

# Benzothiazolo-quinazoline Congeners Function as Anti-microtubule Agents Triggering Mitotic Arrest in Cancer Cells

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## ABSTRACT

**Background:** Antimicrotubule agents effectively block cancer spread and are routinely employed as front-line chemotherapeutics. But patients, who receive Paclitaxel, suffer from peripheral neuropathy, which not only lowers quality of life but also can lead to treatment termination. Therefore, it is essential to identify novel microtubule inhibitors that can act as future front-line chemotherapeutics. In our previous study, we synthesized a cohort of 23 benzothiazolo[3,2-a]quinazoline derivatives employing a tandem copper catalyzed regioselective N-Arylation and aza-michael additions. However, the bioactive potential of the benzothiazolo-quinazoline derivatives are unknown. The synthesized benzothiazolo-quinazoline derivatives structurally resemble the microtubule inhibitors colchicine and CA-4. Therefore in this study; we asked whether benzothiazolo-quinazoline congeners act as microtubule inhibitors. **Materials and Methods:** To answer the question, we treated benzothiazolo-quinazoline congeners in a panel of four cancer cell lines and elucidated the molecular mechanism of action of the hit compounds *in vitro* and *in vivo*. **Results:** We found that benzothiazolo-quinazoline derivatives: 1-3, 6, 10, and 21-23 exhibited superior inhibitory effects with IC<sub>50</sub> values ranging from 4.2-12.1 μM in HeLa cells. We observed that 1-3, 6, 10, and 21-23 arrest cells in G2/M phase and disrupt microtubule network. In addition, we found that the selected congeners increase transcript and protein levels of mitotic markers. As part of mechanism, we observed that the congeners 1-3, 6, 10, and 21-23 promote tubulin depolymerisation similar to nocodazole, a microtubule depolymerising agent. Further, molecular modelling studies have shown that the congeners occupy the colchicines binding pocket in microtubule. We observed that the congeners 1-3, 6, 10, and 21-23 disrupt blood vessel formation in chick eggs and arrest Zebrafish embryo development. **Conclusion:** We conclude that the congeners 1-3, 6, 10, and 21-23 function as antimicrotubule agents. Based on observations, we suggest that the congeners can be lead-optimized to generate next generation of frontline chemotherapeutics.

**Keywords:** Mitosis, Benzothiazolo-quinazoline, Molecular modelling, Tubulin, Zebrafish embryos.

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## INTRODUCTION

Cancer is a malignancy that severely affects human health and well-being. Small molecules are routinely used as a treatment modality to ameliorate cancer and improve lifespan. The small molecules employed as anticancer agents belong into

two categories that are molecules employed as front-line chemotherapy or molecules used as precision medicine. In most developing nations, front-line chemotherapy is a standard set for treating benign or advanced cancers, as it is relatively cost-effective.<sup>1-4</sup> In comparison, precision medicine, offers more targeted therapy with better outcomes and lesser side effects, but is cost-prohibitive.<sup>1</sup> Therefore, it is imperative to discover better front-line chemotherapy to ameliorate cancer. Among the front-line chemotherapy, the microtubule inhibitor Paclitaxel is administered to manage a wide range of cancers.<sup>2</sup> Paclitaxel binds to the taxol binding site in the  $\alpha$ -Paclitaxel binds to the taxol



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binding site in the 5" thor><Year>2020</Year><RecNum>725</RecNum><DisplayText>.2,3 The disruption in tubulin homeostasis triggers cell death termed as mitotic catastrophe.<sup>4</sup> However, Paclitaxel induces peripheral neuropathy as a side effect of chemotherapy.<sup>5</sup> In addition, 60-70% of patients who receive Paclitaxel suffer peripheral neuropathy, which not only lowers quality of life but also usually results in treatment termination.<sup>6</sup> Therefore, it is essential to identify novel microtubule inhibitors that can act as future front-line chemotherapeutics.

In our previous study, we synthesized a cohort of 23 benzothiazolo[3,2-*a*]quinazoline derivatives employing a tandem copper catalyzed regioselective N-Arylation and aza-Michael additions.<sup>7</sup> N-heterocycles have been extensively useful in a multitude of areas ranging from healthcare to energy.<sup>8</sup> Fused heterocycles such as benzothiazole<sup>9</sup> and quinazoline derivatives<sup>10,11</sup> have received much attention as these compounds have exhibited excellent biological activities. Benzothiazolo-quinazolines are a class of N-fused heterocycles, essentially constituted by the fusion of the benzothiazole and the quinazoline moieties. Linearly fused benzothiazolo-quinazolines such as benzothiazolo[2,3-*b*]quinazolinones have proved to display antitumor activities and acted as electrochemical sensors/biosensors<sup>12-14</sup> However, despite their interesting structures, the angularly fused benzothiazolo-quinazolines such as benzothiazolo[3,2-*a*]quinazolines have not been explored in the biological studies. Considering the importance of benzothiazoles and quinazolines as potential anti-cancer therapeutics,<sup>9</sup> we envisaged that the hybrid structures incorporating benzothiazoles and quinazolines moieties may have potential anticancer profiles. The angularly fused benzothiazolo-quinazoline derivatives 1-22 (Chart 1), incorporating a sp<sup>3</sup> carbon tethered with ester/cyano/alcohol group, were synthesized using tandem copper catalyzed N-arylation-aza-Michael addition from the coupling partners 2-aminobenzothiazole and *ortho*-halo cinnamic acid derivatives. The diaryl thioether 23, having cyanamido and acrylate functional groups, was an intermediate in the synthesis of benzothiazolo-quinazoline 2. The syntheses of compounds 1-23 and their structural analyses were reported in our previous study.<sup>7</sup> However, the bioactive potential of the benzothiazolo-quinazoline derivatives is unknown. Moreover, benzothiazolo-quinazoline derivatives structurally resemble the microtubule inhibitors colchicines and CA-4.<sup>15,16</sup> Therefore, in this present study, we asked whether benzothiazolo-quinazoline derivatives act as microtubule inhibitors. To answer the question, we treated benzothiazolo-quinazoline congeners in a panel of four cancer cell lines and elucidated the molecular mechanism of action of the hit compounds *in vitro* and *in vivo*.

## EXPERIMENTAL PROCEDURES

### Cell cultures, maintenance, and antiproliferative evaluation

The cell lines employed to determine IC<sub>50</sub> values and perform molecular mechanism of action studies were obtained from ATCC. All the cell lines were grown in DMEM supplemented with 10% fetal bovine serum in carbon-dioxide incubator maintained at 37°C having 5% CO<sub>2</sub>. For determining IC<sub>50</sub> values, we employed the NCI-60 study protocol. In brief, the benzothiazolo-quinazoline analogues were used in a 5-dose assay wherein the concentrations range from 0.01 M with a fold-increase to 100 μM. The compounds were diluted in DMSO, which also served as vehicle control in our assays. The cell lines were plated at 10,000 cells per well in a 96-well plate using the liquid handling system (Janus, Perkin Elmer) and grown overnight to confluency. Before treatments, for time zero, we performed SRB assay on a 96-well plate. The cells were treated with 5-doses of compounds for duration of 48 hr. After 48 hr, 0.1 mL of 10% TCA was added to the cells and 96-well plates were incubated on ice for 1.5 hr. Later, the medium containing 10% TCA was discarded and the bounds cells were washed 4 times with demineralised water and kept in a hot air oven for 1.5 hr. To check the amount of cells, we stained the 96-well plates using 0.6% Sulforhodamine-B (SRB) obtained from (S4102, Sigma-Aldrich) for 30 min. After SRB staining, the plates were kept in a hot air oven for 1.5 hr and then dissolved in 50 microliters of 10 mM Tris, pH:8 solution. The IC<sub>50</sub> values were derived according to the NCI-60 protocol.<sup>17</sup>

### Cell cycle distribution

HeLa cells were plated onto 60 mm dishes were treated with 1-3, 6, 10, 21-23 and nocodazole (M1404, Sigma) mitosis for 18 hr. Later the cells were washed three times using 1x PBS, trypsinized, and spun down at 1500 rpm for 10 min at 4°C. After centrifugation, the pellets were incubated in 950 microliters of 70% ethanol and incubated on ice for 2 hr. Subsequently, the cells were spun down at 1500 rpm at 4°C and the supernatants were discarded, after 2 times 1x PBS wash, the cell pellets were mixed with propidium iodide- RNAase solution made in 1x PBS. The flow cytometry was performed in Amnis Flowsight and histograms were plotted.<sup>18</sup>

### Immunohistochemistry of tubulin and analysis of nuclear morphology

For immunohistochemistry analysis, HeLa cells were plated on 22 mm glass-coverslips and with 1-3, 6, 10, 21-23 and nocodazole for 18 hr. After incubation, the cells were fixed using 4% paraformaldehyde solution made in 1x PBS for 15 min at room temperature. Later the cells were washed three times using 1x PBS and subjected to permeabilization with 0.5% Triton X-100 and 0.05% Tween-20 solutions made in 1xPBS for a duration of 6 min. Subsequently, the cells were blocked using 5% BSA made in 1x PBS for 2 hr. Primary antibody against tubulin (T6074, Sigma)

was used at 1:500 dilutions for primary staining. The cells were washed three times using 1x PBS and subjected to secondary antibody staining. Here, we used the Cy3 conjugated secondary antibody (AP124C, Sigma) at a dilution of 1:1000 for 2 hr. The cells were again subjected to 1x PBS wash and mounted onto slides using mounting media contacting DAPI (F6057, Sigma). The images of stained cells were captured with Nikon Eclipse Ti microscope.<sup>19</sup>

### Western blot analysis of soluble versus polymerized tubulin

0.1 million HeLa cells were plated on 12-well plates overnight and treated with 1-3, 6, 10, 21-23, nocodazole or Paclitaxel (T7191, Sigma) for 18 hr. The cells were subjected to three times 1x PBS wash, and were ready to collect soluble and insoluble tubulin fractions. For soluble fractions, prewarmed lysis buffer was used as described previously. After incubation, the lysis buffer was aspirated and aliquoted into tubes containing 3x SDS-PAGE buffer. For insoluble fractions, we added 0.3 mL of 1 x SDS-PAGE to each well and harvested the samples. Both the soluble and insoluble fractions were heated in a boiling water bath for 5 mins. We loaded samples equally onto a 8% SDS-PAGE and subsequently transferred onto NC membranes using semi-dry transfer apparatus. 5% non-fat dry milk was added to the membranes and kept for 2 hr as blocking. Later, the membranes were incubated with tubulin antibody (T6074, Sigma) made in 1x TSBT at 1:2500 dilutions for 3 hr. The membranes were washed three times using 1x TBST and later incubated with 1:6000 dilutions of secondary antibody which was conjugated to HRP (A0545, Sigma) for 2 hr. The membranes were subjected to three washes and incubated with ECL solution and proteins were examined using ChemiDoc (LAS500, GE).<sup>20,21</sup> To determine Cyclin B1 (C8831, Sigma) and Aurora A (ZRB1902, Sigma) levels, HeLa cells treated with compounds were washed with 1x PBS, trypsinized, and subjected to Bradford assays. Equal amounts of protein were loaded onto SDS-PAGE gels and subjected to immunoblotting similar to the procedure followed to collect tubulin fractions.

