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
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
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
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
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
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
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
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
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
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
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- ✓ [14]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC3708350/](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC3708350/)  
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
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- ✓ [32]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC9198363/](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC9198363/)  
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- ✓ [35]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC10220775/](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC10220775/)  
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- ✓ [45]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC10010179/](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC10010179/)  
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- ✓ [48]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC8972391/](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC8972391/)  
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- ✓ [49]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC5617838/](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC5617838/)  
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
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
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
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
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
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- ✓ [56]  [en.wikipedia.org/wiki/Oxidative\\_stress](http://en.wikipedia.org/wiki/Oxidative_stress)  
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- ✓ [57]  [www.frontiersin.org/articles/10.3389/fgene.2019.00858/full](https://www.frontiersin.org/articles/10.3389/fgene.2019.00858/full)  
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- ✓ [58]  from a PlagScan document dated 2022-09-20 08:26  
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
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
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
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
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- ✓ [62]  from a PlagScan document dated 2022-02-03 10:28  
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
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
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- ✓ [64]  from a PlagScan document dated 2023-04-17 08:20  
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
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
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- ✓ [66]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC6541445/](https://www.ncbi.nlm.nih.gov/pmc/articles/PMC6541445/)  
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
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- ✓ [67]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC3698477/](https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3698477/)  
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
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
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- ✓ [69]  from a PlagScan document dated 2023-04-24 06:00  
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- ✓ [70]  from a PlagScan document dated 2022-03-03 13:00  
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- ✓ [71]  from a PlagScan document dated 2018-08-30 09:13  
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- ✓ [72]  [en.wikipedia.org/wiki/Riyadh](https://en.wikipedia.org/wiki/Riyadh)  
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- ✓ [73]  [www.ncbi.nlm.nih.gov/pmc/articles/PMC7691860/](https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7691860/)  
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- ✓ [74]  from a PlagScan document dated 2023-04-25 07:53  
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- ✓ [75]  from a PlagScan document dated 2023-04-18 06:18  
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- ✓ [78]  from a PlagScan document dated 2022-06-28 08:49  
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- ✓ [79]  from a PlagScan document dated 2018-06-25 11:06  
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
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- ✓ [80]  [pubmed.ncbi.nlm.nih.gov/34817215/](https://pubmed.ncbi.nlm.nih.gov/34817215/)  
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
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- ✓ [81]  from a PlagScan document dated 2023-07-17 07:35  
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1 Synthesis of 1-(4-chlorobenzoyl)-4-(dimethylamino) pyridin-1-ium chloride and  
2 determination of cytotoxicity and induction of apoptotic protein in human colon  
3 cancer cells

4

5 Abstract

6 Background: In this experiment, we synthesized a new compound 1-(4-chlorobenzoyl)-4-  
7 (dimethylamino) pyridin-1-ium chloride (SM-9) and examined its toxicity and anticancer activity  
8 on human colon cancer (HCT-116) cells. Methods: We uncovered the underlying mechanism of  
9 cell toxicity and apoptosis in HCT-116 cells due to SM-9 compound exposure via MTT assay and  
10 protein profiling array and gene expression through the RT-PCR. Results: <sup>[19]</sup> Our data showed that  
11 the SM-9 compound activated caspase-3, caspase-8, p21, p27, p53 proteins involved in apoptosis  
12 in HCT-116, thereby inducing cytotoxicity, the formation of reactive oxygen species, and  
13 apoptosis. <sup>[19]</sup> The results of this study showed that the SM-9 compound has advantageous qualities  
14 and need to be taken as an anticancer medication. <sup>[19]</sup> This research concluded that through inducing  
15 apoptotic pathways in human colon cancer (HCT-116) cells, the SM-9 compound possesses anti-  
16 cancer capabilities.

17 Keywords: Cytotoxicity; Protein profiling array; Oxidative stress; HCT-116 cells; <sup>[5]</sup> Caspase-3/8,  
18 p21, p53 gene expression

