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## Research Paper

# In-Silico Design, Synthesis and Antiproliferative Evaluation of Thiazolidino-Chromone Derivatives

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## ABSTRACT

Chromones are a group of naturally occurring chemical compound that being existing everywhere in nature, predominantly in plants. The word chromone is derived from the Greek word Chroma, which means colour, which bringing up that most of chromone derivatives can exhibit distinctiveness, in colour. Chromones are the oxygen containing heterocyclic compounds with a benzoannulated  $\gamma$ -Pyrone ring (4H-Chromone-4-one, 4H-benzopyran-4-one). Now a days chromone act as a valid scaffold in medicinal chemistry. Our aim was to develop novel thiazolidino- chromone derivatives which inhibiting Human Topo (II) $\alpha$  ATPase and provide anti-proliferative activity. In-silico design of novel analogues were carried out using ACD labs ChemSketch 12.0. Molinspiration software was used to analyse 'Lipinski Rule of Five' and drug likeness properties. Biological activity was predicted by PASS software. Preliminary docking study was carried out using GLIDE software by SCHRODINGER. Five derivatives which obeyed rule of five and having predicted antitumor activity on Human Topo (II) $\alpha$  ATPase were synthesized by four step process. After the completion of reaction in each step, the compounds were isolated, recrystallised by using suitable solvents, purified by TLC and column chromatography. Analogues were characterized by FT-IR, <sup>1</sup>H NMR, <sup>13</sup>C NMR and Mass Spectroscopy. The Biological evaluation was done by MTT assay using HCT 116 cancer cell lines. The results were compared with standard anticancer drug 5-Fluorouracil.

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The results were compared with standard anticancer drug 5-Fluorouracil. The results of present research work showed that novel thiazolidino- chromone derivatives have comparable antiproliferative effect with that of standard anticancer drug 5-Fluorouracil. This will lead to the development of promising lead compounds for target specific anticancer therapy and encourage further optimization to develop potent antiproliferative agents.

## INTRODUCTION

Cancer is emerging as a first major health problem in developing as well as developed countries. Surpassing cardiac diseases, it is taking number one killer worldwide due to various social, economic and lifestyle factors. There are many chemotherapeutic strategies for cancer treatment have been proposed, tested and in some cases implemented in the past two decades, these diseases still remain deadly. Therefore, there is a desperate need to develop treatments with new chemical entities with novel mechanism of action to combat this disease. One of the most important receptor which is overexpressed in majority of solid tumors is Human Topo (II)  $\alpha$  ATPase. The potential inhibitor of this receptor will definitely prevent this enzyme cascade mechanism and blocks cell division. The chromones are the interesting as structural scaffolds and have been assigned as privileged structures for drug development as probes to study lipid membranes and proteins [8-10]. The versatile biological applicability of chromone derivatives and their potential use in drug discovery implicates the importance of access to efficient synthetic routes to well-designed substituted chromones. 4-Thiazolidinones and their derivatives are an important class of compounds in organic and medicinal chemistry. The 4 thiazolidinone ring system is a core structure in various synthetic pharmaceutical agents, displaying a broad spectrum of biological activities such as, antitubercular, anti-bacterial, anti-HIV, anti-inflammatory, anti-mycobacterial, anti-convulsant, anti-histaminic, anti-cancer, anti-protocol and analgesic. 4- Thiazolidinones are

derivatives of thiazolidine with carbonyl group at the 4th position and formed by the attack of sulphur nucleophile on imine carbon followed by intramolecular cyclization with elimination of water. The rationale behind this research work is that on combining these two therapeutically significant heterocyclic moiety results into a new chemical nucleus which could specifically inhibit Human Topo (II) $\alpha$  receptor, which ultimately leads to the prevention of growth and proliferation of cancerous tissue. So, we aimed to *in-silico* design and development of Thiazolidino chromone derivatives as potential anticancer agents.