### Quantitative RT-PCR

HeLa cells were seeded on 10 cm dishes and treated with DMSO, 1-3, 6, 10, 21-23 and nocodazole for 18 hr. Subsequently, TRI reagent (T9424, Sigma) was used to isolate total RNA as per manufacturer's procedures. The total RNA was run on 1% agarose gel and RNA levels were quantified using NanoDrop 2000, Thermo. cDNA was made using superscript-III synthesis kit and the cDNA was aliquoted into 0.2 mL PCR tubes. Quantitative PCR was conducted in a PCR apparatus (CX96F, BioRad). The  $\Delta\Delta C_t$  method was used to calculate fold change in mRNA levels and GAPDH was used for normalization.<sup>22,23</sup> Plk1 F: 5'-AAGAGATCCCGGAGGTCCTA-3';

R: 5'-AAGAGATCCCGGAGGTCCTA-3'; Nek2A  
F: 5'-AATGAATCGCATGTCCTACAAT-3'; R:

5'-CGTAATAATTTTCATTCAATTCATCAG-3'; GAPDH  
F: 5' AACCTGCCAAGTACGATGACATC-3', R:  
5'-GTAGCCCAGGATGCCCTTGA-3'

### ROS estimation

Compounds 1-3, 6, 10, 21-23 and nocodazole were used to treat HeLa cells which were plated onto 10 cm dishes. After treatments, the compound-treated cells were washed thrice with 1x PBS, trypsinized, and spun down for 2 min at 1500 rpm. The cell pellet were again washed and 20  $\mu$ M DCF-DA (35845, Sigma) was added for 30 min in a CO<sub>2</sub> incubator maintained 37°C. After 30 min, the cells were washed using 1x PBS and fluorescence intensity was measured at  $\lambda_{ex}$ =485 nm/ $\lambda_{em}$ =520 nm in a Amnis Flowsight Flow cytometer and graph was plotted using Graph pad prism software.<sup>18</sup>

### In silico molecular docking and ADMET

The protein tubulin-colchicine complex (PDB ID: 1SAO) was prepared and subjected to minimisation using "protein preparation" wizard from Schrodinger Suite using default parameters. For defining CBP using co-crystal colchicines, a receptor grid was constructed using "Receptor Grid Generation" wizard by taking default parameters and devoid of any constraints with a cube dimension of 25 Å. Simultaneously, the ligands were subjected to "ligprep" to generate possible ionized and minimized conformers employing default parameters. Subsequently the GLIDE software was employed, where the prepared ligands were docked into flexible SP (Standard Precision) and XP (Extra Precision) scoring functions taking the generated grid. Further, MM-GBSA (Molecular mechanics generalized Born surface area) studies were conducted to calculate the binding free energy of each protein complex using default parameters of PRIME module. The drug-like properties of these ligands were determined from data warrior and their respective toxicities were predicted using ProTox-II online server Schrödinger Release 2023-2: Maestro, Schrödinger, LLC, New York, NY, 2023.<sup>24,25</sup>

### Annexin V-FITC

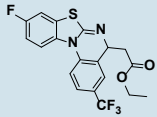
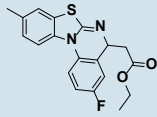
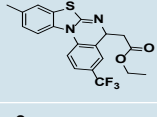
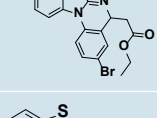
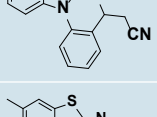
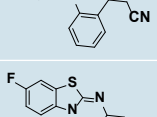
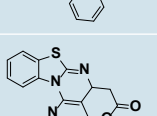
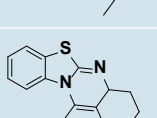
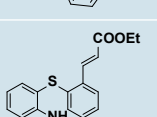
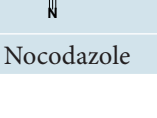
To detect for apoptosis, we employed the Annexin V methodology using the ApoAlert Apoptosis Detection Kit, Clontech. HeLa cells treated with indicated compounds were washed with 1x PBS and incubated with 10  $\mu$ L of Annexin V-FITC and 10  $\mu$ L of propidium iodide for 30 min. The stained cells were washed with 1x PBS and then mounted onto glass slides and images were captured using Nikon Ti Eclipse microscope.<sup>17</sup>

### Measurement of Mitochondrial membrane potential using JC-1

The cationic dye JC-1 (T4069, Sigma) was used to determine mitochondrial membrane potential ( $\Delta\Psi_m$ ). In the mitochondria of healthy cells, JC-1 accumulates and exhibits red fluorescence and when mitochondria are depolarised, JC-1 uptake is restricted

**Table 1: Anti-cancer activity of the compounds 1-23 IC<sub>50</sub> values are listed.**

Compound ID	Compound Structure	HeLa	MDA-MB-231	PC3	DU145
1		4.23±0.25	5.36±1.2	6.81±1.21	10.15±2.56
2		4.75±0.36	5.91±0.86	5.78±1.48	11.56±1.52
3		5.01±1.12	6.58±1.56	9.76±1.63	12.14±1.45
4		27.85±3.4	28.96±4.01	18.96±2.15	35.69±1.54
5		22.63±2.52	24.87±1.89	25.34±2.36	34.52±3.52
6		5.51±0.79	5.74±0.95	7.56±1.25	11.54±1.54
7		24.86±2.65	22.58±1.99	36.84±5.41	45.17±2.69
8		36.11±1.25	28.12±5.63	50.25±7.12	96.32±8.52
9		27.86±2.56	41.75±3.36	18.23±1.23	29.54±5.23
10		8.12±1.12	9.25±1.35	10.52±2.13	18.23±1.79
11		19.56±1.52	25.62±4.56	38.96±4.12	59.65±5.36
12		27.55±1.54	36.52±2.35	27.56±3.65	48.91±3.65
13		26.58±2.36	35.25±8.32	87.63±11.23	65.41±5.87

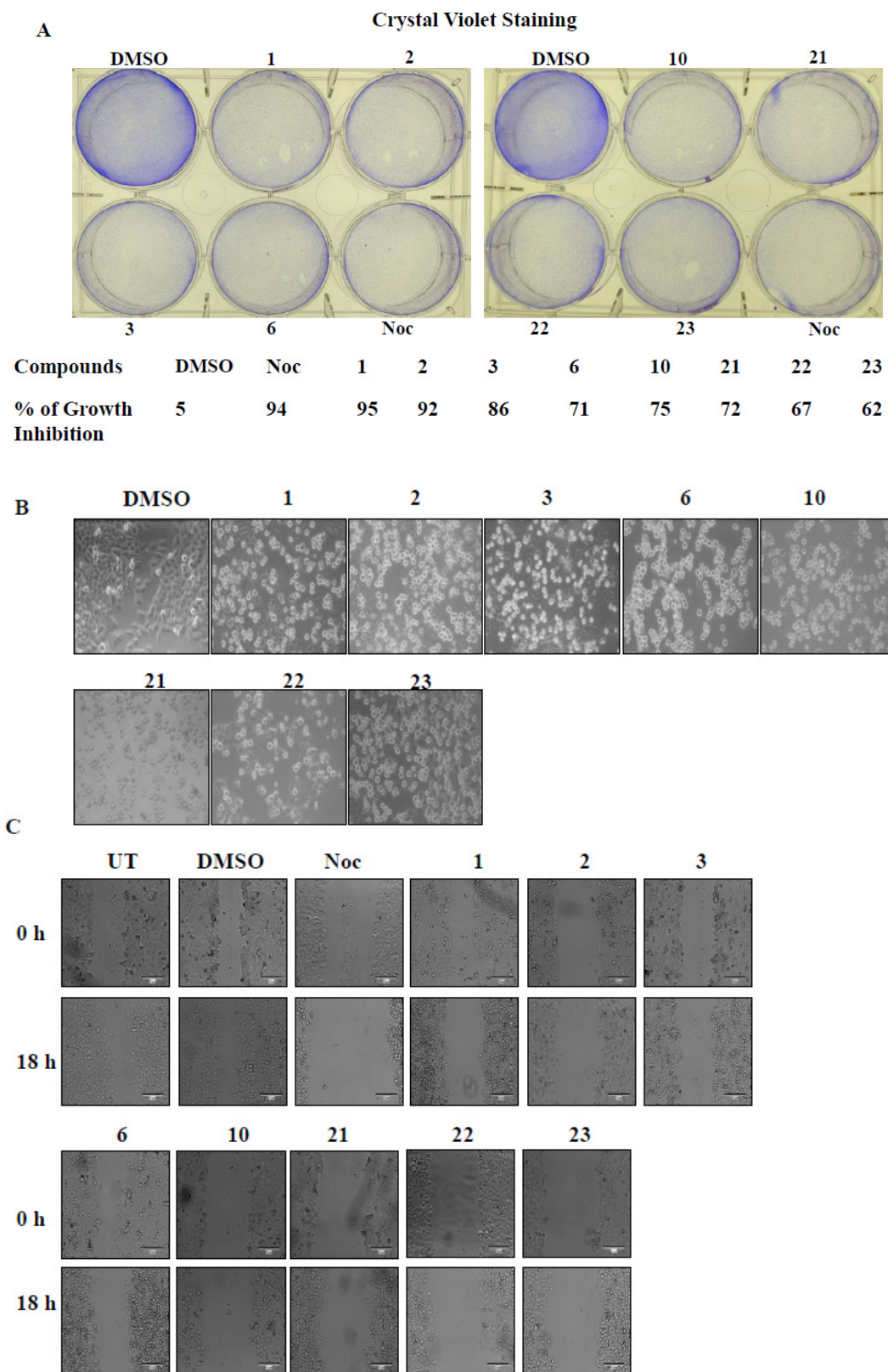
Compound ID	Compound Structure	HeLa	MDA-MB-231	PC3	DU145
14		26.84±3.69	37.23±3.65	35.12±4.63	68.11±4.65
15		35.24±2.36	27.12±1.75	56.58±9.84	47.15±4.52
16		33.58±3.54	35.63±6.48	18.14±1.29	65.23±5.36
17		22.54±1.12	45.65±2.85	36.78±4.21	54.35±7.86
18		26.69±1.75	41.25±1.25	36.89±4.56	78.56±12.56
19		31.25±1.36	30.56±2.69	58.96±7.96	45.63±8.12
20		18.89±1.15	19.38±1.75	20.77±1.45	63.89±6.45
21		8.85±1.65	9.77±1.36	11.23±2.56	15.65±1.42
22		10.12±1.63	14.56±2.10	17.81±1.63	23.56±1.45
23		9.35±1.70	9.89±1.14	10.11±1.25	12.41±1.25
24	Nocodazole	1.2±0.8	1.7±0.5	2.1±0.85	2.2±0.75