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26 Introduction

27Over the past ten years, ionic liquids have drawn a lot of attention due to their special  
28qualities that may be useful for cutting-edge technologies and procedures. Ionic liquids  
29include, among others, corrosion inhibitors (El-Hajjaji et al., 2019), antimicrobials  
30(Albalawi et al., 2018, Titi et al., 2021), and antiviral and anticancer medicines (Titi et  
31al., 2020).<sup>[10]</sup> In this work, we created an ionic liquid (1-(4-chlorobenzoyl)-4-  
32(dimethylamino) pyridin-1-ium chloride, SM-9 compound, SM-9 compound) and tested  
33its cytotoxicity and antioxidant qualities on human colon cancer (HCT116) cells for a  
34duration of 24 hours.<sup>[10]</sup> The toxicity of SM-9 compounds is caused by a number of  
35processes, one of which is the increased generation of ROS in stressed live cells.  
36Mitochondria are the main source of ROS generation in tissue, and the most of ROS  
37produced in the chain of electron transport system (Hansford et al., 1997).<sup>[14]</sup> The electron  
38release from the chain of electron transport system directly to oxygen, and inducing tiny-  
39lived free radicals such as singlet oxygen and superoxide anion etc (Hansford et al.,  
401997).<sup>[7]</sup> In this experiment we have synthesized new chemical compound and investigate  
41the toxic effects of synthesized drug namely SM-9 compound on human colon cancer  
42cells.<sup>[6]</sup> Ali et al., (2023) has reported that biosynthesize cobalt nanoparticles with garlic  
43and onion peel inhibit the growth of bacteria.<sup>[7]</sup> In addition, our findings will be supportive  
44in determining the safety purpose of synthesized compound and it will be helpful as a raw  
45material for manufacturing of anticancer drug. 4-(Dimethylamino)pyridine is a cheap  
46and ingenious chemical induced skin toxicity.<sup>[6]</sup> ROS creates cellular oxidative stress and it  
47leads colon damages and colon cancer (Bardelčíková et al., 2023),<sup>[5]</sup> but at cellular level  
48colon's natural defense system gets damaged in response to chemical compound.<sup>[5]</sup> ROS,

<sup>[0]</sup> 49 which seriously deteriorates bio-membranes, biomolecules, and induces the expression of  
50 matrix metalloproteinases (Tu and Quan, 2016).

<sup>[6]</sup> 51 Drug toxicity includes various mechanisms, mainly the generation of extra reactive  
52 oxygen species (ROS). Schumacker et al., (2014) and Almutairi et al.,<sup>[7]</sup> (2021) have  
53 reported that the damage of the mitochondrial is due to overproduction of ROS in cells.

<sup>[9]</sup> 54 Toxicity induced by contaminant and drug compound in target cells is due to production  
55 of ROS, apoptotic and inflammatory process (El-Sayed et al., 2005). Oxidative stress has  
56 mainly resulted from the generation of ROS. Extra production of ROS induce imbalance  
57 of antioxidant mechanism in cells and as a consequence it leads to peroxidation of lipid  
58 molecules and other oxidative related stress enzymes (Zhuang et al., 2018).<sup>[5]</sup> This is the  
59 first study that reported the adverse effects of synthesized drug SM-9 compound on  
60 human colon cancer cells. <sup>[7]</sup> In this study we are determining the safely effects of SM-9  
61 compound on human colon cancer cells.

<sup>[52]</sup> 62. Materials and methods

<sup>[10]</sup> 63 2.1. Chemical and reagents

64 Trypsin, antibiotic/antimitotic solution (100×), stabilized, 2',7'- I  
65 chlorodihydrofluorescein diacetate (H2DCFDA), MTT dye, and sodium chloride were  
66 purchased from Sigma-Aldrich (St.Louis, Missouri, United States). Human Apoptosis  
67 Array C1(Code: AAH-APO-1-2) Ray Biotech, United States. <sup>[67]</sup> Fetal bovine serum (FBS),  
68 and Dulbecco's Modified Eagle's Medium/Nutrient Mixture F-12 Ham (DMEM), and  
69 from Gibco company USA. All other chemicals such as 4-chlorobenzoyl chloride (1.1  
70 eq), DMSO and ethyl alcohol etc.<sup>[22]</sup> were purchase de from local market Riyadh Saudi  
71 Arabia.

<sup>[0]</sup>72.2. Synthesis new chemical material

73.2.2.1. Synthesis of new chemical compound 1-(4-chlorobenzoyl)-4-(dimethylamino)

74pyridin-1-ium chloride (SM-9)

754-chlorobenzoyl chloride (<sup>[8]</sup>1.1 eq) was added to a solution of 4-dimethylaminopyridine (1.76 eq) in toluene, which was followed by stirring at 80 °C for 24 h. <sup>[13]</sup>The separation of 77viscous liquid from the initially obtained clear and homogeneous mixture of 4-Dimethylaminopyridine and 4-chlorobenzoyl chloride in toluene marked the completion 79of the reaction. <sup>[8]</sup>Extraction was used to separate the product from the unreacted starting 80materials and solvent. <sup>[13]</sup>Finally, all volatile organic compounds were removed from the 81residue by drying it under pressure (Fig.1 a, b).

82.2.2.2. Characterization of SM-9 compound

831-(4-chlorobenzoyl)-4-(dimethylamino) pyridin-1-ium chloride: MP 83–84 °C, <sup>1</sup>H NMR 84(DMSO, 400 MHz,)  $\delta$  8.17 (d, 2H), 7.91 (d, 2H), 7.54 (d, 2H), 6.94 (d, 2H), 3.14 (s, 6H). 85<sup>13</sup>C NMR (DMSO, 100 MHz,)  $\delta$  166.9, 157.4, 139.6, 138.3, 131.6, 130.1, 129.2, 107.4, 8640.); IR ( $\nu_{\max}$  cm<sup>-1</sup>) 3129 (C–H, sp<sup>2</sup>), 1559–1469 (C=C), 1163 (C–N), 1080 (C–O) 87(Fig.1 a, b).



