0.042	< 0.01	0.279	0.031	0.285
OVCAR-5 0.08 0.471 0.259 0.042 5.23 0.347 4.85 OVCAR-8 <0.01	OVCAR-4	15.2	0.735	0.672	0.039	7.19	3.86	>10.0
OVCAR-8 <0.01 0.326 0.124 <0.01 3.98 0.438 3.13 NCI/ADR-RES <0.01	OVCAR-5	0.08	0.471	0.259	0.042	5.23	0.347	4.85
NCI/ADR-RES <0.01 0.040 0.073 <0.01 0.317 0.059 1.47 SK-OV-3 <0.01	OVCAR-8	< 0.01	0.326	0.124	< 0.01	3.98	0.438	3.13
SK-OV-3 <0.01 0.394 0.087 <0.01 2.49 0.626 1.79 Renal cancer 786-0 <0.01	NCI/ADR-RES	<0.01	0.040	0.073	<0.01	0.317	0.059	1.47
Renal cancer 786-0 <0.01	SK-OV-3	<0.01	0.394	0.087	<0.01	2.49	0.626	1.79
786-0 <0.01 0.602 0.137 <0.01 5.84 1.06 2.44 A498 <0.01	Renal cancer							
A498 <0.01 0.040 0.029 <0.01 0.540 0.198 0.417 ACHN <0.01	786-0	< 0.01	0.602	0.137	< 0.01	5.84	1.06	2.44
ACHN <0.01 0.082 0.079 0.217 12.4 0.089 0.518 CAKI-1 <0.01	A498	< 0.01	0.040	0.029	< 0.01	0.540	0.198	0.417
CARI-1 <0.01 0.103 0.035 <0.01 0.492 0.047 0.482 RXF 393 <0.01	ACHN	< 0.01	0.082	0.079	0.217	12.4	0.089	0.518
KXF 393 <0.01 0.151 0.055 <0.01 1.30 0.148 0.551 SN12C <0.01	CAKI-1	< 0.01	0.103	0.035	< 0.01	0.492	0.047	0.482
SN12C <0.01 0.779 0.289 <0.01 0.576 0.315 4.88 TK-10 <0.01	KXF 393	<0.01	0.151	0.055	<0.01	1.30	0.148	0.551
IN-10 <0.01 5.36 20.1 73.6 2.28 0.063 2.78 UO-31 <0.01	SINTZC	<0.01	0.779	0.269	<0.01 72.6	0.950	0.515	4.08
Bit State cancer Solution Solution	10 21	<0.01	0.071	20.1	73.0	2.20	0.005	2.70
MCF7 0.016 0.289 0.039 0.023 0.446 0.034 0.399 MCF7 0.016 0.289 0.039 0.023 0.446 0.034 0.399 MDA-MB-231/ATCC 0.014 0.291 0.294 0.023 0.446 0.034 0.399 MDA-MB-231/ATCC 0.014 0.291 0.294 0.023 1.446 0.034 0.399 MDA-MB-231/ATCC 0.014 0.291 0.294 0.028 1.21 0.992 6.74 HS 578T <0.01	Prostate cancer	<0.01	0.071	0.201	2.29	3.75	0.085	0.079
DU-145 <0.01 0.206 0.207 <0.01 1.30 0.333 1.07 Breast cancer MCF7 0.016 0.289 0.039 0.023 0.446 0.034 0.399 MDA-MB-231/ATCC 0.014 0.291 0.294 0.028 1.21 0.992 6.74 HS 578T <0.01	PC-3	<0.01	0.062	0 105	<0.01	2 41	0 341	2 74
Breast cancer 0.016 0.289 0.039 0.023 0.446 0.034 0.399 MCF7 0.016 0.289 0.039 0.023 0.446 0.034 0.399 MDA-MB-231/ATCC 0.014 0.291 0.294 0.028 1.21 0.992 6.74 HS 578T <0.01	DU-145	< 0.01	0.206	0.207	< 0.01	1.30	0.333	1.07
MCF7 0.016 0.289 0.039 0.023 0.446 0.034 0.399 MDA-MB-231/ATCC 0.014 0.291 0.294 0.028 1.21 0.992 6.74 HS 578T <0.01	Breast cancer		0.200	0.207			0.000	
MDA-MB-231/ATCC 0.014 0.291 0.294 0.028 1.21 0.992 6.74 HS 578T <0.01	MCF7	0.016	0.289	0.039	0.023	0.446	0.034	0.399
HS 578T <0.01 0.133 0.092 <0.01 1.88 1.29 2.01 BT-549 <0.01	MDA-MB-231/ATCC	0.014	0.291	0.294	0.028	1.21	0.992	6.74
BT-549 <0.01 0.0978 0.109 0.029 0.950 12.2 3.45 T-47D 2.50 0.573 95.2 4.34 0.996 nd 1.18 MDA-MB-468 0.022 0.350 0.041 0.022 0.421 0.084 2.04	HS 578T	< 0.01	0.133	0.092	< 0.01	1.88	1.29	2.01
T-47D 2.50 0.573 95.2 4.34 0.996 nd 1.18 MDA-MB-468 0.022 0.350 0.041 0.022 0.421 0.084 2.04	BT-549	< 0.01	0.0978	0.109	0.029	0.950	12.2	3.45
MDA-MB-468 0.022 0.350 0.041 0.022 0.421 0.084 2.04	T-47D	2.50	0.573	95.2	4.34	0.996	nd	1.18
	MDA-MB-468	0.022	0.350	0.041	0.022	0.421	0.084	2.04

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.