and shows green fluorescence. To perform JC-1 assays, HeLa plated on coverslips treated with compounds were washed using 1x PBS and stained employing 5 µg/mL JC-1 dyes. The stained cells were incubated at room temperature in dark for 30 min and later washed using 1x PBS and mounted on coverslips for confocal imaging (Nikon Ti Eclipse).<sup>17</sup>

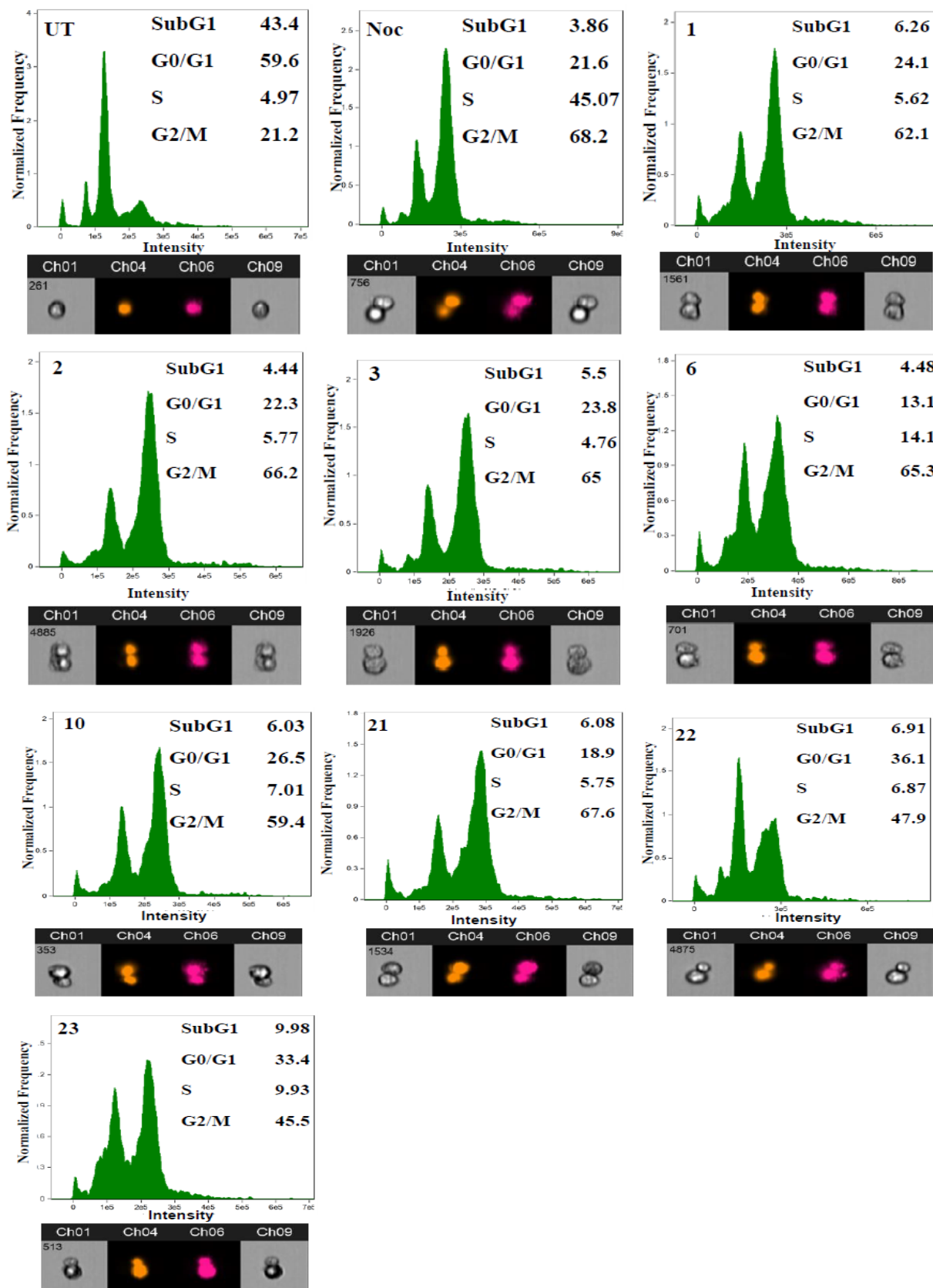
### Chick embryo angiogenesis assay

The assay is extensively employed in diverse applications in biology to investigate alterations in blood vessel creation through a

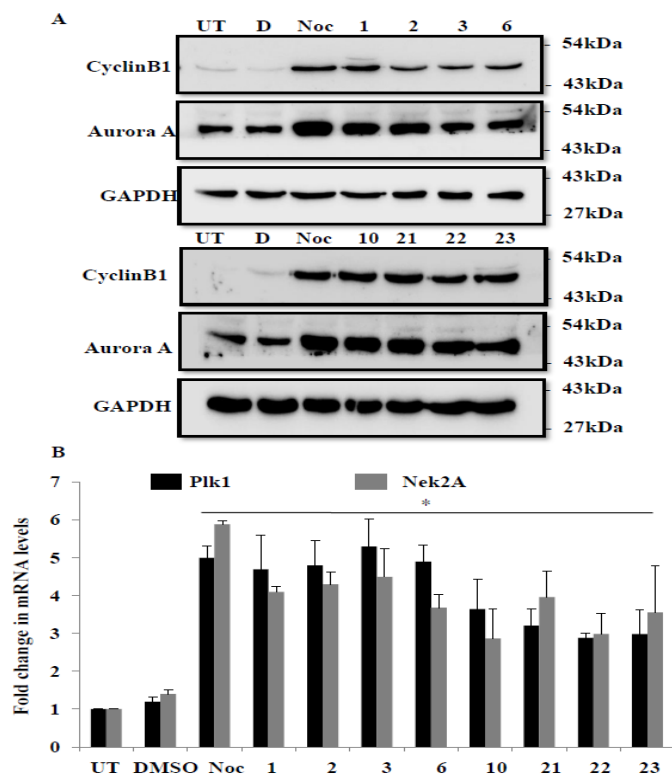
chick's developmental stages. Fertile chicken eggs were obtained from the ICAR-Directorate of Poultry Research, Rajendranagar, Hyderabad, India, in order to conduct the assay. The viable eggs were cultured at 37°C with 60-70% relative humidity. On the air sack side, a tiny opening was created on the fourth day of incubation. After the top of the shell was carefully cracked off, the chosen chemicals were applied. Additionally, the eggs were incubated in the incubator for an additional 4 hr following treatments. Stereomicroscope (S8APO, Leica) photographs of the



**Figure 1:** Benzothiazolo-quinazoline congeners 1-3, 6, 10, and 21-23 reduce HeLa cell growth-A, HeLa cells seeded on six-well plates were treated with 5  $\mu$ M of 1-3, 6, 10, and with 10  $\mu$ M 21-23 for 18 hr. Nocodazole (Noc) was used as a positive control, at 2.5  $\mu$ M concentrations. After 18 hr, cells were washed with 1x PBS, stained with crystal violet dye, and cell density was determined. The images are representative of three independent experiments. B, HeLa cells seeded on six-well plates were treated with 5  $\mu$ M of 1-3, 6, 10, and with 10  $\mu$ M of 21-23 for 18 hr. Images of cells were captured using phase-contrast microscopy,  $n=3$ . C, HeLa cells seeded on six-well plates overnight were scratched using a pipette tip and photographed. After capturing images at 0 hr, the cells were treated with 5  $\mu$ M of 1-3, 6, 10, and with 10  $\mu$ M 21-23 for 18 hr. The images were captured using phase-contrast microscopy,  $n=3$ .



**Figure 2:** Benzothiazolo-quinazoline congeners 1-3, 6, 10, and 21-23 trigger G2/M or mitotic arrest- A, HeLa cells seeded onto 60 mm dishes were treated with 5  $\mu$ M of 1-3, 6, 10, and with 10  $\mu$ M 21-23 for 18 hr. Nocodazole (Noc) was used as a positive control, at 2.5  $\mu$ M concentrations. After 18 hr, cells were washed with 1x PBS, fixed with 70% ethanol, stained with propidium iodide and cell cycle was determined using flow cytometry. The images are representative of three independent experiments. Representative images of captured cells are highlighted with each of the histogram,  $n=3$ .