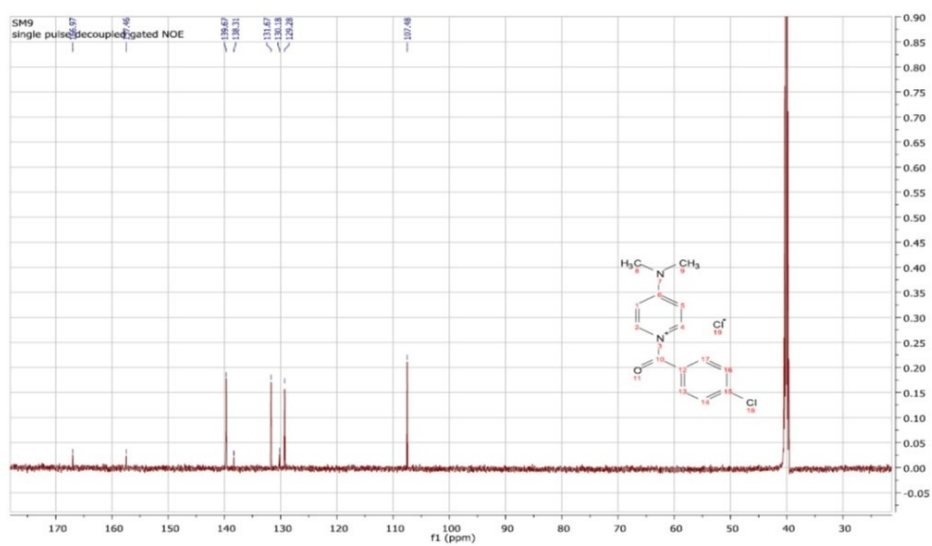


Fig.1 a  $^{13}\text{C}$  NMR spectrum of 1-(4-chlorobenzoyl)-4-(dimethylamino) pyridin-1-ium chloride (SM-9)

88

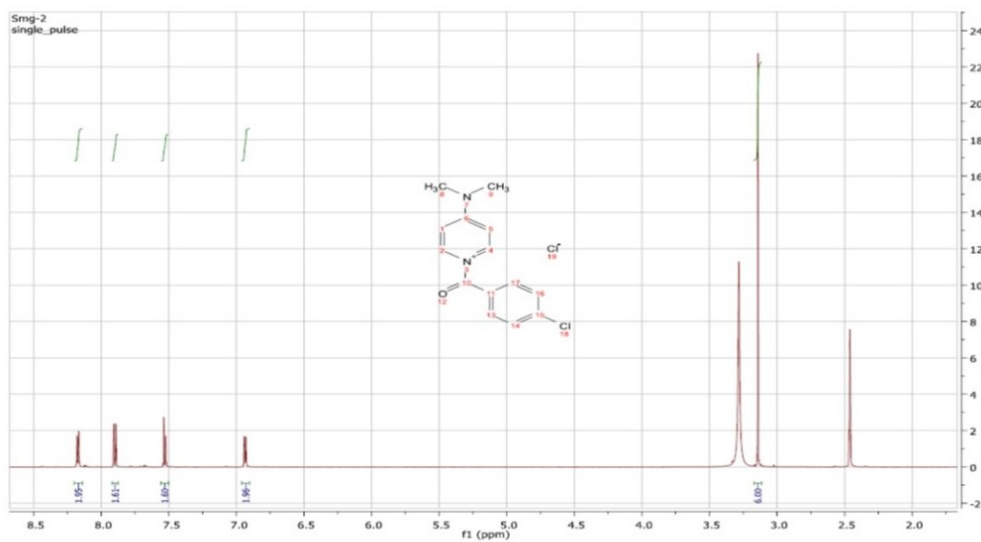


Fig.1 b  $^1\text{H}$  NMR spectrum of 1-(4-chlorobenzoyl)-4-(dimethylamino) pyridin-1-ium chloride (SM-9)

89

90

### 912.3.<sup>[80]</sup> Cell culture

92 The colon cancer (HCT 116) cell lines was bought from ATCC&(American Type  
93 Culture Collection) Manassas Virginia USA.<sup>[0]</sup> The cells were maintained in DMEM with

<sup>[0]</sup>94 fetal bovine serum (10%) and antibiotics & penicillin and streptomycin (10000U/ml) in  
95 CO<sub>2</sub> incubator (5%) at 37°C.

