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### **OPEN** Synthesis and *in-vitro* anti-proliferative with antimicrobial activity of new coumarin containing heterocycles hybrids

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A series of new coumarin-N-heterocyclic hybrids, coumarin-quinolines 7a-e, coumarin-acridines 10b, c and coumarin-neocryptolepines 13b, c were synthesized and evaluated for their anticancer and antimicrobial activities. The structures of all synthesized hybrids were confirmed by FT-IR, <sup>1</sup>H-NMR, <sup>13</sup>C-NMR, and MS spectrometry. The anti-proliferative activity of hybrids 7a-e, 10c and 13c were bio-evaluated using MTT-assay against colon (CaCo-2), lung (A549), breast (MDA-MB-231), and hepatocellular carcinoma (HepG-2) human cancer cell lines using doxorubicin as a reference drug. The results demonstrated that, all hybrids displayed moderate to good anti-proliferative activity against the cell lines. The most active hybrids were 7a–d and 10c against CaCo-2 cancer cell line with  $IC_{50}$ : 57.1, 52.78, 57.29, 51.95 and 56.74 µM, and selectivity index 1.38, 1.76, 2.6, 1.96 and 0.77; respectively. While, 7a,d were potent against A549 cancer cell line with IC<sub>50</sub>: 51.72, 54.8 µM and selectivity index 1.5, 0.67; respectively. Moreover, 7c showed the most potency against MDA-MB-231 cancer cell line with IC\_{50}: 50.96  $\mu M$  and selectivity index 2.20. Interestingly, docking results revealed that binding energy of the current compounds showed marked affinity values ranging from -6.54 to -5.56 kcal with interactions with the reported key amino acid SER 79. Furthermore, the antimicrobial activity of the synthesized hybrids 7a-e, 10b,c, 13b and 13c were evaluated against Gram-positive and Gramnegative bacterial and fungal strains. The hybrids 10b, 13b, 10c, and 13c exhibited broad-spectrum antibacterial activity against E.coli, S. mutans, and S. aureus with MIC from 3.2 to 66 µM, this hybrids also displayed antifungal activity against C. albicans with MIC values ranging from 0.0011 to 29.5 µM. In-silico investigation of the pharmacokinetic properties indicated that tested hybrids had high GI absorption, low Blood Brain Barrier (BBB) permeability in addition to cell membrane penetrability.

In recent years, the number of people suffering from cancer and multi-resistant infections has increased, such that both diseases are already seen as current and future major causes of death. Moreover, chronic infections are one of the main causes of cancer, due to the instability in the immune system that allows cancer cells to proliferate. Likewise, the physical debility associated with cancer or with anticancer therapy itself often paves the way for opportunistic infections. Though advances in cancer therapy and diagnosis have considerably improved life expectancy<sup>1</sup>, the overall survival rate of patients remains poor<sup>2</sup>. Disseminated cancer at the time of diagnosis and acquisition of tumor resistance are two main reasons. The growing knowledge of the biochemical pathways

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involved in a disease process increases the possibility to develop new approaches to treat this disease<sup>3,4</sup>. However, there are two main challenges to using chemotherapy successfully in cancer treatment are tumor cell resistance and the need for tumor cell specificity to prevent toxicity to normal tissues. Consequently, targeting pathways that selectively inhibit cancer cells growth and metastasis may provide successful anticancer therapy<sup>5</sup>. Due to the possibility of life-threatening infections developing during anticancer treatment for cancer patients which are associated with prolonged hospital stay and poor quality of life<sup>6</sup>, many researchers are working to synthesize new drugs that have both anticancer and antimicrobial activity<sup>7–9</sup>.