## MATERIALS AND METHODS

### Materials and instrumentation

All the chemicals and reagents used in this research work were of analytical or synthetic grade from Sigma Aldrich, E-Merck (Germany) and S D Fine Chemicals (India). All the chemicals were dried and purified according to standard methods before use, wherever necessary. Software used in this study include ACD Labs Chemsketch, Chemdraw Ultra 8.0, Molinspiration, PASS online, The synthetic procedures were carried out by using both conventional and microwave method. All the reaction courses and product mixtures were routinely monitored by TLC plates and visualized with UV light or iodine chamber. Melting point of synthetic compounds was determined on a Labindia MR-VIS visual melting point apparatus and is uncorrected. Absorbance values against wavelength were taken on a Systronic double beam UV-166 spectrophotometer. The FT-IR spectra were recorded using FT-IR (Agilent Cary 630 FT-IR spectrophotometer using KBr pellet). <sup>1</sup>H NMR spectra were recorded using NMR spectrophotometer (Bruker 400 ultra shield DPX 400) and chemical shifts are expressed as  $\delta$  (ppm) using TMS as an internal standard in DMSO-*d*<sub>6</sub>. Mass spectra of the compounds were done with



mass spectrometer (micromass-O-TOF-MS ES+). Anticancer evaluation was done by MTT assay using the HCT 116 cell lines.

### ***In-silico* methods**

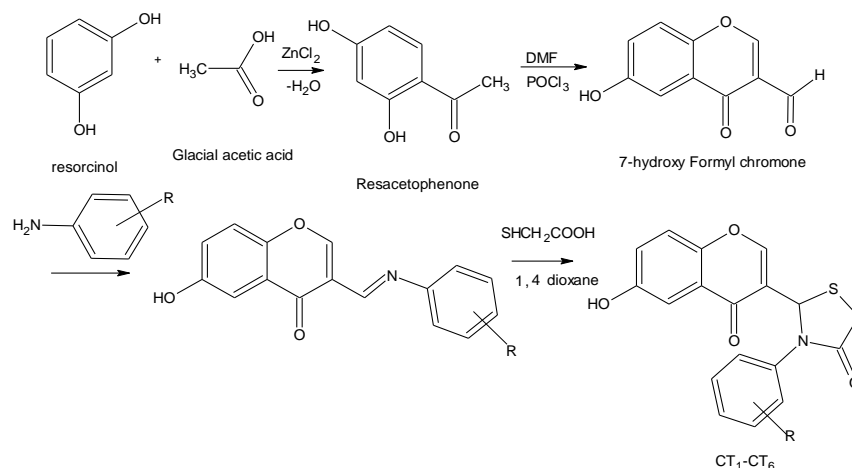
#### ➤ ***In-silico* molecular modeling**

*In silico* methods in drug discovery can be used for the identification and quantification of the physico-chemical properties of the new drug candidate and also to analyse whether any of these properties of the new drug candidate and also to analyse whether any of these properties having significant effect on its biological activity. *In silico* designing of drug candidate will help to identify the possible target of the drug candidate and predicts its biological activity. The physico-chemical properties of the drug candidate were discovered by using various computer software, in which the electronic, lipophilic and steric parameters can be determined by using ACD Lab chemsketch. Molinspiration software will help to determine the drug likeness by analyzing lipinsky rule of five. The approach used in PASS is based on the suggestion that the affinity = f (structure).

#### ➤ **Docking studies**

Docking is the computational simulation of a ligand binding to a receptor, which helps to predict the binding orientation of small molecule drug candidates to their protein targets in order to predict the affinity and the activity of the small molecule. Docking is a very important tool in the rational design of drugs. Schrodinger is a comprehensive software suite for analyzing and modeling molecular structures, biological macromolecules (proteins and nucleic acid). The selected analogues were docked onto the binding pocket Human Topo (II) $\alpha$  ATPase Receptor (PDB ID. 1ZXM). These docking studies give the best matching between two molecules: designed thiazolidino chromone and the binding pocket of target protein. Different steps involved in docking studies include; preparation of ligand and protein, docking methods, scoring of docking results and analysis, refinement and filtering tools.

#### **Synthetic methodology**



**Scheme 1: Synthesis of Thiazolidino chromone (CT1-CT6)**

#### **Step 1 Synthesis of resacetophenone (2,4-dihydroxyacetophenone)**

2,4-dihydroxyacetophenone was synthesized by reported method. Briefly, the synthesis was carried out by dissolving freshly fused and powdered zinc chloride (0.24 mole) in 32 ml of glacial acetic acid

by heating in sand bath. Dry resorcinol (0.2 mole) was added with stirring at 140<sup>o</sup>C. The solution was heated until it just begins to boil and kept for 20 minutes at 150<sup>o</sup>C. Dilute HCl (1:1) was added to the mixture and the solution was cooled to 5<sup>o</sup>C. The separated product was filtered and washed

with dilute HCl. The product was recrystallised from hot water.