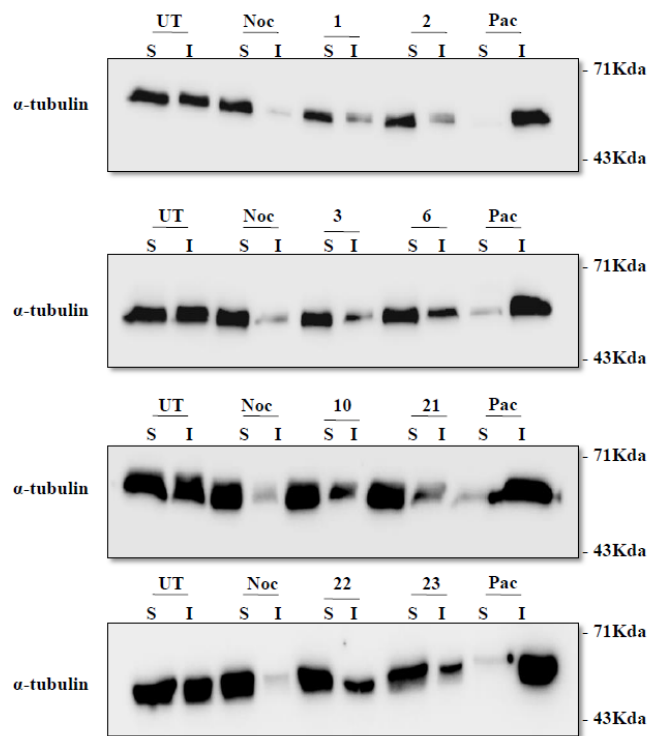


**Figure 3:** Benzothiazolo-quinazoline congeners induce accumulation of protein and transcripts of mitotic markers-A, HeLa cells seeded onto 60 mm dishes were treated with 5  $\mu\text{M}$  of 1-3, 6, 10, and with 10  $\mu\text{M}$  21-23 for 18 hr. Nocodazole (Noc) was used as a positive control, at 2.5  $\mu\text{M}$  concentrations. After 18 hr, cells were washed with 1x PBS, lysed using 1x RIPA buffer, and quantified with Bradford assays. Equal amounts of protein were resolved in 10% SDS-PAGE, and the protein levels of Cyclin B1 and Aurora A were detected using specific antibodies. GAPDH was used as normalization control. The images are representative of three independent experiments. B, under similar treatment conditions as A, the mRNA levels of Plk1 and Nek2A were analyzed by qPCR. After normalization to GAPDH, the gene expressions were represented as normalized mRNA fold change (mean $\pm$ SD), \* $p$ <0.05, and  $n$ =3.

treated and untreated eggs were taken at 0 and 4 hr, respectively. The ImageJ program was used to quantify the blood vessels.<sup>26,27</sup>

### Zebrafish screening

Wild type zebrafish (*Danio rerio*) were bred and kept under conditions of 28.5°C with 14 h:10 hr light-dark cycles. Three pairs of zebrafish mated simultaneously, and the resulting embryos were combined. After dechlorination, three to four embryos were placed in each well of a 48-well plate using 0.4 mL of E3 embryo medium. The embryos were incubated at 28.5°C after being treated with specific drugs at 10  $\mu\text{M}$  and 20  $\mu\text{M}$ , nocodazole at 2.5  $\mu\text{M}$  and 5  $\mu\text{M}$ , or DMSO (1%) at 5 hr after fertilization. In biological replicates, roughly 6-7 embryos were examined for every chemical. They were examined using bright-field microscopy (Olympus, IX73) 28 hr after conception. Only substances that produced comparable phenotypic abnormalities in the majority of the embryos were regarded as active.<sup>28,29</sup>



**Figure 4:** Benzothiazolo-quinazoline congeners act as microtubule depolymerising agents- HeLa cells were plated onto 12-well plates and treated with 5  $\mu\text{M}$  of 1-3, 6, 10, and with 10  $\mu\text{M}$  21-23 for 18 hr. Nocodazole (Noc) was used at 2.5  $\mu\text{M}$  concentrations and Paclitaxel (Pac) as at 1  $\mu\text{M}$  concentrations for 18 hr. The distribution of tubulin in soluble and insoluble fractions was analyzed by immunoblotting. The fractions containing soluble and polymerised tubulin were collected and separated by SDS-PAGE. Tubulin was detected by immunoblot analysis,  $n$ =3.

### Statistical analysis

The student's  $t$ -test was used to calculate statistical differences. Using a two-tailed  $t$ -test,  $p$  values for pair-wise comparisons were deduced,  $p$ <0.05 was considered significant.

## RESULTS

### Structure-activity Relationship of benzothiazolo-quinazoline congeners

In this study, *in vitro* inhibition of the benzothiazolo-quinazoline derivatives 1-22 and diaryl thioether 23 were tested towards HeLa, MDA-MB-231, PC3 and DU145 cell lines. The  $\text{IC}_{50}$  values are summarized in Table 1. The methyl 2-(5H-benzo[4,5]thiazolo[3,2-a]quinazolin-5-yl)acetate 1 exhibited excellent inhibition with  $\text{IC}_{50}$  values of 4.2  $\mu\text{M}$ , 5.4  $\mu\text{M}$ , 6.8  $\mu\text{M}$  and 10.5  $\mu\text{M}$  towards HeLa, MDA-MB-231, PC3 and DU145 cell lines, respectively. Similarly, 2-(5H-benzo [4, 5]thiazolo[3,2-a]quinazolin-5-yl)acetates 2 and 3 exhibited

excellent inhibition of HeLa, MDA-MB-231, PC3 and DU145 in the range of 4.7-12.1  $\mu\text{M}$  ( $\text{IC}_{50}$ ). It may be noted that the compounds 1-3 showed similar inhibitory effects across the four cell lines. However, benzothiazolo-quinazolines 4 and 5, containing isopropyl and methoxy substituents on the benzene ring of the benzothiazole moiety, exhibited inferior inhibitory effects. It must be noted that benzothiazolo-quinazoline 6, having two methyl substituents performed well with  $\text{IC}_{50}$  values of 5.5-11.5  $\mu\text{M}$ , which are similar to the compounds 1-3. Benzothiazolo-quinazolines 7-9, bearing electron withdrawing groups (F,  $\text{OCF}_3$ ) provided weak inhibition towards the four cell lines. Interestingly, benzothiazolo-quinazoline 10, with two fluorine groups on the benzene ring of the benzothiazole moiety displayed superior pan-inhibitory effects with  $\text{IC}_{50}$  values ranging from 8.1-18.2  $\mu\text{M}$ . Compounds having electron withdrawing groups (F,  $\text{CF}_3$ ) on the benzene ring of the quinazoline moiety such as 11 and 12 returned much lower  $\text{IC}_{50}$  values. Benzothiazolo-quinazolines 13-17, bearing substituent's on both the benzene rings of the benzothiazole and quinazoline moieties, also exhibited lower inhibition across the four cell lines. In the place of the ester group of the benzothiazolo-quinazoline, we employed a cyano group. Accordingly, cyano tethered benzothiazolo-quinazolines 18-20 were examined to observe low inhibitory activities. Note that N-doped benzothiazolo-quinazoline 21 worked well in this study to furnish promising  $\text{IC}_{50}$  values ranging from 8.8-15.6  $\mu\text{M}$ . Furthermore, alcohol tethered benzothiazolo-quinazoline 22, derived from the reduction of the ester group of 2, provided  $\text{IC}_{50}$  values in the range of 10.1-23.6  $\mu\text{M}$ . Lastly, we tested the intermediate 23, ethyl (E)-3-(2-((2-cyanamidophenyl)thio)phenyl)acrylate, to observe higher inhibitory values in the range of 9.3-12.4  $\mu\text{M}$ . Thus, we show that benzothiazolo-quinazoline derivatives: 1-3, 6, 10, and 21-23 exhibited superior inhibitory effects with  $\text{IC}_{50}$  values ranging from 4.2-12.1  $\mu\text{M}$ . To check that 1-3, 6, 10, and 21-23 indeed affected HeLa cell growth, we performed crystal violet staining and examined cell morphology. We found

that the congeners indeed decreased crystal violet staining and altered cell morphology, indicating that the potency of the congeners (Figure 1A-B). Further, the benzothiazolo-quinazoline congeners also blocked cell migration (Figure 1C). Overall, these observations prompted us to elucidate the in-depth mechanism of action of 1-3, 6, 10, and 21-23 compounds.

### Benzothiazolo-quinazoline congeners trigger G2/M arrest

The SAR analysis revealed that compounds 1-3, 6, 10, and 21-23 potently reduce the growth of cancer cells. A characteristic feature of antiproliferative compounds is that they alter cell cycle progression;<sup>30</sup> therefore, we asked whether these compounds also affect cell cycle progression. To answer this question, we treated HeLa cells 1-3, 6, 10, and 21-23 for 18 hr and performed cell analysis using imaging flow cytometry. As controls, we also treated HeLa cells with DMSO or nocodazole, a G2/M blocker for similar duration.<sup>21</sup> Here, we found that the 1-3, 6, 10, and 21-23 effectively arrested cell cycle progression in G2/M phase similar to nocodazole treatments. A representative imaging analysis also shows that benzothiazolo-quinazoline congeners treated cells have doubled DNA content compared to untreated or DMSO-treated cells (Figure 2). Thus, benzothiazolo-quinazoline congeners arrest cells in G2/M phase.