Panel/cell line	21	22	23	24	25
	μM	μM	μM	μM	μM
Leukemia					
CCRF-CEM	0.058	2.74	0.042	0.064	1.41
HL-60(TB)	0.048	4.57	0.031	0.059	1.03
K-562	0.040	1.65	0.043	0.044	0.513
MOLT-4	0.131	3.99	0.049	0.198	3.82
KPMI-8226	0.231	3.96	0.065	0.241	2.47
SK Lung cancer	0.022	1,45	0.055	0.027	0.081
A549/ATCC	0.078	5.16	0.061	0.086	0.665
EKVX	0.596	18.5	0.478	0.485	na
HOP-62	0.082	8.69	0.273	0.133	6.98
HOP-92	1.01	5.26	0.068	0.083	3.02
NCI-H226	0.194	20.9	8.18	0.170	6.20
NCI-H322M	nd	243	0.230 nd	0.101	4.58
NCI-H460	0.041	4.51	0.039	0.037	0.839
NCI-H522	0.037	3.05	0.027	0.086	0.261
Colon cancer					
COLO 205	0.474	5.31	0.047	0.051	0.449
HCC-2998	0.107	12.6	0.207	0.119	4.13
HCT-15	0.048	3 70	0.042	0.045	0.317
HT29	0.038	3.32	0.038	0.038	0.394
KM12	0.043	4.04	0.040	0.044	0.586
SW-620	0.047	3.53	0.044	0.046	0.504
CNS Cancer					
SF-268	2.54	18.9	0.078	2.14	4.79
SF-295 SF-539	0.042	4.92 8.42	0.045	0.042	0.446
SNB-19	0.035	24.2	0.062	0.089	3.63
SNB-75	0.042	6.04	0.034	0.043	0.769
U251	0.050	4.97	0.045	0.050	0.605
Melanoma					
LOX IMVI	0.075	5.32	0.078	0.074	0.993
M14	12.5	3.75	0.042	5.85 0.052	0.868
MDA-MB-435	0.024	3.95	0.042	0.027	0.278
SK-MEL-2	na	4.70	0.051	na	0.304
SK-MEL-28	0.090	6.25	0.081	0.092	1.61
SK-MEL-5	0.042	4.27	0.048	0.036	0.336
UACC-257	11.3	11.9	nd	10.7	nd
Ovarian cancer	0.048	0.96	0.049	0.059	0.519
IGROV1	0.417	21.0	0.347	0.379	3.06
OVCAR-3	0.033	5.68	0.031	0.034	0.277
OVCAR-4	0.338	22.1	3.52	1.35	2.61
OVCAR-5	10.6	>100	0.197	0.516	3.33
OVCAR-8	0.121	7.38	0.103	0.148	2.90
SK-OV-3	0.059	1.70	0.035	0.057	0.582
Renal cancer	0.051	11.5	0.117	0.050	1.7 1
786-0	0.090	14.1	0.077	0.085	6.11
A498	0.035	3.36	0.035	0.025	0.458
ACHN	1.76	23.7	nd	1.66	1.94
CAKI-1	0.058	4.34	0.035	0.072	0.650
KAF 393 SN12C	0.036	4.48 8.10	0.083	0.033	0.496
TK-10	11.1	13.2	10.3	20.1	2.10
UO-31	1.41	14.3	0.663	1.26	2.89
Prostate cancer					
PC-3	0.054	7.85	0.075	0.054	1.84
DU-145 Proget concor	0.132	22.6	0.130	0.153	1.88
Breast cancer MCF7	0.034	3 10	0.041	0.036	0376
MDA-MB-231/ATCC	0.245	7.18	0.644	0.299	1.71
HS 578T	0.131	12.9	0.611	0.124	2.03
BT-549	1.94	5.38	0.049	0.348	0.927
T-47D	0.058	6.83	nd	0.065	0.811
MDA-MB-468	0.041	na	0.051	0.035	0.237

na: Not analyzed, nd: not determined, ^aGl₅₀: 50% growth inhibition, concentration of drug resulting in a 50% reduction in net cell growth as compared to cell numbers on day 0.

Growth inhibition $(GI_{50}/\mu M)^a$ data for (*Z*)-indol-2-yl (**8 11**), (*Z*)-benzo[*b*]furan-2-yl (**12, 13**), and (*Z*)-benzo[*d*]thiazol-2-yl (**15 17**) cyano combretastatin analogues.

ovary, renal, prostate, and breast cancer cell lines. The compounds were considered for progression to full five-concentration assay if they reduced the growth of cancer cells to 60% or more in at least eight of the 60 cell lines screened. The single dose results are expressed as the percent growth inhibition of treated cells at the test concentration of 10⁻⁵ M following 48 h of incubation. From the preliminary screens, compounds 8, 9, 11-13, 15, 17 and 21-25 were selected as leads for more comprehensive studies designed to determine GI₅₀, TGI and LC₅₀ values, which represent the molar drug concentration required to cause 50% growth inhibition, total growth inhibition, and the concentration that kills 50% of the cells, respectively. The compounds were evaluated using five different concentrations at 10-fold dilutions (10^{-4} M, 10^{-5} M, 10^{-6} M, 10^{-7} M and 10^{-8} M) and incubations were carried out over 48 h exposure to drug. Trans-cyanostilbenes analogues which contained the 3,4,5trimethoxyphenyl (8, 11, 12, 15, 21, 24 and 25) and dimethoxyphenyl (9, 13, 17, 22 and 23) moieties exhibited low micromolar level growth inhibition in subsequent five dose screening assays against all 60 human cancer cell lines in the panel. The growth inhibition results of the most potent of these compounds are presented in Tables 1 and 2. We hypothesized that the cytotoxic activity of these novel analogues was likely due to their interaction with the colchicine binding site on tubulin.

2.2.2. Indol-2-yl analogues

Compound **8** [(*Z*)-3-(1*H*-indol-2-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile] exhibited GI_{50} values ranging from <0.01 μ M to 1.16 μ M in 95% of the cancer cell lines screened, and showed potent growth inhibition ($GI_{50} = <0.01 \ \mu$ M) in 74% of the cancer cells in the 60-cell panel (Table 1).

The 3,4-dimethoxyphenyl acrylonitrile analog of **8**, compound **9**, which lacks the 5-methoxy group on the phenyl ring, also exhibited potent growth inhibition against 93% of the cancer cell lines in the panel, with GI_{50} values ranging from 0.021 to 0.779 μ M, and afforded an average GI_{50} value of 0.80 μ M against all the cancer cell lines in the panel (Table 1). This compound exhibited potent growth inhibition against MDA-MB-435 melanoma cancer cells with a $GI_{50} = 0.021 \ \mu$ M (Table 1).