### Step 2 Synthesis of formyl chromone

POCl<sub>3</sub> (0.49 mol) was added drop wise to dimethylformamide (DMF) (121 ml) with stirring at 30-35° C, after the addition, the mixture was stirred at 50°C for 1 h. Then the solution of acetophenone derivatives (0.12 mol) in least amount of DMF was added drop wise with stirring to the above mixture. After that the mixture was stirred at 45-55°C for 2 hours, kept over the night at room temperature and slowly poured over mixture ice and water(200g). Product was stirred for 6 hours, then filtered off and recrystallized from ethanol.

### Step 3 Synthesis of Schiff base

A 50 ml borosilicate glass beaker was charged with formyl chromone (188 mg, 0.001mol) and aromatic amine (93 mg, 0.001 mol) in methanol (5 mL). The reaction mixture was irradiated inside a microwave oven for 10 min at an output of 340 watts power. Progress of the reaction was monitored by TLC. After completion of the reaction, the reaction mixture was poured into the ice water and the product obtained was separated by filtration. The product was purified by recrystallization from ethanol.

### Step 4 Synthesis of thiazolidinone substituted chromone derivatives

To the mixture of schiff base (0.01 mol) and mercapto acetic acid (0.01) mol dissolved in dioxane (20 ml), anhydrous zinc chloride (0.0004 mol) was added and refluxed, for 8 hrs. The reaction mixture were poured into ice water, filtered, washed with 10% sodium carbonate solution and recrystalized using methanol.

### Antiproliferative evaluation (MTT Assay)

Determination of cell growth rates is widely used in the testing of drug action, cytotoxic agents and screening other biologically active compounds. MTT assay is a colorimetric assay that measures the reduction of yellow 3-(4,5-dimethylthiazol-2-

yl)-2,5-diphenyltetrazolium bromide (MTT) by mitochondrial succinate dehydrogenase. The MTT enters the cells and passes into the mitochondria where it is reduced to an insoluble, colored (dark purple) formazan product. The cells are then solubilized with an organic solvent (e.g. Dimethylsulfoxide) and the released, solubilized formazan reagent is measured spectrophotometrically at 540 nm. Since reduction of MTT can only occur in metabolically active cells the level of activity is a measure of the viability of the cells. The cells were washed with 1x PBS and then added 30 µl of MTT solution to the culture (MTT- 5mg/mL dissolved in PBS). It was then incubated at 37°C for 3h. MTT was removed by washing with 1x PBS and 200µl of DMSO was added to the culture. Incubation was done at room temperature for 30 minutes until the cell got lysed and color was obtained. The solution was transferred to centrifuge tubes and centrifuged at top speed for 2minutes to precipitate cell debris. Optical density was read at 540 nm using DMSO as blank in a ELISA microplate reader.

**% viability = (OD of Test/ OD of Control) X 100**

**Percentage mortality= 100 - %viability**

## RESULTS AND DISCUSSION

### *In-silico* molecular modeling studies

The *In-silico* molecular modeling studies of novel thiazolidino chromone derivatives were carried out successfully with the aid of different software for selection of suitable drug candidates prior to wet lab synthesis. *In-silico* studies were performed on designed 12 derivatives by means of ACD Lab ChemSketch 12.0, Chem Draw 8.0, Molinspiration, PASS, Schrodinger software. Among the 12 designed analogues, six analogues were found to obey Lipinski rule of five and their drug likeness were predicted by Molinspiration software. The analogues which are having desired physico-chemical properties were chosen for wet lab synthesis (**Table 1, Table 2, Table 3, Table 4**).



**Table 1: Molecular descriptors for designed analogues generated by ACD Labs Chems sketch 12.0**

| Compound | Mol. wt | Molar volume cm <sup>3</sup> | Parachor Cm <sup>3</sup> | Surface tension Dyne/cm | Polarisability (10 <sup>-24</sup> cm <sup>3</sup> ) |
|----------|---------|------------------------------|--------------------------|-------------------------|---|
| CT1      | 339.336 | 227±3.0                      | 663.5±6.0                | 72.8±30                 | 35.60±0.5   |
| CT2      | 384.364 | 238.9±3.0                    | 720.6±6.0                | 82.7±30                 | 38.20±0.5   |
| CT3      | 384.364 | 238.9±3.0                    | 720.6±6.0                | 82.7±30                 | 38.20±0.5   |
| CT4      | 373.811 | 239±3.0                      | 700.7±6.0                | 73.8±30                 | 37.55±0.5   |
| CT5      | 355.336 | 225.5±3.0                    | 678.7±6.0                | 82.0±30                 | 36.35±0.5   |