#### 96 2.4. Exposure of SM-9 compound

97 HCT-116 cells were sub-cultured for 24 h before exposure to SM-9 compound. Stock  
98 suspension of SM-9 compound (50 mg /ml DMSO) was prepared in slightly warm  
99 DMSO. The stock suspension of drug was diluted as per experimental concentration. <sup>[57]</sup>  
100 In this experiment we applied different concentrations of compound (control, solvent  
101 control, 50, 100, 200, 300, 400 and 500 µg/ml) to determine the toxicity of compound.  
102 <sup>[21]</sup> The untreated cells were considered as the control.

#### 103 2.5. MTT assay

104 Cell viability of HCT-116 cells were determined by using by MTT test (Alarifi et al., <sup>[57]</sup>  
105 2015) after exposure to various concentration of compound (control, solvent control, 50,  
106 100, 200, 300, 400 and 500 µg/ml) over 24 hrs. After adding, MTT dye (<sup>[9]</sup>0.05 mg/ml) to  
107 each well the plate was incubated for 4 hrs. <sup>[45]</sup> The formazan crystal was solubilized in  
108 organic solvent and after 15-minute incubation at room temperature. <sup>[23]</sup> The optical density  
109 of culture palate was measured at 570 nm by using a microplate reader (BioTek  
110 Instruments, Winooski, VT, USA) with Gen5 software (version 1.09).

111 The half-maximal inhibitory concentration (IC<sub>50</sub>) 24 hr of SM-9 compound was  
112 determined on the basis of MTT test result. Based on IC<sub>50</sub> 24 hr of SM-9 compound, we  
113 have fixed 3 sub-lethal concentrations for further experiment (Table 1).

114 <sup>[0]</sup> Table 1. The half-maximal inhibitory concentration (IC<sub>50</sub>) and concentrations for SM-9  
115 compound was used on the HCT-116 cells.

IC50 -24 h = 400 µg/ml SM-9 compound	
Percentage %	Concentrations
1/8 of IC <sub>50</sub>	50 µg/ml
½ of IC <sub>50</sub>	100 µg/ml
2/3 of IC <sub>50</sub>	266 µg/ml

#### 1172.6. Evaluation of ROS generation

118 Due to exposure of SM -9 compound (50, 100, 266 µg/ml) to HCT-116 cells the  
 119 production of ROS levels was evaluated as per Ali et al., (2021) method.<sup>[11]</sup> The cells  
 120 (3x10<sup>4</sup>) was sub cultured in a black bottom culture plate (96 well) and kept in a CO<sub>2</sub>  
 121 incubator at 37°C for 24 h.<sup>[0]</sup> Later treatment to compound the culture plates were  
 122 incubated for 24 hrs.<sup>[0]</sup> Later 24 h, DCFH-DA (10µM) was mixed/ well for 35 minute at  
 123 37°C.<sup>[77]</sup> Later incubation, plate was wash away with chilled PBS and fluorescence intensity  
 124 was evaluated at 485 nm excitation and 520 nm emissions using by a microplate reader  
 125 with Gen5 software (version 1.09)<sup>[34]</sup> (Bio-Tek Instruments, Winooski, VT, USA).  
 126 Obtained results was expressed in percent of fluorescence intensity with compare with  
 127 control.<sup>[23]</sup> A separate set of experiments were carried out to assess for generation of ROS  
 128 through qualitative analysis method (Ali et al., 2021).

#### 1292.6. Protein array for detecting apoptosis

130 To find out the mechanism of apoptosis induced by SM-9 compound, a protein array  
 131 (Human Apoptosis Array C1(Code: AAH-APO-1-2) Ray Biotech, United States) was  
 132 applied for finding the part of proteins related to apoptosis. HCT-116 cells was exposed  
 133 with the SM-9 compound (0, 50, 100 and 266 µg/ml) for 24h. Extracted proteins (200 µg)  
 134 from control and exposed cells was incubated 5 h with the human apoptosis antibody  
 135 array.<sup>[0]</sup> The protein The apoptosis array membranes was scanned in a Biorad Chemi Doc

<sup>[10]</sup> 136 XRS+ Imaging System (Tide Mill Road, Units 3, 4, USA). <sup>[18]</sup> The obtained data was  
137 analyzed by using Image Lab software (Biorad Chemi Doc XRS+), and the protein signal  
138 intensity of all samples were assessed according to the kit's ((Code: AAH-APO-1-2) Ray  
139 Biotech, USA) instructions.

## 140 2.7. Gene expression

### 141 2.7.1. <sup>[64]</sup> RNA isolation and cDNA synthesis

142 The fresh cell lysate of control and SM-9 compound treated HCT cells were prepared. <sup>[21]</sup> As  
143 per manufactures protocol of RNeasy Mini Kit (Qiagen, Germany), we have isolated total  
144 RNA from the above cells. <sup>[21]</sup> The quality of RNA was determined by means of NanoDrop  
145 2000c spectrophotometer (Thermo Fisher Scientific, USA). We have synthesized cDNAs  
146 by using cDNA Synthesis Kit (BioRad, USA) as per kit protocol.