The introduction of an *N*-(4-cyanobenzyl) group into the structure of compound **8** afforded compound **11** [(*Z*)-4-((2-(2-cyano-2-(3,4,5-trimethoxyphenyl)vinyl)-1*H*-indol-1-yl)-methyl) benzonitrile], which exhibited GI₅₀ values ranging from 0.024 μ M to 0.672 μ M in 95% of the cancer cell lines screened and showed potent growth inhibition properties in all five leukemia cell lines, with GI₅₀ values in the range 0.031–0.274 μ M. Compound **11** also showed potent growth inhibition against the MDA-MB-435 melanoma cell line (GI₅₀ = 0.024 μ M), and exhibited growth inhibition < 1 μ M against 93% of the cancer cells in the 60-cell panel (Table 1).

2.2.3. Benzofuran-2-yl analogues

Substitution of the 2-indolyl moiety in compound **8** for a benzofuran-2-yl moiety afforded compound **12** [(*Z*)-3-(benzofuran-2-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile], which exhibited good growth inhibition against all the cancer cells in the 60-cell panel with GI_{50} values ranging from <0.01 to 73.6 μ M. This compound exhibited remarkably potent growth inhibition against 70% of the cancer cell lines screened with $GI_{50} = <0.01 \ \mu$ M (Table 1).

Substitution of the 3,4,5-trimethoxyphenyl group in **12** for the 3,4-dimethoxyphenyl moiety afforded compound **13** [(*Z*)-3-(benzofuran-2-yl)-2-(3,4-dimethoxyphenyl)acrylonitrile], which exhibited good growth inhibition against 54% the cancer cells in the panel with GI₅₀ ranging from 0.229 to 0.996 μ M. This compound exhibited potent growth inhibition of MDA-MB-435 melanoma cancer cells with a GI₅₀ value of 0.229 μ M (Table 1). The average GI₅₀ value of this compound against all the cancer cell lines

screened was 2.59 µM.

2.2.4. Benzo[d]thiazol-2-yl analogues

The substitution of a benzofuran-2-yl moiety in compound **12** for a benzo[d]thiazol-2-yl moiety afforded compound **15** [(*Z*)-3-(benzo[*d*]thiazol-2-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile], which exhibited potent growth inhibition against 94% of the cancer cells in the 60-cell panel, with Gl₅₀ ranging from 0.021 to 12.2 μ M. The average Gl₅₀ value for this compound against all the cancer cell lines screened was 0.93 μ M. This compound exhibited potent growth inhibition against 73% of the cancer cell lines screened with Gl₅₀ = <1 μ M (Table 1) and exhibited potent growth inhibition against MDA-MB-435 melanoma cancer cells with a Gl₅₀ of 0.021 μ M (Table 1).

Substitution of the 3,4,5-trimethoxyphenyl group in **15** for the 3,5-dimethoxyphenyl moiety afforded compound **17** [(*Z*)-3-(benzo [*d*]thiazol-2-yl)-2-(3,5-dimethoxyphenyl)acrylonitrile], which exhibited potent growth inhibition against 94% of the cancer cells in the 60-cell panel with GI₅₀ ranging from 0.199 to 8.39 μ M. The average GI₅₀ value for this compound against all the cancer cells in the panel was 1.87 μ M. This compound exhibited potent growth inhibition against only 38% of the cancer cell lines screened with GI₅₀ = <1 μ M (Table 2). This compound exhibited potent growth inhibition against MDA-MB-435 melanoma cancer cells with a GI₅₀ value of 0.199 μ M (Table 1).

2.2.5. Indol-3-yl analogues

The indol-3-yl analog (**21**) of compound **8** exhibited good growth inhibition in the 60-cell screen with GI_{50} values ranging from 0.022 μ M to 12.5 μ M against all the cancer cell lines screened. Compound **21** showed potent growth inhibition against SR leukemia cells ($GI_{50} = 0.022 \,\mu$ M), and exhibited <1 μ M growth inhibition against 83% of the cancer cell lines screened (Table 2).

The indol-3-yl analog (**22**) of compound **9** exhibited growth inhibition against all the cancer cell lines screened with GI_{50} values ranging from 1.45 μ M to >100 μ M (Table 2).

Replacement of the 3,4-dimethoxyphenyl group in compound **22** with a 3,5-dimethoxyphenyl moiety afforded compound **23** [(*Z*)-3-(1*H*-indol-3-yl)-2-(3,5-dimethoxyphenyl)acrylonitrile], which exhibited good growth inhibition against all the cells in the panel with GI₅₀ values ranging from 0.020 μ M to 10.3 μ M. This compound showed potent growth inhibition against melanoma MDA-MB-435 cancer cells with a GI₅₀ value of 0.02 μ M. Compound **23** exhibited <1 μ M growth inhibition against 85% of the cancer cell lines screened (Table 2).

Replacement of the indol-3-yl group in compound **21** with a 5methoxyindol-3-yl moiety afforded compound **24** [(*Z*)-3-(5methoxy-1*H*-indol-3-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile], which exhibited good growth inhibition against all the cancer cells in the panel, with GI_{50} values ranging from 0.025 to 20.1 μ M. This compound exhibited potent growth inhibition against A498 renal cancer cells with a GI_{50} value of 0.022 μ M. Compound **24** exhibited <1 μ M growth inhibition against 88% of the cancer cell screened (Table 2).

2.2.6. Benzofuran-3-yl analogues

Replacement of the indol-3-yl group in compound **21** with a benzofuran-3-yl moiety afforded compound **25** [(*Z*)-3-(benzofuran-3-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile], which exhibited good growth inhibition against all the cancer cells in the panel, with GI₅₀ values ranging from 0.237 to 19.1 μ M. The average GI₅₀ value of this compound against all the cancer cell lines screened was 1.99 μ M. This compound exhibited potent growth inhibition against 52% of the cancer cell lines screened with GI₅₀ values <1 μ M (Table 2).