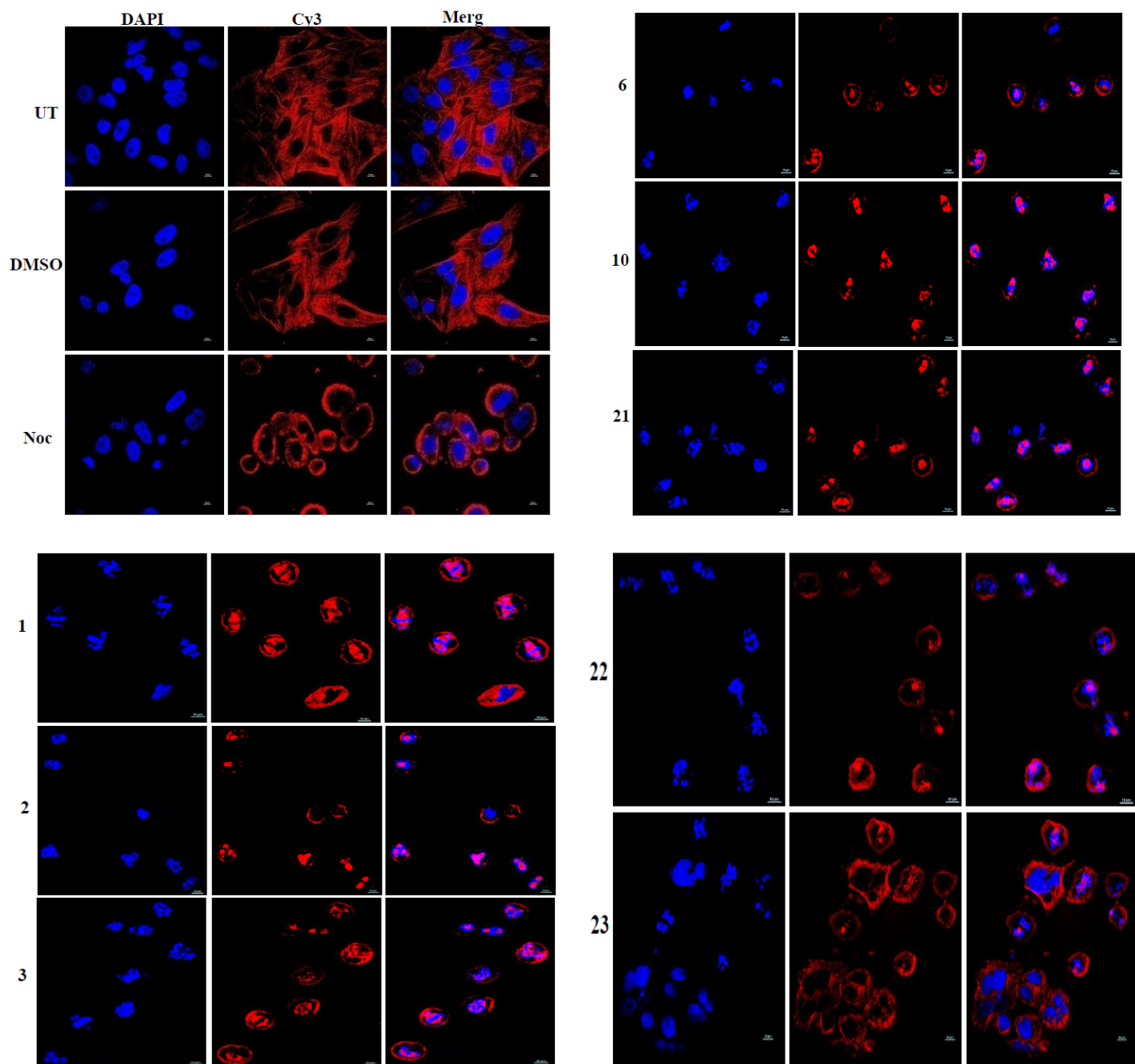
### Benzothiazolo-quinazoline congeners treated cells accumulate protein and transcript levels of mitotic markers

Cyclin B1/Cdk1 and Aurora A phosphorylates mitotic proteins to ensure the orderly progression of cell division. Further, accumulation of Cyclin B1 and Aurora A is a typical feature of mitosis;<sup>31</sup> therefore, we checked if 1-3, 6, 10, and 21-23 affect Cyclin B1 and Aurora A protein levels. We found that the selected benzothiazolo-quinazoline congeners increase Cyclin

**Table 2: Summarised *in silico* parameters of compounds, including ADMET properties.**

Sl. No.	Comp	<sup>a</sup> DS (kcal/mol)	<sup>b</sup> GEM (kcal/mol)	<sup>c</sup> $\Delta\text{G}_{\text{bind}}$ (kcal/mol)	ADME properties (Lipinski rule of five)						Toxicity (Protox-II server)		
					<sup>d</sup> MW	<sup>e</sup> HBD	<sup>f</sup> HBA	<sup>g</sup> cLogP	<sup>h</sup> TPSA	Breach of rule	<sup>i</sup> LD <sub>50</sub> (mg/kg)	Toxicity class	Toxicity target
1	1	-7.34	-71.69	-105.00	324.403	0	4	3.168	67.2	0	600	4	None
2	2	-7.22	-66.10	-76.51	310.376	0	4	2.7617	67.2	0	700	4	None
3	3	-6.77	-68.74	-105.49	338.43	0	4	3.5119	67.2	0	700	4	None
4	6	-6.8	-65.36	-70.39	352.457	0	4	3.8558	67.2	0	2900	5	None
5	10	-7.38	-68.74	-49.52	360.383	0	4	3.3696	67.2	0	700	4	None
6	21	-7.32	-61.34	-78.38	325.391	0	5	2.5185	80.09	0	1100	4	None
7	22	-7.19	-59.26	-98.28	324.403	1	4	3.4396	87.42	0	4000	5	None
8	23	-6.74	-68.85	-71.93	282.366	1	3	2.6834	61.13	0	2900	5	None

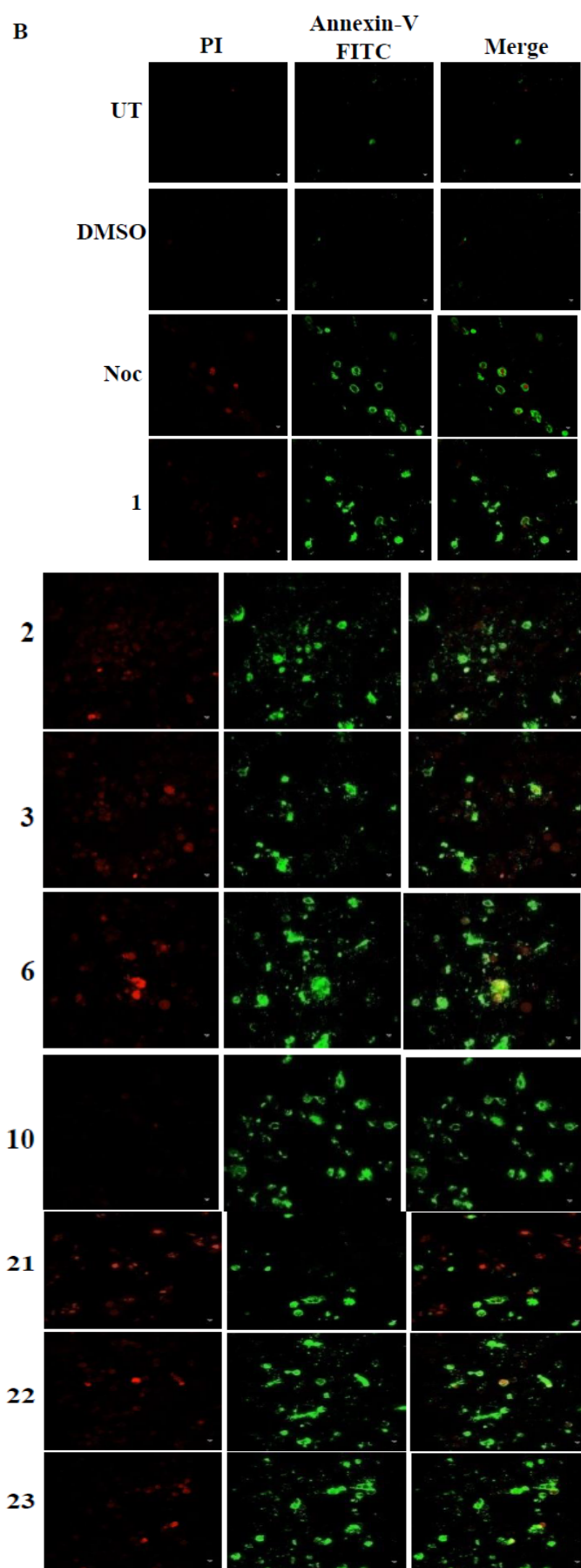
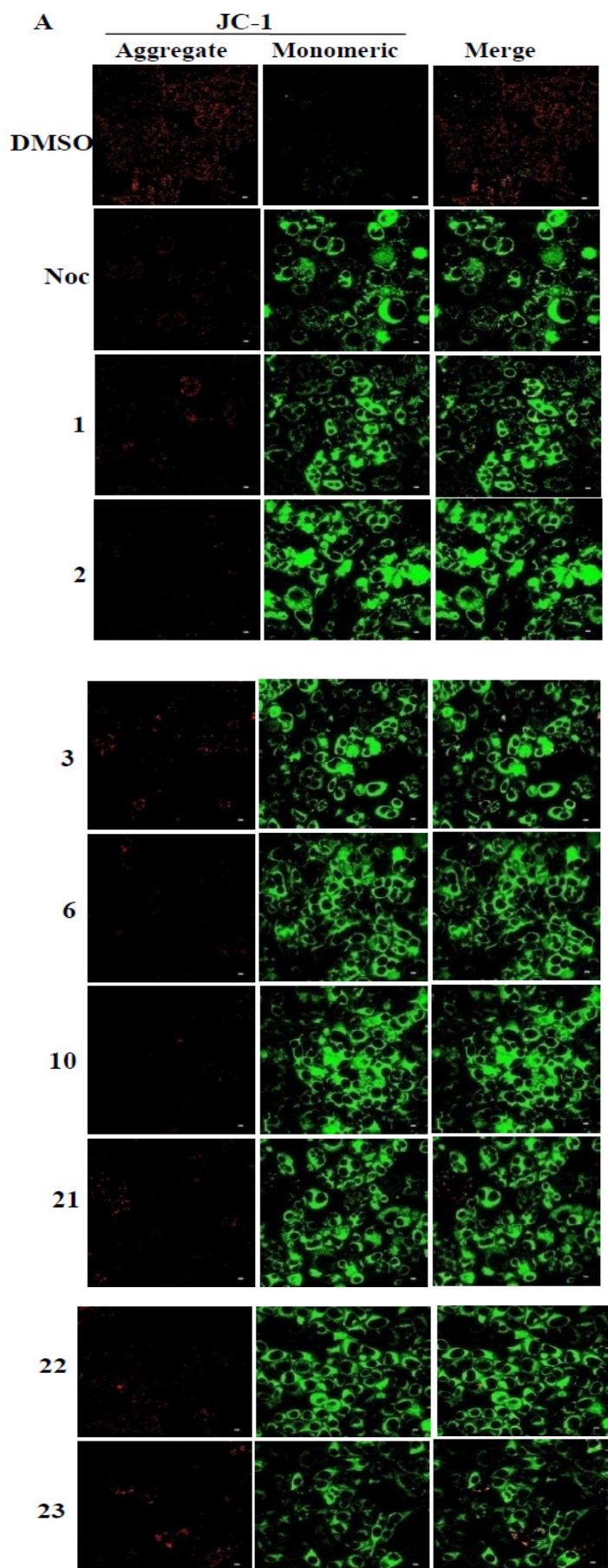
<sup>a</sup>Dock score, <sup>b</sup>Glide Emodel, <sup>c</sup>MM-GBSA or binding free energy, <sup>d</sup>Molecular Weight, <sup>e</sup>H-bond donor, <sup>f</sup>H-bond acceptor, <sup>g</sup>Hydrophilicity, <sup>h</sup>Topological Polar Surface Area, <sup>i</sup>lethal dose, 50%.