### 147 2.7.2. <sup>[29]</sup> Reverse transcription (RT-PCR)

148 The apoptotic genes such as Bax, Bcl2, and Tp53 expression in HCT cells were analyzed  
149 by using PowerUp™ SYBR™ Green Master Mix (Applied Biosystems, USA) and Light  
150 cycler 480 (Roche, Basel, Switzerland) as per kit information. We have done all  
151 experiment in duplicate. <sup>[29]</sup> The following primers for apoptotic genes expression in SM-9  
152 compound treated HCT cells was used as in Table 2. <sup>[29]</sup> GAPDH used as a housekeeping  
153 gene, and the fold change in relative quantification were determined by using this  
154 formula  $2^{-\Delta\Delta Ct}$ .

155 Table 2. List of primer sequences of apoptotic genes.

Gene	Primer F sequence	Primer R sequence
Bcl2	ATGTGTGTGGAGAGCGTCAA	GGGCCGTACAGTTCCACAAA
Bax	TGAAGCGACTGATGTCCCTG	GGGCCGTACAGTTCCACAAA
TP53	TGAAGCGACTGATGTCCCTG	CAAAGATGGTCACGGTCTGC
GAPDH	GGGAAGCTTGTCAATCAATGG	GAGATGATGACCCTTTTGGC

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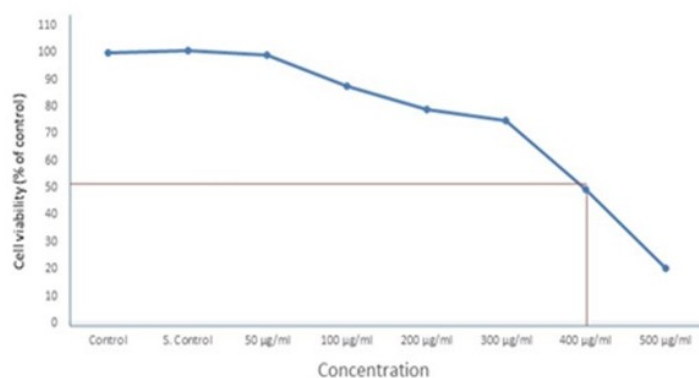
## 1572.8. Statistical analysis

158 Results were analyzed by SPSS 26.0<sup>[68]</sup> software (IBM) and expressed as mean  $\pm$  standard159 deviation (SD). \*p value 0.05, \*\*p value 0.01<sup>[7]</sup> were considered statistically

160 significant.

161

## 1623. Results

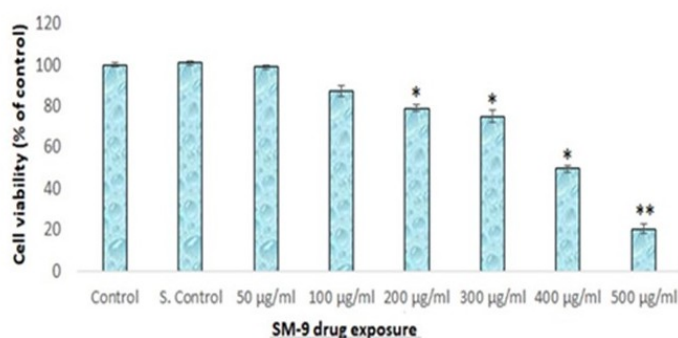
1633.1.<sup>[24]</sup> Determination IC<sub>50</sub> value of SM-9 compound164 We have calculated the median inhibitory concentration (IC<sub>50</sub> for 24 h) of SM-9 compound on165 HCT-116 cells by using MTT test result through software OrigenPro 8.5 program. The IC<sub>50</sub> value166 of SM-9 compound on HCT-116 cells was 400  $\mu$ g/ml (Figure 2, Table 1).

**Figure. 2.** Determination of IC<sub>50</sub> 24 h for SM-9 drug in HCT-116 cells Each value represents the mean  $\pm$ SE of five experiments. n= 3, \*p < 0.05 \* \*p < 0.01 vs. control.

167

1683.2.<sup>[0]</sup> Cytotoxicity

<sup>[0]</sup>▶ Cytotoxicity of SM-9 compound was determined in HCT cells by using MTT assay.<sup>[0]</sup>▶ SM-1709 compound induced toxicity in a dose-dependent manner. SM-9 compound induced high cell death in HCT cells than untreated cells. A significant toxic effect of SM-9 compound was seen in HCT cells at 500 µg/ml of SM-9 compound (Figure 3).