From structure-activity relationships (SAR) it is evident of that the heterocyclic series of *trans*-cyano CA-4 analogues containing benzo[*b*]thiophene-2-yl [11], indole-2-yl, benzofuran-2-yl and benzothiazole-2-yl moieties exhibited more growth inhibitory activity than the corresponding isomeric benzo[*b*]thiophene-3-yl, indole-3-yl, benzofuran-3-yl analogues. Also, analogues containing a 3,4,5-trimethoxyphenyl group generally exhibited better growth inhibition properties than analogues containing a 3,4dimethoxyphenyl or 3,5-dimethoxyphenyl moiety. Importantly, introduction of a 2-benzo[*b*]thiophenyl or 2-indolyl heterocyclic moiety in place of the 3-fluoro-4-methoxyphenyl moiety of *trans*cyano CA-4 [9] (Fig. 1, III) dramatically improved anti-cancer activity.

2.3. Anti-leukemic activity

Previously, we have reported on the synthesis of (*Z*)-benzo[*b*] thiophen-2-yl acrylonitriles (**29–31**) (Fig. 2) [11] as potent anticancer agents and we have evaluated the biological activity of these compounds against PC3 prostate cancer cell lines [11]. Compound **29** was also screened for its *in vitro* inhibitory activity on tubulin polymerization in these prostate cancer cells utilizing both an immunofluorescence assay and by using antibody against tubulin [11]. These benzo[*b*] thiophen-2-yl acrylonitrile compounds were also able to overcome P-glycoprotein (P-gp)-mediated resistance in PC3-DR prostate cancer cell lines and also exhibited a concentration-dependent anti-tubulin interaction in both PC3 and PC3-DR prostate cancer cells [11]. We have also reported that the thiophen-2-yl analogue **29** exhibits more potent growth inhibition than the isomeric thiophen-3-yl analogue **31** [11].

Compounds **29** and **30** both exhibited low nanomolar range growth inhibition ($GI_{50} < 10 \text{ nM}$) against all six leukemia cell lines in the NCI 60 cell line panel [11]. These results indicate that (*Z*)-benzo[*b*]thiophene analog **29** is a potential anti-leukemic compound.

In the current study, the novel cyanocombretastatin heteroaromatic analogues **8**, **11–17**, **21–25** and the previously reported (*Z*)-benzo[*b*]thiophene CA-4 analogs **29–31** [11] were evaluated for their anti-leukemic activity against the MV4-11 AML cell line (Table 3). From these studies, analogues **8**, **12** and **29** were determined to be the most active compounds against this leukemia cell line.

Compound **29** was further screened against a panel of 12 different primary leukemia cell lines, and showed a dose-dependent toxicity against most of the cell lines tested (Fig. 3). The average LD₅₀ value for the 12 different leukemia cell lines was 132 nM (range: 18.0–271 nM). Specifically, MV4-11 cells exhibited the most sensitivity to compound **29** (LD₅₀ = 18 nM), whereas THP-1 and MLL-ENL cells were the most resistant cell line (LD₅₀ = 227.0 and 271 nM, respectively) (Fig. 3).

We also tested the activity of **29** against primary blast crisis chronic myeloid leukemia (CML) cells and found that the response to **29** was also time-dependent (Fig. 4.). We observed that at 96 h more than 50% of the cells were apoptotic/dead after exposure to drug concentrations as low as 25 nM. We have also found that the activity of **29** was mediated via the activation of caspase cascades (MLG/HZ personal communication).



Fig. 3. Activity of compound 29 on 12 different leukemia cell lines.



Fig. 4. Activity of compound 29 in a primary leukemia sample (blast crisis of chronic myeloid leukemia) at 24 h, 48 h and 96 h time-points.

2.4. Tubulin polymerization inhibition

Based upon the 60-cell screening results from the leukemia cell lines (Tables 1 and 2) and the data from the *in vitro* toxicity studies on cultured AML cells (Table 3), seven compounds: **8**, **12**, **21**, **23**, **25**, **29** and **31**, were screened for their *in vitro* inhibitory activity on tubulin polymerization utilizing both an immunofluorescence assay and by using antibody against tubulin (a marker for dynamic microtubules) (Fig. 5) [36].

MV4-11 cells were treated with the above seven compounds at doses of 25, 50 and 100 nM for 2 h and cell-based tubulin depolymerization assays performed. The polymerized tubulin in the pellet (P) and unpolymerized tubulin in the supernatant (S) were



Fig. 5. Effect of *trans*-cyano CA-4 analogues 8, 12, 21, 23, 25, 29 and 31 to inhibit tubulin polymerization in MV4-11 cells.

Table	3

Anti-leukemic activity (LD₅₀) of the most potent compounds against the MV4-11 AML cell line.

	5 (50)	1 1	8					
Compound	8	11	12	13	14	15	16	17
LD ₅₀ (µM)	0.044	0.369	0.047	>20	1.169	0.233	1.223	4.339
Compound	21	22	23	24	25	29	30	31
LD ₅₀ (µM)	0.565	>20	0.375	0.467	4.529	0.018	6.063	0.180

detected by immunoblotting using antibody against tubulin. Compounds **8**, **12**, and **29** all demonstrated >50% inhibition of tubulin polymerization at 50 nM concentration (Fig. 5).

2.5. Molecular docking

The binding modes of three of the most active compounds (**8**, **12**, and **29**) were determined at the colchicine binding site on tubulin using *in silico* molecular docking protocols. Chemical structures of the molecules were drawn using Marvin Sketch (ChemAxon). Atomic coordinates for the α , β -tubulin heterodimer were derived from the PDB file 1SA0 of the crystal structure of tubulin-colchicine complex. Input coordinate files for both the protein and the compounds were generated using the Dock Prep module in the UCSF-Chimera software.