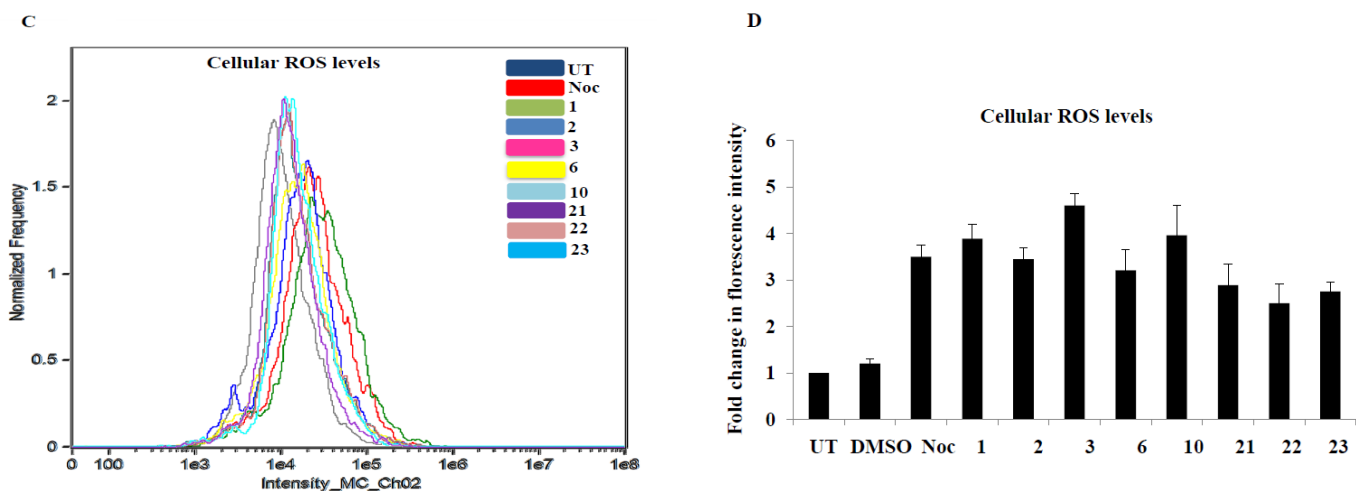


**Figure 5:** Benzothiazolo-quinazoline congeners disrupt microtubule network- A, HeLa cells seeded on 18 mm coverslips and were treated with 5  $\mu$ M of 1-3, 6, 10, and with 10  $\mu$ M 21-23 for 18 hr. Nocodazole (Noc) was used as a positive control, at 2.5  $\mu$ M concentrations After 18 hr, cells were washed with 1x PBS, followed by fixation, permeabilization, and indirect immunofluorescence analysis was performed. The cells were incubated with anti- $\alpha$ -tubulin primary antibody and stained using Cy3-conjugated secondary antibody. Nuclei were stained with DAPI. The merged images of cells stained for tubulin and DAPI are shown. The images are representative of three independent experiments.

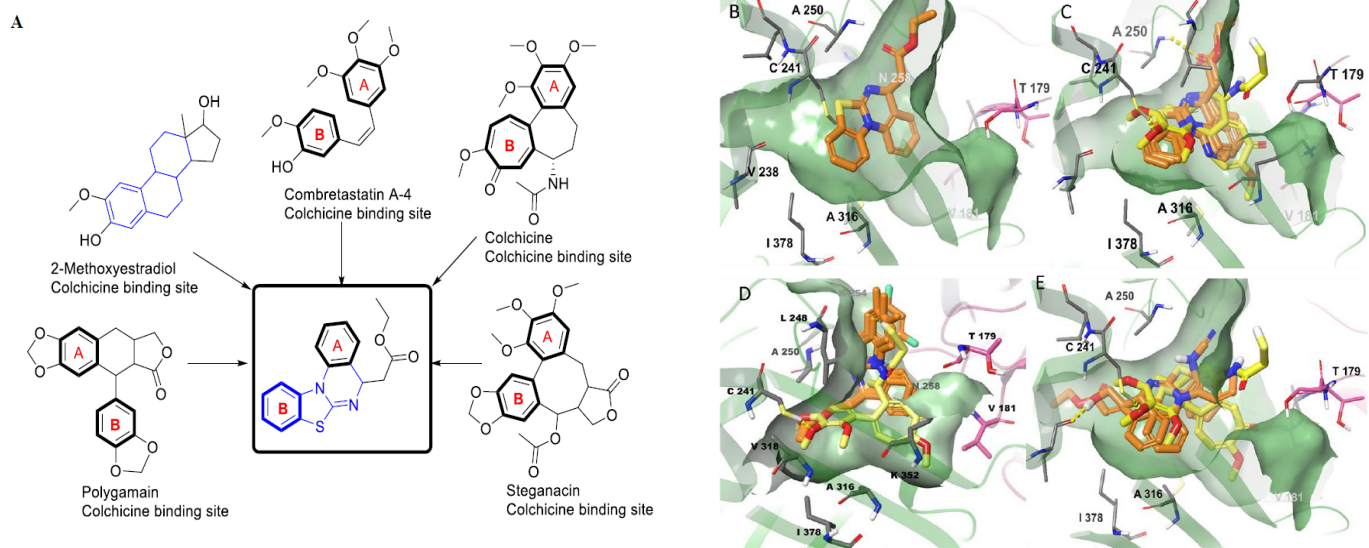
B1 and Aurora A protein levels similar to nocodazole (Figure 3A). To corroborate that benzothiazolo-quinazoline congeners indeed arrest cells in G2/M, we examined the transcript levels of mitotic markers including Plk1 and Nek2A in HeLa cells treated with selected congeners.<sup>32</sup> We observed that the selected

benzothiazolo-quinazoline congeners strongly increase the transcript levels of mitotic markers similar to nocodazole-treated cells (Figure B). Overall, the compounds 1-3, 6, 10, and 21-23 increase Cyclin B1 protein and transcript levels of mitotic markers.





**Figure 6:** Benzothiazolo-quinazoline congeners increase mitochondrial depolarization, induce Annexin V, and activate cellular reactive oxygen species (ROS) levels and staining A, HeLa cells seeded on 18 mm coverslips and were treated with 5  $\mu\text{M}$  of 1-3, 6, 10, and with 10  $\mu\text{M}$  21-23 for 18 hr. Nocodazole (Noc) was used as a positive control, at 2.5  $\mu\text{M}$  concentrations. After 18 hr, the cells were treated with JC-1 and confocal microscopy was performed to determine mitochondrial polarization. Depolarized mitochondria emit green fluorescence and polarized mitochondria emit red fluorescence. PI represents Propidium iodide. Representative images from three independent experiments are shown  $n=3$ . B, under similar treatment conditions of A, apoptosis was assessed using the ApoAlert Annexin V Apoptosis Detection Kit, Clontech according to the manufacturer's instructions. Representative images from three independent experiments are shown  $n=3$ . C-D, under similar treatment conditions of A, after treatments, the cells were centrifuged at 1000 rpm for 1 min. The centrifuged cells were washed using 1 x PBS and incubated with 20  $\mu\text{M}$  DCF-DA for 20 min at 370C in a  $\text{CO}_2$  incubator. The incubated cells were washed with 1x PBS and fluorescence intensity was measured at  $\lambda_{\text{ex}}=485 \text{ nm}/\lambda_{\text{em}}=520 \text{ nm}$  in an Amnis Flowsight Flow cytometer and the mean fluorescence intensity was calculated. C is a representative image of ROS intensity calculated using flow cytometry, and D, is the graph plotted as fold change of fluorescence intensity.