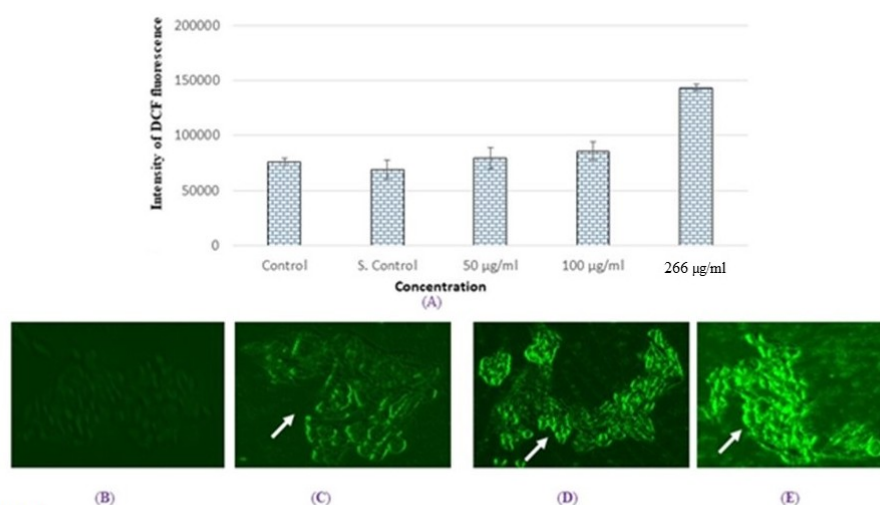


**Figure 3.** Cytotoxicity of SM-9 drug in HCT-116 cells over 24 h, as evaluated by MTT assays. Each value represents the mean  $\pm$ SE of three experiments. n= 3, \*p < 0.05 \*\*p < 0.01 vs. control.

173

### 1743.3.<sup>[44]</sup>▶ Intracellular reactive oxygen species (ROS)

175 The production of ROS was increased as exposure of compound was increased and we  
 176 have observed high intensity of green fluorescence at 266 µg/ml of SM-9 compound  
 177 (Figure 4A, E).<sup>[7]</sup>▶ In this study we have determined the production of intracellular ROS by  
 178 measuring DCF intensities in SM-9 compound exposed cells (Figure 4).<sup>[44]</sup>▶ The maximum  
 179 production of ROS was found at concentration of SM-9 compound 266 µg/ml (Figure 4  
 180A).



**Figure. 4** Production of intracellular ROS in HCT cells for 24 h due to SM-9 drugs exposure (A). Percent of DCF fluorescence intensity and generation of green fluorescence in HCT cells for 24 h (B). Control HCT cells (C). HCT cells at 50 µg/ml (D). HCT cells at 100 µg/ml (E). HCT cells at 266 µg/ml. Each value represents the mean  $\pm$ SE of three experiments. \* $p < 0.05$  vs. control. Arrow indicate green fluorescent intensity in cell as marker of ROS generation.

181

#### 1823.4. Apoptotic protein expression

183 Due to treatment of various concentration of SM-9 compound to HCT cells for 24 h, our  
 184 team member analyzed specific proteins related apoptosis and toxicity using Human  
 185 apoptosis protein array (Fig. 5a,b,c).<sup>[11]</sup> As indicated in Fig. 5, we found changes in those  
 186 apoptotic proteins.<sup>[82]</sup> Many proteins were down regulated including Bad, Bcl2, CD40, and  
 187 upregulated Bax, Caspase-3, Caspase-8, CD40L, cytochrome C, Fas, Fas, ligand, Hsp70,  
 188 HTRA, IGF-1sr, IGF-2, IGFBP5, IGFBP6, livin, P21, P27, P53, survivin, TNF- $\beta$ , TNFR-  
 189 II, TRAIL-1, TRAIL-2, and TRAIL-3, depending on its role in apoptosis pathway (Fig. 5<sup>[24]</sup>  
 190 a, b, c).<sup>[0]</sup>

	A	B	C	D	E	F	G	H	I	J	K	L	M	N
1	Pos	Pos	Neg	Neg	Blank	Blank	bad	bax	bcl-2	bcl-w	BID	BIM	Caspase3	caspase8
2	Pos	Pos	Neg	Neg	Blank	Blank	bad	bax	bcl-2	bcl-w	BID	BIM	Caspase3	caspase8
3	CD40	CD40L	ciAP-2	cytoC	DR6	Fas	FasL	Blank	HSP27	HSP60	HSP70	HTRA	IGF-I	IGF-II
4	CD40	CD40L	ciAP-2	cytoC	DR6	Fas	FasL	Blank	HSP27	HSP60	HSP70	HTRA	IGF-I	IGF-II
5	IGFBP-1	IGFBP-2	IGFBP-3	IGFBP-4	IGFBP-5	IGFBP-6	IGF-1sR	livin	p21	p27	p53	SMAC	Survivin	STNF-R1
6	IGFBP-1	IGFBP-2	IGFBP-3	IGFBP-4	IGFBP-5	IGFBP-6	IGF-1sR	livin	p21	p27	p53	SMAC	Survivin	STNF-R1
7	sTNF-R2	TNF-alpha	TNF-beta	TRAILR-1	TRAILR-2	TRAILR-3	TRAILR-4	XIAP	Blank	Blank	Neg	Neg	Neg	Pos
8	sTNF-R2	TNF-alpha	TNF-beta	TRAILR-1	TRAILR-2	TRAILR-3	TRAILR-4	XIAP	Blank	Blank	Neg	Neg	Neg	Pos