Docking was performed using SwissDock (http://www. swissdock.ch/), which employs the EADock DSS algorithm to generate binding modes [37], estimate CHARMM energies [38], account for solvent effects using the FACTS implicit solvation model [39], and rank binding modes with the most favorable energies.

The top binding poses for the above three compounds at the colchicine-binding site of tubulin are shown in Fig. 6. The molecules are structurally very similar, differing only in the nature of the heterocyclic moiety in the molecule. All three of these heterocyclic moieties are $10-\pi$ electron bicyclic aromatic ring systems. The structural difference between the three compounds is due to the nature of the heteroatoms in the fused 5-membered ring of the heterocycle: i.e. indole (8), benzofuran (12), and benzothiophene (29). Thus, an additional interest in carrying out the molecular docking studies was to assess the effect of switching the heterocyclic moiety in these compounds on tubulin binding characteristics. None of the compounds make any polar contacts with tubulin residues. Instead, all of them occupy the same hydrophobic pocket at the α,β -interface where colchicine binds, and are stabilized through numerous van der Waals' interactions with residues from both subunits. Compound 8 is stabilized through Van der Waal's interactions with Gln11, Thr179 and and Val181 of α-tubulin, and Lys352, Val318, Cys241, Leu248, Lys254 of β-tubulin (Fig. 6A). The two compounds 12 and 29 show identical binding modes (Fig. 6B and C) that involve interactions with the same set of residues. In both cases, the ring containing the heteroatom (benzofuran in 12 and benzothiophene in 29) is stabilized through Van der Waal's interactions with Asn101, Ser178, Thr179 and Val181 of α-tubulin (but not Gln11 as seen in compound 8), and Asn258 and Lys352 of β-tubulin. However, the heteroatom-ring in the two compounds is rotated by 180° with respect to one another. The 3,4,5trimethoxyphenyl moiety is stabilized by Cys241, Leu242, Leu248, Asp251, Lys254, Leu255, Met259, Val315 and Val318. Thus, the binding modes of compounds **12** and **29** are exactly superimposable on each other, except for their oppositely oriented heteroatoms ('O' in **12** and 'S' in **29**). The orientation of compound **8**, on the other hand, is not superimposable on those of **12** and **29**, although **8** also occupies the same binding pocket at the interface of α- and β-tubulin.

In summary, our molecular docking results were able to explain the trend in potencies observed for the three chosen compounds. This is further reflected in the free energy (Δ G) values of -8.12, -7.74 and -7.65, kcal/mol for compounds **29**, **12** and **8**, respectively.

3. Conclusions

Novel heterocyclic cyanocombretastatin A-4 analogues have been synthesized and evaluated for their anticancer activity against a panel of 60 human cancer cell lines. Compounds containing a trimethoxyphenyl moiety or a dimethoxyphenyl moiety showed potent growth inhibition, with GI_{50} values generally <1 μ M against most of the cancer cell lines used, with 8 and 12 being the most potent compounds. The novel cyanocombretastatin heteroaromatic analogues 8, 11–17, 21–25 and the previously reported (*Z*)-benzo[*b*] thiophene CA-4 analogs 29-31 were evaluated for their antileukemic activity against the MV4-11 AML cell line. Compounds 8, 12 and 29 also showed potent anti-leukemic activity against leukemia MV4-11 cell lines ($LD_{50} = 44$ nM, 47 nM, and 18 nM, respectively). The most active compound from the series, 29, was also screened against a variety of 12 different leukemia cell lines and exhibited LD₅₀ values <300 nM against all 12 leukemia cell lines. Compounds 8, 12, 21, 23, 25, 29 and 31 were also screened for their in vitro inhibitory activity on tubulin polymerization in MV4-11 cells and compounds 8, 12 and 29 all demonstrated >50% inhibition of tubulin polymerization at 50 nM concentrations. The binding modes of the three most active compounds, 8, 12 and 29 at the colchicine binding site on tubulin have been investigated utilizing molecular docking studies are consistent with the rank potency of these compounds as inhibitors of tubulin polymerization. From the cell screening and molecular docking results, we consider compounds 8 and 29 as lead compounds in the development of new anticancer agents that target tubulin. Compounds 8 and 29 are



Fig. 6. Binding modes of A) compound **8**; B) compound **12**; C) compound **29**; at the colchicine-binding site of tubulin. The inhibitors are shown as *purple* ball-and-sticks and the tubulin residues are shown as *orange* (α -tubulin) and *yellow* (β -tubulin) sticks. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

considered lead compounds suitable for further development as anti-leukemic drugs.

4. Experimental section

4.1. Chemistry

Melting points were recorded on a Kofler hot-stage apparatus and are uncorrected. TLC controls were carried out on pre-coated silica gel plates (F 254 Merck). ¹H and ¹³C NMR spectra were recorded on a Varian 400 MHz spectrometer equipped with a Linux workstation running on vNMRj software. All the spectra were phased, baseline was corrected where necessary, and solvent signals (CDCl₃) were used as reference for both ¹H and ¹³C spectra. HRMS data was obtained on an Agilent 6210 LCTOF instrument operated in multimode.

4.2. Methodology for the in vitro 60 human cancer cell screen

The methodology for the anti-cancer screening assay was carried out as per the reported literature procedure [40], which is also available at http://dtp.nci.nih.gov/branches/btb/ivclsp.html http:// dtp.nci.nih.gov/branches/btb/ivclsp.html.

4.3. Methodology for anti-leukemic activity determination

MV4-11 cells were cultured in Iscove's Modified Dulbecco's Media (IMDM) supplemented with 100 U/ml penicillin, 100 U/ml streptomycin, and 10% fetal bovine serum (Life technologies). Cells were seeded at 5 × 105/ml, and treated with the test compounds. At 24 and 48 h post treatment, cells were stained with annexin V-FITC (BD Biosciences) and 1 μ g/ml 7-AAD (Life Technologies). Percent cell dead was determined by flow cytometry as the percent of annexin V+ cells. Data were analyzed using Flowjo 9.3.2 for Mac OS X (TreeStar). The cell death was represented relative to vehicle control (DMSO).