**Table 2: Analysis of Lipinski rule of five for selected thiazolidino-chromone derivatives analogues**

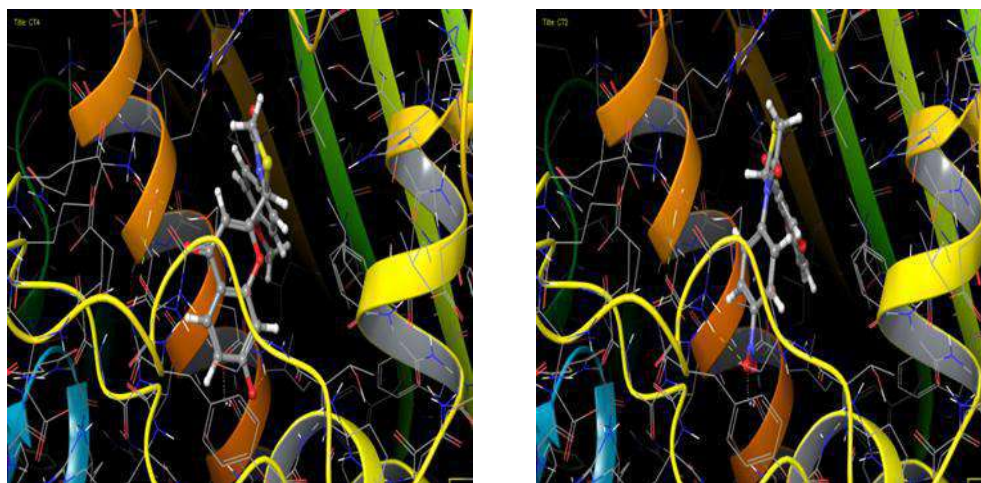
| Compound | Log P | Molecular Weight | No. of Hydrogen bond acceptors | No. of Hydrogen bond donors | No. of Rotatable Bonds | Violations |
|----------|-------|------------------|--------------------------------|-----------------------------|------------------------|------------|
| CT1      | 2.62  | 339.37           | 5                              | 1                           | 2                      | 0          |
| CT2      | 2.53  | 384.37           | 8                              | 1                           | 3                      | 0          |
| CT3      | 2.53  | 384.37           | 8                              | 1                           | 3                      | 0          |
| CT4      | 2.68  | 369.40           | 6                              | 1                           | 3                      | 0          |
| CT5      | 2.14  | 335.37           | 6                              | 2                           | 2                      | 0          |
| CT6      | 3.79  | 389.43           | 5                              | 1                           | 2                      | 0          |

**Table 3: Analysis of drug likeness score for selected derivatives**

| compound | GPCR ligand | Ion channel modulator | Kinase inhibitor | Nuclear receptor Ligand |
|----------|-------------|-----------------------|------------------|-------------------------|
| CT1      | -0.39       | -0.65                 | -0.85            | -0.57                   |
| CT 2     | -0.48       | -0.62                 | -0.90            | -0.59                   |
| CT3      | -0.49       | -0.66                 | -0.93            | -0.77                   |
| CT4      | -0.40       | -0.82                 | -0.79            | -0.70                   |
| CT5      | -0.37       | -0.63                 | -0.81            | -0.54                   |
| CT6      | -0.26       | -0.63                 | -0.67            | -0.60                   |

**Table 4: Prediction of Biological activity of proposed analogues using PASS software**

| Compounds | Effect     | Pa    | Pi    |
|-----------|------------|-------|-------|
| CT1       | Anticancer | 0.291 | 0.023 |
| CT2       | Anticancer | 0.271 | 0.031 |
| CT3       | Anticancer | 0.294 | 0.022 |
| CT 4      | Anticancer | 0.299 | 0.020 |
| CT5       | Anticancer | 0.283 | 0.026 |
| CT6       | Anticancer | 0.284 | 0.025 |



**Figure 1: Docking images of Thiazolidino chromone derivatives (A. CT2 and B CT4)) on binding pocket of Human TOPO (ii) A**

**Table 5: Docking scores of selected derivatives with target protein**

| Target           | PDB ID | Compound Name | GLIDE Score |
|------------------|--------|---------------|-------------|
| Human TOPO(II) A | 1ZXM   | CT1           | -5.7        |
|                  |        | CT 2          | -6.43       |
|                  |        | CT3           | -4.1        |
|                  |        | CT4           | -10.17      |
|                  |        | CT5           | -5.92       |
|                  |        | CT6           | -5.94       |