During the last few years molecular hybridization became a powerful tool in drug design and discovery offering an attractive approach to obtain better drugs for the treatment of a large variety of human diseases including cancer and microbial illness. One of the methods used for the construction of hybrid molecules combines two or more drug pharmacophores in a single multi-functional molecule using a linker chain. The main goals of this pharmacophore merging approach consist in the interaction of the resulting molecule with dual or multiple targets, amplifying the biological activity and specificity, reducing the known side effects associated with each hybrid part, reducing the drug-drug interactions when compared with conventional classic drugs<sup>10-12</sup>. In this context, coumarin constitutes a unique motif for the construction of various classes of biologically active analogues with wide variety of pharmacological activities<sup>13</sup>, such as antidepressants<sup>14</sup>, antimicrobials<sup>15–18</sup>, antioxidants<sup>19–21</sup>, anti-inflammatories<sup>20,22</sup>, antinociceptives<sup>23</sup>, antitumors<sup>24–26</sup>, antiasthmatics<sup>27</sup>, antivirals<sup>28</sup>, antifungal<sup>29</sup> and anti-coagulant activities.<sup>30,31</sup> Previous studies showed that the hybridization of coumarin at C-4 and C-3 positions with heterocycles have synergistic effect on biological activity.<sup>32–38</sup> Possible modifications of the coumarin core structures with respect to improving their anticancer and antimicrobial activities would be the installation of different side chains at position 3 and the identity of the spacer influences the flexibility of the structure and hence weakens or enforces interaction with certain targeted biomolecules. Such substitution patterns would affect and improve biological activity. Based on the forementioned information, it might be assumed that the installation of biologically effective N-heterocyclic scaffolds with natural origin such as quinoline<sup>39</sup>, acridine<sup>40</sup> and neocryptolepine<sup>41</sup> motifs into the coumarin core structure would improve the coumarin pharmacological profile as anticancer and antimicrobial agent with synergistic effects when compared with each moiety separately as depicted in Fig. 1. Herein, a series of new coumarin based heterocyclic hybrids are synthesized, and their anti-proliferative and antimicrobial activity are evaluated. In addition, a molecular docking study was executed to confirm binding of hybrids with the target protein structures.

#### Materials and methods Materials

4,7-Dichloroquinoline 97%, ethylenediamine 99%, hydrazine hydrate 98%, 4,4'-methylenedianiline 97%, triethylamine 99%, coumarin-3-carboxylic acid 99%, were purchased from Sigma Aldrich, MO, USA. Also, reagents as 1,3-Diaminopropane 99%, 1,4-Phenylenediamine, thionyl chloride 97% and solvents as ethanol 98%, dichloromethane 98%, and petroleum ether with 60–80 °C, were supplied from LOBA Chemie, Mumbai, India and used without any further purification. The prepared starting materials such as 9-chloroacridne and 11-chloroneocryptolepine and their corresponding free amines were prepared as reported<sup>42,43</sup>.

#### Cell lines and microbial strains

*Cell lines*; WISH cell line (Human amnion-derived normal epithelial cells, CCL-25), *MDA-MB-231* cell line (Human, mammary gland/breast adenocarcinoma, HTB-26), *CaCo-2* cell line (Human Colorectal



Figure 1. Design of coumarin N-heterocyclic hybrids as potent antiproliferative and antimicrobial agents.

adenocarcinoma, epithelial cells, HTB-37), *A549* cell line (Human Lung adenocarcinoma, epithelial cells, CCL-185) cell lines and *HepG-2* (liver hepatocellular carcinoma, HEPG2, HB-8065) were obtained from ATCC, originally with license and serial numbers as mentioned.

Microbial strains: Escherichia coli, pseudomonas aeruginosa, Staphylococcus aureus, Streptococcus mutans and Candida albicans were collected from the microbial culture collection of Biopharmaceutical Products Research Department, Genetic Engineering and Biotechnology Research Institute, SRTA-City, New Borg El-Arab city, Alexandria, Egypt.

#### Molecular docking

Molecular docking studies were effectuated using and visualized on *Biovia DS*-2021 online free software to examine its affinity to topoisomerase II(Topo-II). Topo-II protein (pdb: 3FOE) was downloaded from protein data bank<sup>44</sup>. Protein preparation was done by deleting water molecules, selecting the co-crystalized ligand to be used as a current selection for defining the active site of the Top II and create binding site sphere attributes which was then extended to cover the whole protein. The ligand was then deleted, polar hydrogen was added, and files are saved as optimized protein. Ligand optimization and docking was performed by selecting dock ligand in lib-dock to prepare the ligand automatically in the program using the input site sphere attributes which was created upon protein optimization and minimize energy for the macromolecules before docking. The selected minimization force field was CHARMm.

#### Instrumental characterization

<sup>1</sup>H-NMR and <sup>13</sup>C-NMR spectroscopic analyses were carried out with Bruker Germany; 400 and 100 MHz; respectively. Reports of chemical shifts were made in parts per million (ppm) as it relates to the respective solvent (DMSO-d<sub>6</sub>). Also, **FT-IR** spectra were performed with Alpha, Bruker Germany at faculty of Science, Zagazig University. Thermo-Scientific GCMS model ISQ, USA, was used for the mass-spectra, which were conducted on the direct inlet part; at the regional center for Mycology and Biotechnology (RCMB). Exploring the characteristic fragmentation using Electron Impact mode and expected molecular weight at 70 eV. Melting point was recorded using scientific melting point apparatus without correction.