4.4. Methodology for blast crisis chronic myeloid leukemia analyses

Primary blast crisis chronic myeloid leukemia (CML) samples were obtained with informed consent and IRB approval from Weill Cornell Medical College. The cryopreserved primary samples were thawed and cultured as previously described [41]. Cells were treated with indicated doses of compound. At 48 h post treatment, cells were stained with CD45-APC-H7 (BD Biosciences), followed by annexin V-FITC and 7-AAD staining. Cell dead was determined by flow cytometry. The percent cell dead of treated cells is represented by percent of annexin V+ blasts normalized to DMSO control.

4.5. Tubulin polymerization inhibition assay

MV4-11 cells were treated with indicated doses of compounds for 2 h. Cells were then lysed in microtubule-stabilizing buffer (100 mM Pipes, 1 mM EGTA, 1 mM MgSO₄, 30% glycerol, 5% DMSO, 1 mM DTT, 0.02% NaN₃, 0.125% NP-40, pH 6.9) at 37 °C. Free tubulin (supernatants, S) and polymerized tubulin (pellets, P) were examined by immunoblotting using tubulin antibody. Both microtubulestabilizing and microtubule-destabilizing drugs inhibit hypoxiainducible factor-1 alpha accumulation and activity by disrupting microtubule function [42].

4.6. General synthetic procedure: synthesis of (Z)-heterocyclic cyanocombretastatins (8–17 and 21–27)

A mixture of carbaldehyde (1.0 mole) and the appropriate

substituted phenylacetonitrile (1.1 mole) in 5% sodium methoxide/ methanol was heated under reflux for 3–6 h. The resulting solution was cooled to room temperature and poured into ice-cold water to afford a crude yellow solid. The solid was filtered off, washed with water, and finally washed with cold methanol. The obtained crude solid was recrystallized from methanol to afford the desired condensation product as a pure yellow crystalline solid.

4.6.1. (Z)-3-(1H-indol-2-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile (**8**)

mp: 147–148 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.83 (s, 6H), 3.93 (s, 3H), 6.73 (s, 2H), 6.77 (s, 1H), 7.08–7.25 (m, 4H), 7.57 (d, J = 8.0 Hz, 1H), 7.90 (brs, 1H, NH) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.23, 60.92, 102.78, 105.90, 111.54, 111.79, 112.73, 119.88, 120.97, 121.11, 121.48, 121.74, 127.37, 129.14, 130.67, 130.92, 132.26, 138.10, 138.99, 153.69 ppm. HRMS (ESI) m/z calcd for C₂₀H₁₉N₂O₃ [M+H]⁺ 335.1390; Found 335.1399.

4.6.2. (Z)-2-(3,4-Dimethoxyphenyl)-3-(1H-indol-2-yl)acryloni-trile (9)

mp: 170–172 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.92 (s, 3H), 3.97 (s, 3H), 6.90 (s, 1H), 6.92 (s, 1H), 7.11–7.16 (m, 1H), 7.22 (dd, *J* = 2, 8.4 Hz, 1H), 7.29–7.41 (m 1H), 7.42–7.44 (m, 2H), 7.62 (d, *J* = 8.4 Hz, 1H), 9.47 (brs, 1H, NH) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.04, 56.06, 105.89, 108.14, 111.60, 112.11, 118.57, 119.98, 120.94, 121.49, 125.36, 126.34, 127.41, 129.56, 132.52, 137.98, 149.42, 149.95 ppm. HRMS (ESI) *m*/*z* calcd for C₁₉H₁₇N₂O₂ [M+H]⁺ 305.1285; Found 305.1276.

4.6.3. (*Z*)-2-(3,5-Dimethoxyphenyl)-3-(1H-indol-2-yl)acrylonitrile (**10**)

mp: 195–197 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.84 (s, 6H), 6.46 (t, J = 2.0 Hz, 1H), 6.77 (d, J = 2.4 Hz, 2H), 6.94 (d, J = 2 Hz, 1H), 7.12 (t, J = 8.4 Hz, 1H), 7.29–7.33 (m, 1H), 7.42 (d, J = 8.8 Hz, 1H), 7.50 (s, 1H), 7.62 (d, J = 7.6 Hz, 1H), 9.48 (brs, 1H, NH) ppm. ¹³C NMR (100 MHz, CDCl₃): δ 55.54, 100.96, 103.70, 105.84, 111.70, 113.06, 119.85, 121.03, 121.66, 125.69, 127.34, 131.62, 132.22, 135.45, 138.17, 161.31 ppm. HRMS (ESI) m/z calcd for C₁₉H₁₇N₂O₂ [M+H]⁺ 305.1285; Found 305.1290.

4.6.4. (Z)-4-((2-(2-Cyano-2-(3,5-dimethoxyphenyl)vinyl)-1H-indo1-yl)methyl)benzonitrile (**11**)

mp: 190–192 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.84 (s, 6H), 3.85 (s, 3H), 5.51 (s, 2H, –CH₂), 6.64 (s, 2H), 7.11 (d, *J* = 8 Hz, 2H), 7.18–7.29 (m, 4H), 7.58 (d, *J* = 8.4 Hz, 2H), 7.74 (d, *J* = 8 Hz, 1H), 7.79 (s, 1H) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 46.33, 56.27, 60.99, 103.25, 107.35, 107.33, 109.27, 110.92, 111.98, 118.07, 118.14, 121.44, 122.50, 125.11, 126.61, 127.58, 127.88, 129.72, 132.73, 132.88, 138.23, 142.59, 153.62 ppm. HRMS (ESI) *m*/*z* calcd for C₂₈H₂₄N₃O₃ [M+H]⁺ 450.1826; Found 450.1821.