Blank: Blank spot

Neg: Negative control spot

Pos: Positive control spot

(a)



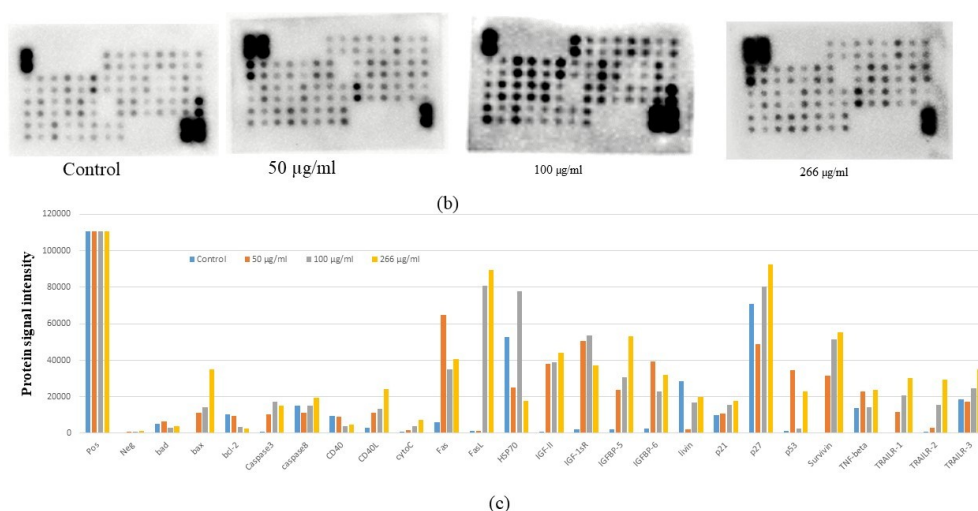


Fig. 5. Analysis of apoptosis-related proteins in HCT-116 cells due to SM-9 compound exposure (a) Template showing the location of apoptosis-related antibodies spotted on the Human Apoptosis Array C1 kit. (b) Equal amounts of protein extracts from control and treated HCT-116 cells were analyzed using the antibody array. The chemiluminescent intensities were quantified by densitometry. A positive control was used to normalize the results from different membranes. (c) Representative bar graph of the apoptotic upregulated and downregulated proteins.

192

### 193.5. Apoptotic gene expression in HCT-116 cells

194 To confirm the mechanism of toxicity induced by SM-9 compound in HCT-116 cells we

195 have determined the expression of some specific genes such as Bcl2, Bax and p53. RT-PCR

196 analysis was performed to determine the expression level of the apoptotic genes e.g.

197 Bcl2, Bax and p53 in HCT cells. The higher expression of gene expression was observed

198 at 200 µg/ml of SM-9 compound after 24 h of exposure (Figure 6 A, B, C). The p53 gene

199 was downregulated at 100 µg/ml of SM-9 compound but upregulated at 50 µg/ml and

200 266 µg/ml of SM-9 compound in cells (Figure 6 C). Other apoptotic genes, such as Bcl2

201 and Bax were expressed in HCT cells (Figure 6 A, B).

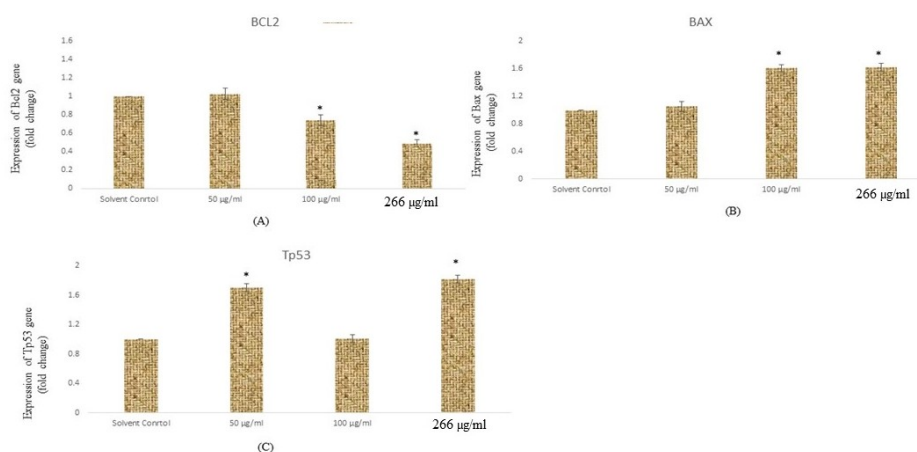


Figure 5. Expression of apoptotic genes in HCT cells exposed to SM-9 drug for 24 h. Expression of (A) Bcl2 (B) Bax, (C) p53, gene in HCT cells for 24 h. Results are expressed in average  $\pm$  SD of triplicate experiments. \* $p < 0.05$  vs. solvent control