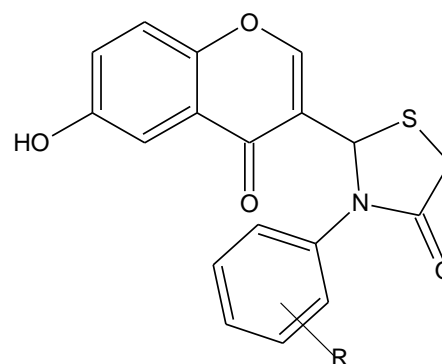
### Molecular docking

All the proposed derivatives were subjected to flexible docking on to the binding pocket of Human TOPO(II)  $\alpha$  (Pdb ID: 1ZXM) using GLIDE Programme of Schrodinger. The docking scores were calculated on the basis of Glide score (**Figure 1, Table 5**).

### Synthetic methods

The analogues which were designed by *in-silico* studies were selected for wet lab synthesis based on Lipinski rule of five, PASS value and docking energy score. The synthetic scheme involved was a five-step reaction. After the isolation of product in each step the products were recrystallised and purified by TLC. The structure of proposed analogue is shown in **Figure 2**. Five new

derivatives were synthesized by conventional both conventional and microwave method). The percentage yield of the reaction, melting point, and Rf value of each compounds were calculated and shown in **Table 6**.



**Figure 2: General structure of thiazolidino chromone derivatives.**

**Table 6: Characterization data of synthesized acetidino-quinazoline derivatives**

| Compound code | Molecular formula   | Molecular Weight (g) | Melting point | % yield | Rf value |
|---------------|---|----------------------|---------------|---------|----------|
| CT 2          | C <sub>18</sub> H <sub>12</sub> N <sub>2</sub> O <sub>6</sub> S | 384.364              | 240-242       | 70      | 6.9      |
| CT 3          | C <sub>18</sub> H <sub>12</sub> N <sub>2</sub> O <sub>6</sub> S | 384.364              | 226-228       | 72      | 6.7      |

|      |   |         |         |    |     |
|------|---|---------|---------|----|-----|
| CT 4 | C <sub>18</sub> H <sub>12</sub> N <sub>2</sub> O <sub>6</sub> S | 373.811 | 221-223 | 68 | 6.5 |
| CT 5 | C <sub>18</sub> H <sub>13</sub> NO <sub>5</sub> S               | 355.336 | 214-217 | 55 | 6.1 |
| CT 6 | C <sub>22</sub> H <sub>17</sub> NO <sub>4</sub> S               | 391.441 | 291-293 | 77 | 7.1 |

**Table 7: Characteristic FT-IR, 1HNMR and Mass spectral analysis of synthesized analogues**

| Compound                     | IR (KBr $\nu$ cm <sup>-1</sup> )  |
|------------------------------|---|
| STEP 1                       | 3648.098 OH stretch, 3047.792 aromatic CH, 3002.975 Aliphatic CH, 1516.987 Aromatic C=C, 1202.628 OH-bend.  |
| STEP 2                       | 3592 OH stretch, 3040.44 aromatic CH, 1274.282 OH-bend, 3592 OH stretch, 2918.855 aliphatic CH, 1740.571 C=O stretch (aldehyde), 1692.347 C=O stretch, 1441.747 aromatic C=C, 1274 and 1142 C-O-C |
| STEP 3                       | 3600.217 OH stretching, 2999.543 aromatic CH, 1691.809 C=O, 1569.291 and 1349 NO stretch, 1428.341 aromatic C=C, 1102.945 C-O, 800.00 C-N.  |
| STEP 4(CT2)                  | 3600.337 OH stretching, 3111.241 aromatic CH, 1645.809 C=O, 1560 and 1349.907 NO stretch, 1428.341 aromatic C=C, 1102.945 C-O, 782.00 C-N, 650.00 C-S-C,  |
| 1HNMR (CT2)                  | 3.3112(S, CH <sub>2</sub> ), 5.0210(S' Aromatic C-OH), 5.3160(S, CH), 6.6208-7.9631(M, Aromatic proton)   |
| Mass spectral Analysis (CT2) | 384.3 (Molecular ion peak), 161.02 (Base peak)  |