4.6.5. (Z)-3-(Benzofuran-2-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile (**12**)

mp: 102–104 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.90 (s, 3H), 3.91 (s, 6H), 6.89 (s, 2H), 7.26–7.31 (m, 1H), 7.36–7.40 (m, 1H), 7.41 (s, 1H), 7.50 (s, 1H), 7.53 (dd, *J* = 1.2, 10 Hz, 1H), 7.63–7.65 (m, 1H) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.53, 61.25, 103.30, 110.89, 111.10, 111.71, 117.60, 122.16, 123.81, 126.94, 127.66, 128.29, 129.17, 139.50, 151.21, 153.68, 155.20 ppm. HRMS (ESI) *m/z* calcd for C₂₀H₁₈NO4 [M+H]⁺ 336.1230; Found 336.1224.

4.6.6. (Z)-3-(Benzofuran-2-yl)-2-(3,4-dimethoxyphenyl)acrylonitrile (**13**)

mp:111–113 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.91 (s, 3H), 3.95 (s, 3H), 6.89 (d, *J* = 8 Hz, 2H), 7.14 (d, *J* = 2.4 Hz, 1H), 7.24–7.29 (m, 1H),

7.34–7.38 (m, 2H), 7.45 (s, 1H), 7.51 (d, J = 8.4 Hz, 1H), 7.60 (d, J = 7.6 Hz, 1H) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.17, 108.50, 110.43, 110.87, 111.45, 111.64, 117.70, 119.33, 122.05, 123.72, 126.36, 126.45, 126.70, 128.40, 149.52, 150.56, 151.54, 155.18 ppm. HRMS (ESI) m/z calcd for C₁₉H₁₆NO₃ [M+H]⁺ 306.1125; Found 306.1131.

4.6.7. (Z)-3-(Benzofuran-2-yl)-2-(3,5-dimethoxyphenyl)acrylonitrile (14)

mp: 142–144 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.84 (s, 6H), 6.49 (s, 1H), 6.81 (d, J = 2.0 Hz, 2H), 7.25 (t, J = 7.6 Hz, 1H), 7.37 (t, J = 7.6 Hz, 1H), 7.45 (s, 1H), 7.49 (s, H), 7.53 (d, J = 8.8 Hz, 1H), 7.62 (d, J = 8.0 Hz, 1H) ppm. ¹³C NMR (100 MHz, CDCl₃): δ 55.54, 101.65, 104.08, 110.79, 111.36, 111.63, 117.41, 122.08, 123.66, 126.89, 128.15, 128.48, 135.43, 151.07, 155.22, 161.27 ppm. HRMS (ESI) *m/z* calcd for C₁₉H₁₆NO₃ [M+H]⁺ 306.1125 Found 306.1126.

4.6.8. (*Z*)-3-(*Benzo*[*d*]*thiazo*l-2-*yl*)-2-(3,4,5-trimethoxyphenyl) acrylonitrile (**15**)

mp: 110–111 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.90 (s, 3H), 3.93 (s, 6H), 6.98 (s, 2H), 7.50–7.57 (m, 2H), 7.95–8.00 (m, 2H), 8.10 (d, J = 7.6 Hz, 1H) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.33, 61.0, 103.74, 110.0, 116.74, 116.97, 121.85, 123.88, 126.98, 127.05, 128.11, 133.82, 135.49, 140.37, 140.57, 152.51, 153.72, 161.33 ppm. HRMS (ESI) *m/z* calcd for C₁₉H₁₇N₂O₃S [M+H]⁺ 353.0960; Found 353.0968.

4.6.9. (Z)-3-(Benzo[d]thiazol-2-yl)-2-(3,4-dimethoxyphenyl)ac-rylonitrile (**16**)

mp: 130–131 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.94 (s, 3H), 3.95 (s, 3H), 6.93 (d, *J* = 8 Hz, 1H), 7.22 (s, 1H), 7.38 (d, *J* = 8 Hz, 1H), 7.48–7.55 (m, 2H), 7.91–7.96 (m, 2H), 8.08 (d, *J* = 8 Hz, 1H) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.08, 56.09, 108.39, 111.34, 116.79, 117.19, 120.47, 121.83, 123.59, 125.39, 126.88, 127.08, 131.90, 135.22, 149.58, 151.46, 152.02, 161.75 ppm. HRMS (ESI) *m*/*z* calcd for C₁₈H₁₅N₂O₂S [M+H]⁺ 323.0854; Found 323.0842.

4.6.10. (*Z*)-3-(*Benzo*[*d*]*thiazo*l-2-*y*l)-2-(3,5-*dimethoxypheny*l)acrylonitrile (**17**)

mp: 149–151 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.84 (s, 6H), 6.55 (s, 1H), 6.89 (s, 2H), 7.50–7.57 (m, 2H), 7.95–7.97 (d, J = 8 Hz, 1H), 8.04 (s, 1H), 8.11 (d, J = 8 Hz, 1H) ppm. ¹³C NMR (100 MHz, CDCl₃): δ 55.60, 103.01, 104.55, 116.67, 117.32, 121.87, 123.86, 127.09, 127.14, 134.54, 134.58, 135.45, 152.15, 161.29, 161.37 ppm. HRMS (ESI) *m/z* calcd for C₁₈H₁₅N₂O₂S [M+H]⁺ 323.0849; Found 323.0850.

4.6.11. (*Z*)-3-(1*H*-Indol-3-yl)-2-(3,4,5-trimethoxyphenyl)acrylonitrile (**21**)

mp:178–180 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.89 (s, 3H), 3.95 (s, 6H), 6.87 (s, 2H), 7.24–7.31 (m, 2H), 7.45 (d, *J* = 7.6 Hz, 1H), 7.77 (s, 1H), 7.78 (s, 1H), 8.42 (d, *J* = 2.8 Hz, 1H), 8.94 (brs, 1H, NH) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.32, 61.02, 102.79, 104.99, 111.77, 111.88, 117.95, 120.01, 121.27, 123.46, 126.25, 127.22, 130.71, 133.13, 135.54, 138.24, 153.58 ppm. HRMS (ESI) *m*/*z* calcd for C₂₀H₁₉N₂O₃ [M+H]⁺ 335.1390; Found 335.1387.