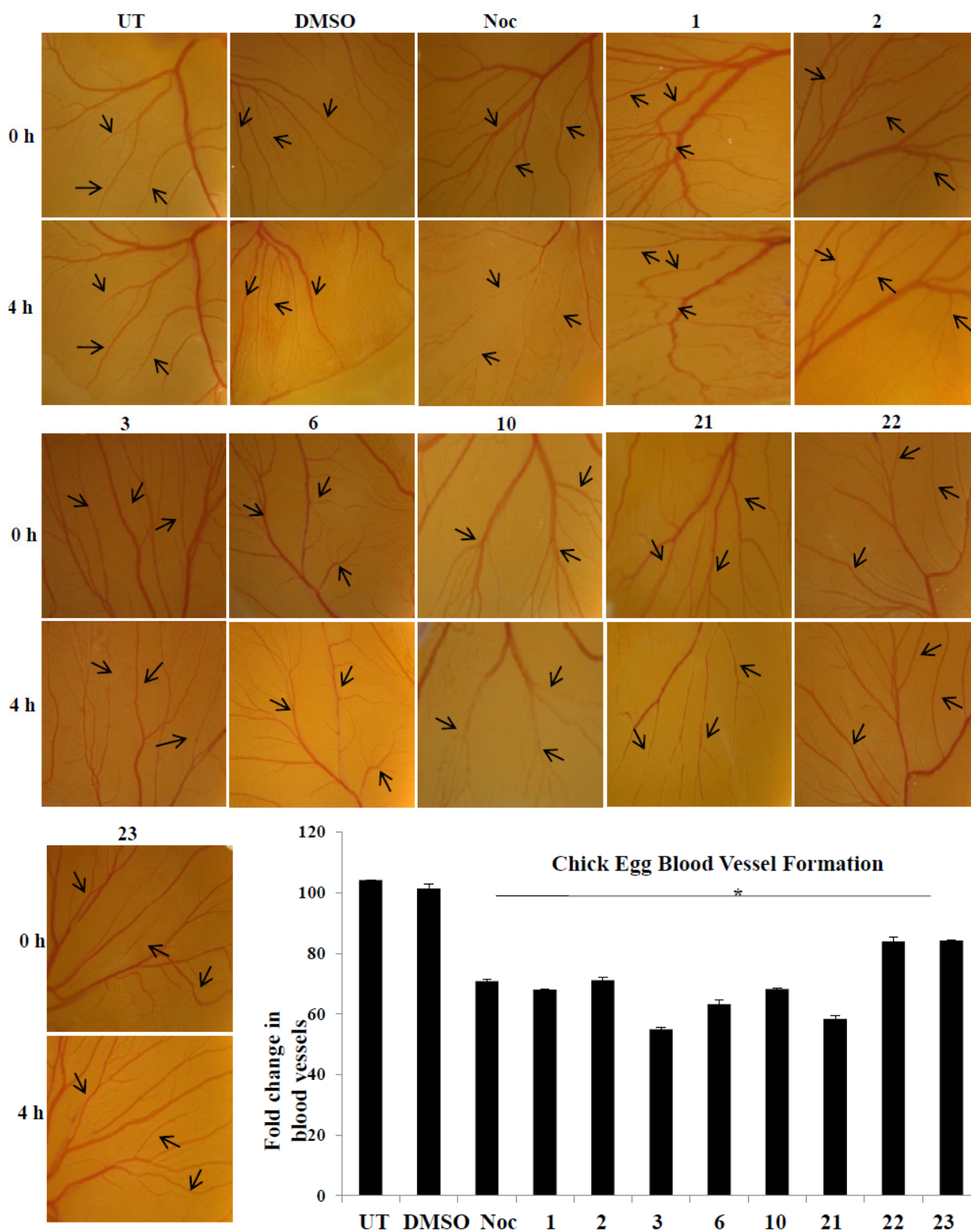


**Figure 7:** Benzothiazolo-quinazoline congeners occupy the colchicine binding site of tubulin. A, Structural similarities of compound 1 with other prominent colchicine binding agents. B, compound 1 in colchicine binding pocket. C, series-1 overlap with colchicine. D, series-2 overlap with colchicine. E, compound 22 and 23 overlap with colchicine. Here, yellow sticks represent Colchicine, orange sticks represent ligands, the  $\beta$ -tubulin residues are represented by grey thin sticks, and  $\alpha$ -tubulin residues are pink thin sticks.

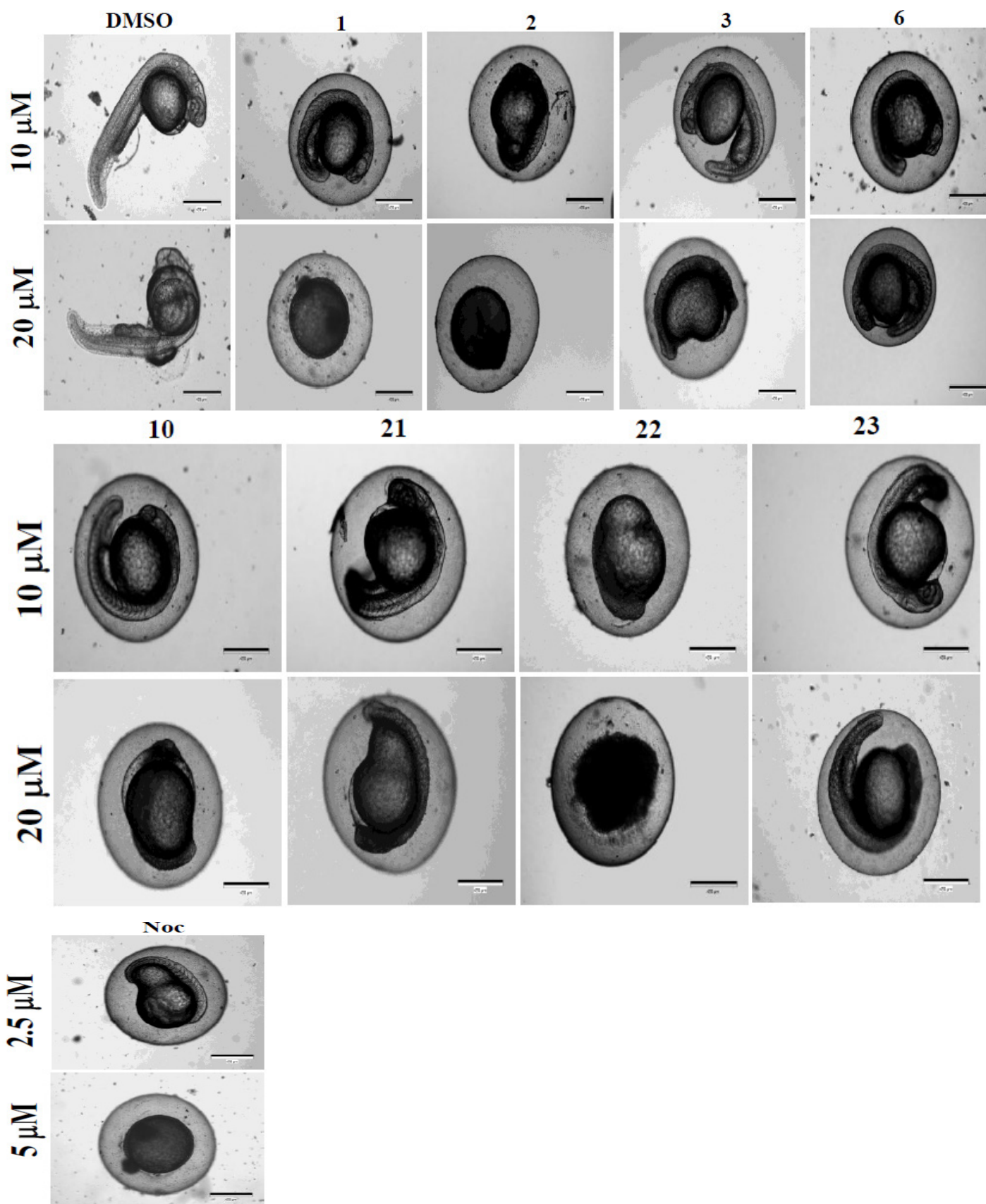
### Benzothiazolo-quinazoline congeners act as microtubule depolymerising agents

As the selected congener's phenocopy nocodazole, a microtubule depolymerising agent by arresting cells in G2/M and increasing mitotic markers, so we checked whether the benzothiazolo-quinazoline congeners show similar mechanism of action. To check, we treated HeLa cells with 1-3, 6, 10, and 21-23

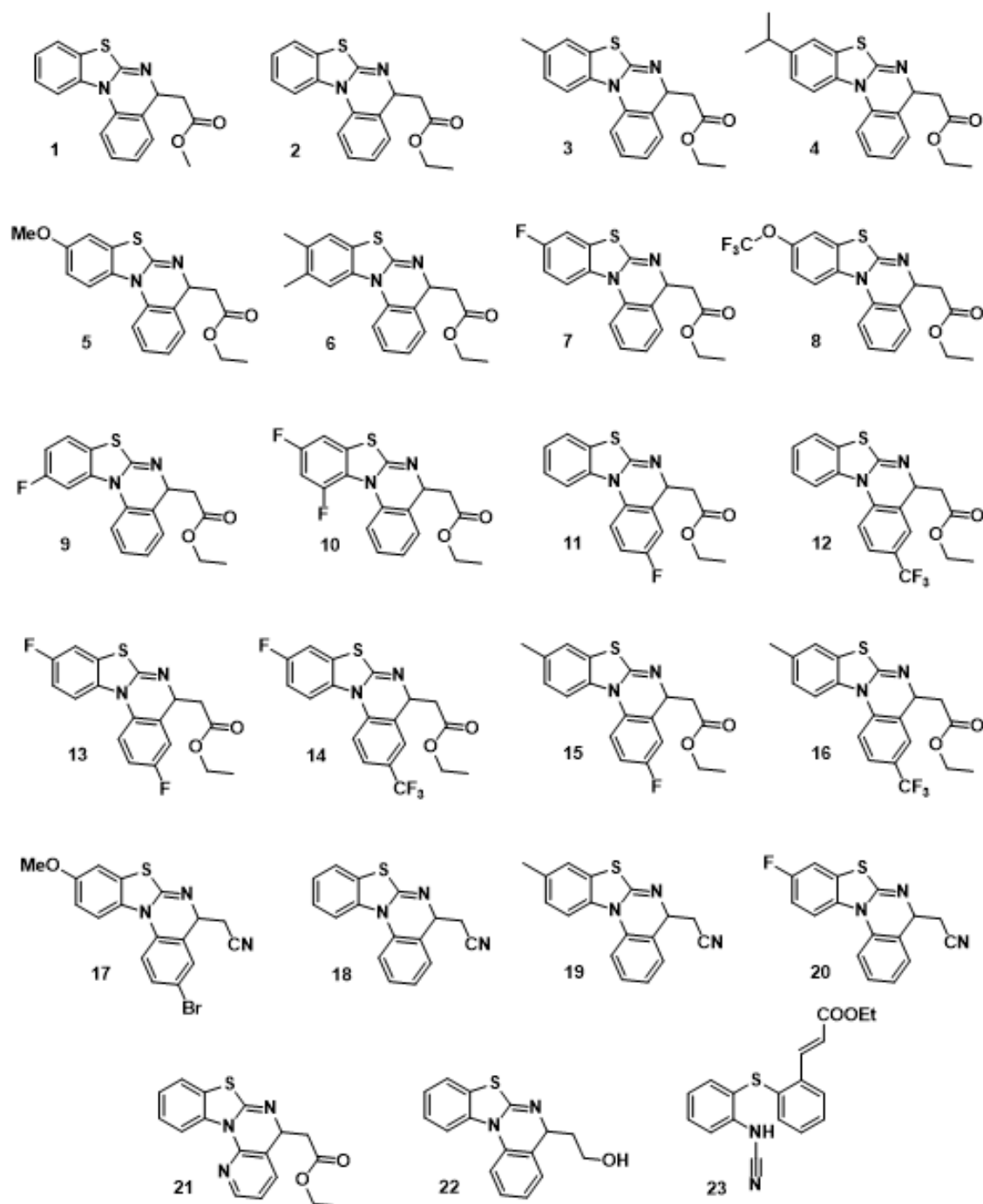
for 18 hr and then collected soluble and insoluble fractions for immunoblot analysis. As controls, we also treated HeLa cells with DMSO, nocodazole, or paclitaxel, a microtubule polymerising agent for similar duration. We observed that the selected benzothiazolo-quinazoline congeners have more soluble tubulin levels similar to nocodazole. In comparison, DMSO-treated cells had fairly equal levels of tubulin indicating homeostasis, whereas Paclitaxel-treated cells had more insoluble tubulin levels (Figure



**Figure 8:** Benzothiazolo-quinazoline congeners block angiogenesis in chick embryos- Chick embryos were treated with 5  $\mu$ M of 1-3, 6, 10, and with 10  $\mu$ M 21-23 at 0 hr and 4 hr. Nocodazole (Noc) was used as a positive control, at 2.5  $\mu$ M concentrations. Images were quantified using Image J software, and plotted as fold change in blood vessel formation and represented as mean $\pm$ SD. Significant differences from UT embryo were observed (\* $p$ <0.05),  $n$ =3.