**Table 8: Comparative evaluation of Antiproliferative effect of CT2, CT4 and CT6 on HCT116 cells**

| Concentration | Percentage viability (%) |       |       |       |
|---------------|--------------------------|-------|-------|-------|
|               | standard                 | CT4   | CT6   | CT2   |
| 10 $\mu$ g/ml | 59.32                    | 66.92 | 79.88 | 80.44 |
| 20 $\mu$ g/ml | 55.86                    | 59.45 | 78.10 | 78.32 |
| 30 $\mu$ g/ml | 53.85                    | 54.27 | 72.51 | 69.60 |
| 40 $\mu$ g/ml | 44.58                    | 51.92 | 65.69 | 66.92 |

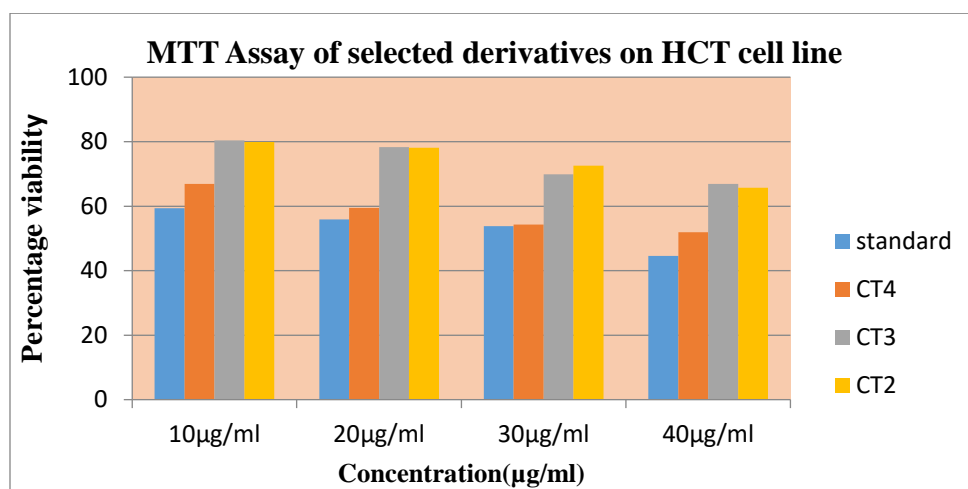
### Spectral characterization of Thiazolidino chromone derivatives

The newly synthesized novel Acetidinoquinazoline derivatives were further characterized by FT-IR, 1HNMR, and Mass spectral studies. The complete spectral analysis of prototype lead molecule CT2 is shown in Table 7.

### Evaluation of antiproliferative effect (MTT Assay)

After the preliminary in-silico molecular modeling studies followed by the docking studies on the binding pocket of Human TOPO (II)A receptor, thiazoliino chromone derivatives were selected for wet lab synthesis. The synthesized compounds

were purified and characterized by FT-IR, 1HNMR and Mass spectral studies. Docking studies proved that the CT2, CT4 and CT6 were more effectively bind with the receptor based on glide score. Four concentrations of the test compounds were used for MTT assay. The results were compared with that of standard drug 5-fluorouracil. The concentrations used were 10, 20, 30 and 40  $\mu$ g/mL. The cell lines used was HCT 116 cell line. Test results showed that the anticancer activity of the proposed derivatives is less than that of the standard drug 5-fluorouracil. The compound CT4 shows more activity than CT6 and Cl (Table 8, Figure 3)



**Figure 3: Comparative evaluation of antiproliferative effect of CT2, CT4 and CT6**

## CONCLUSION

The present work led to the development of novel antitumor molecules containing Thiazolidino chromone pharmacophore. This research work was focused on the structure-based drug design and development of novel Thiazolidino chromone derivatives and their antiproliferative evaluation. We have designed 12 new analogues and after *in-silico* molecular modeling and docking studies, selected six analogues for wet lab synthesis. These derivatives were spectrally characterized by FT-IR, <sup>1</sup>HNMR, mass spectroscopy. The antiproliferative evaluation of three derivatives was done against HCT116 cell line. The compounds CT2 and CT4 have shown significant activity against HCT 116 cell line and compared with that of standard drug 5-fluorouracil. Thus this work presents a potent antiproliferative effect of synthesized analogues. The Activity prediction by *in-silico* methods are very well correlated with biological activity. From this study, this can be concluded that the synthesized Thiazolidino chromone derivatives can be lead candidate to be developed into useful antiproliferative agents that could lead further research work on this potent nucleus. An extensive study is also warranted to determine additional physicochemical and biological parameters to have deeper insight into

SAR and optimize the effectiveness of these lead molecules.

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