4.6.12. (*Z*)-2-(3,4-Dimethoxyphenyl)-3-(1H-indol-3-yl)acrylonitrile (**22**)

mp: 170–172 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.92 (s, 3H, –OCH₃), 3.97 (s, 3H, –OCH₃), 6.91 (d, J = 8.4 Hz, 1H), 7.15 (s, 1H), 7.24–7.21 (m, 3H), 7.44 (d, J = 7.6 Hz, 1H), 7.75 (s, 1H), 7.78 (s, 1H), 8.40 (s, 1H), 8.75 (brs, 1H, NH) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 55.97, 56.12, 104.92, 108.44, 108.55, 111.47, 111.73, 111.89, 117.90, 118.22, 120.14, 121.08, 121.24, 123.31, 125.90, 127.24, 127.83, 131.89, 135.53, 149.24 ppm. HRMS (ESI) m/z calcd for C₁₉H₁₇N₂O₂ [M+H]⁺ 305.1285; Found 305.1278.

4.6.13. (Z)-2-(3,5-Dimethoxyphenyl)-3-(1H-indol-3-yl)acrylonitrile (**23**)

mp: 190–192 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.86 (s, 6H), 6.45 (s, 1H), 6.82 (s, 2H), 7.25–7.29 (m, 2H), 7.44 (d, *J* = 8.0 Hz, 1H), 7.75 (d, *J* = 7.2 Hz, 1H), 7.87 (s, 1H), 8.44 (s, 1H), 8.83 (brs, 1H, NH) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 55.52, 100.10, 103.67, 111.79, 118.00, 119.95, 121.38, 123.50, 126.40, 127.28, 133.78, 135.45, 136.84, 161.21 ppm. HRMS (ESI) *m/z* calcd for C₁₉H₁₇N₂O₂ [M+H]⁺ 305.1285; Found 305.1297.

4.6.14. (Z)-3-(5-Methoxy-1H-indol-3-yl)-2-(3,4,5-trimethoxy-phenyl)acrylonitrile (**24**)

mp: 183–185 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.88 (s, 3H, –OCH₃), 3.89 (s, 3H, –OCH₃), 3.94 (s, 6H, –OCH₃), 6.85 (s, 2H), 6.93 (d, J = 8.8 Hz, 1H), 7.18 (s, 1H), 7.33 (d, J = 8.4 Hz, 1H), 7.69 (s, 1H), 8.38 (s, 1H), 8.60 (brs, 1H, NH) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 55.98, 56.37, 61.00, 100.47, 102.95, 104.74, 111.72, 112.52, 113.21, 120.02, 126.78, 127.93, 130.52, 130.80, 133.17, 153.60, 155.39 ppm. HRMS (ESI) m/z calcd for C₂₁H₂₁N₂O₄ [M+H]⁺ 365.1496; Found 365.1502.

4.6.15. (*Z*)-3-(*Benzofuran*-3-y*l*)-2-(3,4,5-trimethoxypheny*l*)ac-rylonitrile (**25**)

mp: 144–146 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.91 (s, 3H), 3.96 (s, 6H), 6.89 (s, 2H), 7.37–7.41 (m, 2H), 7.55 (s, 1H), 7.57 (d, J = 8.4 Hz, 1H), 7.72 (d, J = 7.6 Hz, 1H), 8.67 (s, 1H) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.82, 61.47, 103.72, 112.10, 112.48, 116.53, 119.01, 119.37, 123.97, 125.99, 126.67, 129.83, 129.98, 139.70, 146.35, 154.14, 155.35 ppm. HRMS (ESI) *m*/*z* calcd for C₂₀H₁₈NO₄ [M+H]⁺ 336.1236; Found 336.1218.

4.6.16. (Z)-3-(Benzofuran-3-yl)-2-(3,4-dimethoxyphenyl)acry-lo nitrile (**26**)

mp: 135–137 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.92 (s, 3H), 3.96 (s, 3H), 6.90 (d, J = 8.8 Hz, 1H), 7.13 (d, J = 2.4 Hz, 1H), 7.24 (dd, J = 2.4, 8.4 Hz, 1H), 7.33–7.40 (m, 2H), 7.49 (s, 1H), 7.54 (d, J = 8.4 Hz, 1H), 8.62 (s, 1H) ppm; ¹³C NMR (100 MHz, CDCl₃): δ 56.04, 56.08, 108.64, 111.38, 111.48, 111.96, 116.18, 118.66, 118.82, 118.92, 123.42, 125.42, 126.32, 126.57, 128.11, 145.55, 149.38, 150.13, 154.84 ppm. HRMS (ESI) m/z calcd for C₁₉H₁₆NO₃ [M+H]⁺ 306.1125; Found 306.1126.

4.6.17. (*Z*)-3-(*Benzofuran*-3-*y*l)-2-(3,5-dimethoxyphenyl)acry lo nitrile (**27**)

mp:121–123 °C, ¹H NMR (400 MHz, CDCl₃): δ 3.86 (s, 6H), 6.50 (s, 1H), 6.81 (s, 2H), 7.34–7.42 (m, 2H), 7.56 (d, *J* = 7.2 Hz, 1H), 7.62 (s, 1H), 7.69 (d, *J* = 8 Hz, 1H), 8.68 (s, 1H); ppm; ¹³C NMR (100 MHz, CDCl₃): 55.54, 101.06, 104.11, 111.48, 111.98, 116.03, 118.50, 118.91, 123.57, 125.52, 126.24, 130.40, 135.65, 146.12, 154.85, 161.29 ppm. HRMS (ESI) *m*/*z* calcd for C₁₉H₁₆NO₃ [M+H]⁺ 306.1125; Found 306.1133.

Acknowledgments

We are grateful to NCI/NIH (Grant Number CA 140409) and to the Arkansas Research Alliance (ARA) for financial support, M.L.G is funded by the US National Institutes of Health (NIH) through the NIH Director's New Innovator Award Program, 1 DP2 OD007399-01 and to the NCI Developmental Therapeutic Program (DTP) for screening data.

Appendix A. Supplementary data

Supplementary data related to this article can be found at http:// dx.doi.org/10.1016/j.ejmech.2014.12.050.

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