202

203

#### 2044. Discussion<sup>[39]</sup>

205 The production of novel chemical compounds has increased significantly in recent years

206 on a global scale, and it is fascinating to use this technology to the pharmaceutical

207 industry for the benefit of human health.<sup>[39]</sup> However, a significant and concurrent

208 consequence of these growing chemical-based uses could be detrimental to the

209 ecosystem, as people may not be aware of the risk of drug exposure or the different ways

210 that chemicals can infiltrate biological systems (Yang et al., 2012). This study looked at

211 the biological reactions that occurred when the SM-9 compound was exposed to human

212 colon cancer (HCT-116) cells. We found that the SM-9 compound caused dose-

213 dependent toxicity, reactive oxygen species, and apoptosis in HCT-116 cells. The

214 finding that HCT-116 cells are susceptible to the SM-9 chemical is more significant.<sup>[11]</sup> The

215 bioactivity of SM-9 compound may be due to presence of 1-(4-chlorobenzoyl)-4-

216 (dimethylamino) group is worthy for further analysis.<sup>[0]</sup> The production level of ROS was

217 higher in HCT-116 cells at maximum concentration of SM-9 compound exposure

218 demonstrate that higher dose of this drug exposure will be more effective to minimize the  
 219 growth of cancerous cells. ROS has a major role in various cellular mechanisms, such as  
 220 cell cycle, cell proliferation, and gene expression, and ultimately the mechanism of cell  
 221 growth was stopped or cell death occurred (Almutairi et al., 2020).<sup>[6]</sup> Our investigation  
 222 showed that the SM-9 chemical was advantageous for usage as an anti-cancer medication  
 223 and sensitive to HCT-116 cells. By using RT-PCR analysis to assess the apoptotic  
 224 potential of the SM-9 compound in these cells, it was found that higher compound  
 225 concentrations had a greater apoptotic effect. Due to the correlation between the  
 226 induction of cytotoxicity, oxidative stress, and apoptotic gene expression, it was evident  
 227 that exposure to a more dose of SM-9 chemical was more effective. In HCT-116 cells, all  
 228 of these results proved the SM-9 compound's anticancer capabilities.<sup>[25]</sup> The untreated HCT-  
 229 116 cells showed decreased protein signal fluorescence intensity of various proteins such  
 230 as bad, bax, bcl2, p53, p21, Hsp70, p27, TRAIL-1, TRAIL-2, TRAIL-3 and TNF  $\beta$  with  
 231 comparison to the SM-9 compound treated HCT-116 cells. The current finding indicated  
 232 the effect of SM-9 compound to cause inhibition of the bad, bcl2 signaling pathway in the  
 233 HCT-116 cells. These findings suggest that bad, bcl2 regulation could be chunked via  
 234 SM-9 compound.<sup>[6]</sup> The results revealed that both types of apoptotic pathways may be  
 235 involved in the induction of the apoptosis in HCT-116 cells by upregulating caspase-8  
 236 and p53 and reducing bad, bcl2 into the cytoplasm and nucleus.<sup>[10]</sup> The concerns of the  
 237 mitochondrial-intrinsic actions indicated by mRNA and the upregulation of p53, bax, and  
 238 cytochrome c could elicit mitochondrial damage and dysfunction by enhancing the levels  
 239 of caspase-8.  
 240 Protein arrays are useful because it can provide a map of known cell apoptotic and

241 signaling proteins.<sup>[12]</sup>

242 Protein arrays have shown a unique ability to analyze signaling pathways using small numbers of cultured cells or cells (Grubb et al., 2003). Using this approach, untreated and treated HCT-116 cells of protein lysate is arrayed onto nitrocellulose-coated slides. Main technological components of this method offer unique advantages over to detect multi-protein expression.

247 In this study a useful methodology for the synthesis of a novel chemical compound was described.<sup>[7]</sup> The results showed that SM-9 compound has high efficacy and should be given particular consideration in anticancer activity.<sup>[7]</sup> Furthermore, this finding demonstrated that SM-9 compound killed the colon cancer cells and induced apoptosis through various activated proteins, gene such as caspase-8, p53, NF- $\beta$ , and bax.<sup>[23]</sup> The results established that SM-9 compound may be applied as a medicinal drug alone or in combination with other chemotherapeutics for considering different types of cancer cells.<sup>[35]</sup> In the future, we will investigate about the mechanism of toxicity due to SM-9 compound in vivo experiments

256 Declaration of conflicting interests

257 None

258 Data Availability Statement

259 The original contributions presented in the study are included in the article.

260 Funding Statement

261 None

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265<sup>[24]</sup>Supplementary Material

266Associated with this article are the <sup>1</sup>H and <sup>13</sup>C NMR spectra of SM-9 compound.

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