**Figure 9:** Benzothiazolo-quinazoline congeners inhibit zebrafish embryo growth-Zebrafish embryos were treated with 10 μM and 20 μM of 1-3, 6, 10, 21-23 at 5 hr post fertilization. Nocodazole (Noc) was used as a positive control, at 2.5 μM and 5 μM concentrations After 28 hr, the embryos were imaged using microscopy to observe morphological changes.



**Chart 1:** Benzothiazolo-quinazoline congeners, 1-23 were employed in the study.

4). Taken together, the benzothiazolo-quinazoline congeners act microtubule depolymerising agents by increasing soluble tubulin levels.

### Benzothiazolo-quinazoline congeners disrupt microtubule network

Microtubule network is disrupted during cell division, where in microtubules bind to and aid in the separation of

chromosomes. As benzothiazolo-quinazoline congeners arrest cells in G2/M and increase mitotic markers, we checked if the benzothiazolo-quinazoline affects cellular phenotype. To check, we treated HeLa cells with the selected benzothiazolo-quinazoline congeners for 18 hr and stained for tubulin protein. Immunofluorescence analysis revealed that HeLa cells treated with selected benzothiazolo-quinazoline congeners have spindle microtubules with centrally located chromosome (Figure

5). Taken together, the selected benzothiazolo-quinazoline derivatives trigger a change in HeLa cells from a normal phenotype to mitosis.

### Benzothiazolo-quinazoline congeners depolarise mitochondria, induce ROS and promote mitotic cell death

Anticancer agents deplete cellular glutathione levels and increase ROS levels to trigger ROS-dependent cell death.<sup>33</sup> As benzothiazolo-quinazoline congeners 1-3, 6, 10, and 21-23 reduce cell growth and arrest cells in G2/M; thus, we determined if benzothiazolo-quinazoline congeners depolarise mitochondria and activate cell death. For mitochondria, we performed JC-1 staining, and found that benzothiazolo-quinazoline congener treated cells harboured depolarised mitochondria, as evidenced by strong JC-1 aggregate staining (Figure 6A). For cell death, we treated HeLa cells with compounds and examined for Annexin-V staining. We found that benzothiazolo-quinazoline congeners activate mitotic cell death as evidenced by strong Annexin-V staining (Figure 6B). Next, we checked for another apoptotic parameter, cellular Reactive Oxygen Species (ROS) levels; we treated HeLa cells with selected congeners and performed cytometry analysis. We found that selected congeners increase cellular ROS levels, suggesting that the congeners trigger ROS-dependent cell death (Figure 6C). Taken together, the benzothiazolo-quinazoline congeners depolarise mitochondria, induce ROS and promote mitotic cell death in HeLa cells.

### Benzothiazolo-quinazoline congeners occupy the colchicines binding site of tubulin

Colchicine is a natural product binding at the interface of  $\alpha$  and  $\beta$  subunits in the Colchicine Binding Pocket (CBP) that binds tubulin and inhibits tubulin polymerisation. The CBP is constructed by the strands S8 and S9, loop T7, and helices H7 and H8; the pocket is divided into three zones (zone-1 to 3). The peripheral residues of  $\alpha$  and  $\beta$  tubulins form zone-1, zone-2, and zone-3 are situated in  $\beta$ -tubulin, where zone-2 forms the main pocket and zone-3 extends further into the  $\beta$ -tubulin forming an extended pocket.<sup>34-36</sup> The colchicine binds in the main cavity (zone-2) and extends into zone-1. The ring-A methoxy groups interact with  $\beta$ C241 and the ring-B associate with  $\alpha$ T179, and  $\alpha$ V181 residues.<sup>37,38</sup> The docking of these compounds obtained good binding modes ( $\sim$ -7.0 kcal/mol) (Table 2) and interaction analysis revealed three different binding patterns. The set-1 compounds (1, 2, and 21) occupied zone-2 of CBP similar to colchicine in set-1 compounds, three rings (quinazoline and thiazole) were homologous to three rings of colchicine and the remaining fourth ring occupied the cavity designated for methoxy groups. The most active compound 1 exhibited a dock score of -7.34 kcal/mol and has a high Glide Emodel value of -71.69 kcal/

mol, along with increase in binding free energy ( $\Delta G_{\text{bind}} = -105.00$  kcal/mol), showing good affinity and excellent occupancy (Figure 7B). We found that the ester tail extended into zone-1 similar to the amide residue of colchicine (Figure 7C). Further, we observed that 2 has a good dock score (-7.22 kcal/mol), binding free energy (-76.51 kcal/mol), and Emodel score (-66.10 kcal/mol), and the ester carbonyl of 2 formed an H-bond with  $\beta$ A250 (backbone), possibly contributing to better cytotoxicity. Finally, 21 also displayed good affinity (-7.32 kcal/mol), though relatively less cytotoxic. The set-2 (3, 6, and 10) on the other hand, displayed an inverted mode of binding where the aromatic rings occupied the sidechain region of colchicine while extending the ester tail into zone-2 (Figure 7D). Furthermore, difluoro substituted aromatic ring (ring-B) of 10 also formed  $\pi$ -cation interaction with  $\beta$ K254 (side chain), which might have contributed to its high binding efficiency (-7.38 kcal/mol). Nevertheless, its poor binding free energy (-49.52 kcal/mol) indicates forced binding. Interestingly, set-1 seems to have more affinity towards tubulin binding, which is clearly depicted by excellent binding efficiencies (Table 2). Compound 22 occupied zone-2 and the alcohol sidechain slightly extended into zone-3, towards the methoxy region of colchicine with a moderate binding (-6.74 kcal/mol); similarly, 23 also extends into zone-3 (Figure 7E). The polar nature of these sidechains might cause the decrease in cytotoxicity as the polar 22 is relatively less active, coinciding with the decrease in dock score. The (*S*)-enantiomer of the ligands showed preferable affinity towards tubulin compared to their counterparts. The ADME study identified the ligands as drug like that obeyed Lipinski rule of five. Additionally, the toxicity prediction from ProTox-II established their non-toxic nature. The Table 2 summarised all the *in silico* parameters and docking parameters of the compounds, respectively.

### Benzothiazolo-quinazoline congeners block angiogenesis in chick embryos

Anti-microtubule agents such as combretastatins, BNC105, colchicines, and phenylahistin were shown to occupy the colchicine site in the microtubule and disrupt blood vessel formation in tumors. These set of compounds were classified as vascular disrupting agents.<sup>39,40</sup> As benzothiazolo-quinazoline congeners 1-3, 6, 10, and 21-23 also disrupt tubulin polymerization and occupy colchicine site, therefore, we asked whether benzothiazolo-quinazoline congeners can function as vascular disrupting agents. To answer this question, we performed the chick embryo angiogenesis assays and examined the formation of the blood vasculature. We found that the selected benzothiazolo-quinazoline congeners effectively disrupt blood vasculature, but not in DMSO-treated eggs (Figure 8). Finally, these results show that benzothiazolo-quinazoline congeners can be employed as vascular disrupting agents.

## Benzothiazolo-quinazoline congeners inhibit zebrafish embryo growth

Zebrafish (*Danio rerio*) is an emerging model system to elucidate chemical genetics.<sup>41</sup> To check whether benzothiazolo-quinazoline congeners can affect zebrafish embryos development, we treated embryos with benzothiazolo-quinazoline congeners 1-3, 6, 10, and 21-23 and examined the treated embryos. We found that benzothiazolo-quinazoline congeners delayed embryo development when compared to DMSO-treated or control embryos. The control embryos traversed to the prim6 stage, whereas the benzothiazolo-quinazoline congener's embryos were arrested at the 18-cell somite stage. As a control, we found that nocodazole-treated embryos also show similar abnormalities in development (Figure 9). Based on these observations, we show that benzothiazolo-quinazoline congeners can function as potential chemotherapeutic leads.

## CONCLUSION

In the present study, we show that that eight benzothiazolo-quinazoline congeners act as anti-microtubule agents. The congeners are amenable for further structural modifications and can be lead-optimised into next generation anti-microtubule agents.

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## CONFLICT OF INTEREST

The authors declare that there is no conflict of interest.

## ABBREVIATIONS

**DMEM:** Dulbecco's modified Eagle's medium; **ROS:** Reactive oxygen species; **SDS-PAGE:** Sodium dodecyl sulfate-polyacrylamide gel electrophoresis; **PBS:** Phosphate buffer saline; **ECL:** Enhanced chemiluminescence; **Noc:** Nocodazole; **ATCC:** American Type Culture Collection.

## AUTHORS' CONTRIBUTION

Supriya Bhukya: Conceptualization, Methodology, Validation, Data curation, Writing-original draft, Writing-review and editing, Sonam Swain, Darna Mounika Methodology, Validation, Data curation. Jyoti Honnanayakanavar, Battu Harish, Purna Chandra Behera, Validation, Data curation. Vidya Jyothi Alli, Methodology, Validation Supervision, Surender Singh Jadav, Writing-review and editing, Investigation, Methodology, Validation, Data Anthony Addlagatta, Suriseti Suresh, Nishant Jain, Writing-original draft, Writing-review and editing, Conceptualization, Investigation, Project administration, Funding acquisition.

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