#### General synthesis of coumarin- N-heterocyclic hybrids 7a–e, 10b, c and 13b, c.

Coumarin-3-carbonyl chloride **3** (0.30 g, 1.27 mmol), appropriate amines **6a–e**, **9b,c** and **12b,c** (1.27 mmol) dissolved in  $CH_2Cl_2$  (2 mL), triethylamine (0.39 g, 3.81 mmol) was added dropwise with stirring at room temperature (25°C). Progress of the reaction was observed by thin layer chromatography (TLC) till starting materials were consumed (12 h.). The reaction mixture was poured into ice/water, and extracted three times with  $CH_2Cl_2$ , the organic layer was collected, dried followed by removing of the solvent residue using rotatory evaporator. The precipitated solid was filtered off, dried and recrystallized from ethanol to give pure **7a–e**, **10b,c and 13b,c** in good yields.

#### N'-(7-Chloroquinolin-4-yl)-2-oxo-2H-chromene-3-carbohydrazide (7a)

Red solid, yield (0.37 g, 80%) m.p: 294–296 °C, **FT-IR** (KBr) cm<sup>-1</sup> v: 3265 (NH), 1700 (C–C=O), 1655 (C–C=O), 1603 (C=C<sub>Ar</sub>), 1532 (C=N), 1202 (C–C). <sup>1</sup>**H-NMR**  $\delta$  ppm: 7.44–8.25 (m, 8H, CH<sub>Ar</sub>), 8.42 (s, 1H, CH<sub>Ar</sub>), 8.89 (s, 1H, CH=N<sub>Ar</sub>), 10.82 (br.s, 1H, NH), 14.67 (br.s, 1H, NH). <sup>13</sup>**C-NMR**  $\delta$  ppm: 116.25, 116.73, 118.36, 119.45, 122.11, 123.71, 125.04, 125.25, 125.69, 125.90, 128.18, 128.79, 130.22, 147.27, 151.97, 153.84. **EI-MS**, m/z (C<sub>19</sub>H<sub>12</sub>ClN<sub>3</sub>O<sub>3</sub>) calcd, 365.06; found, 365.48 [M<sup>+</sup>]=13.36%, 367.56 [M+2]<sup>+</sup>.

#### *N-(2-((7-Chloroquinolin-4-yl) amino) ethyl)-2-oxo-2H-chromene-3-carboxamide (7b)*

Yellowish solid, yield (0.46 g, 92%) m.p: 274–276 °C, **FT-IR** (KBr) cm<sup>-1</sup> v: 3409 (NH), 2954(CH), 1685 (C–C=O), 1648 (O–C=O), 1610 (C=C<sub>Ar</sub>), 1538 (C=N), 1240 (C–C). <sup>1</sup>**H-NMR**  $\delta$  ppm: 3.64 (m, 2H, CH<sub>2</sub>), 3.89 (m, 2H, CH<sub>2</sub>), 7.47–8.41 (m, 8H, CH<sub>Ar</sub>), 8.54 (s, 1H, CH<sub>Ar</sub>), 8.91 (s, 1H, CH=N<sub>Ar</sub>), 13.32 (s, 1H, NH). <sup>13</sup>**C-NMR**  $\delta$  ppm: 29.00, 37.81, 116.13, 117.43, 118.75, 123.96, 124.19, 125.14, 127.49, 130.31, 133.42, 134.16, 147.76, 150.14, 151.89, 153.92, 160.19, 161.70. **EI-MS**, m/z (C<sub>21</sub>H<sub>16</sub>ClN<sub>3</sub>O<sub>3</sub>) calcd, 393.09; found, 393.32[M<sup>+</sup>]=11.18%, 395.43 [M+2]<sup>+</sup>.

#### N-(3-((7-Chloroquinolin-4-yl) amino) propyl)-2-oxo-2H-chromene-3-carboxamide (7c)

White solid, yield (0.42 g, \$1%) m.p: 191-1946 °C, **FT-IR** (KBr) cm<sup>-1</sup> v: 3331 (NH), 2925(CH), 1710 (C–C=O), 1640 (O–C=O), 1609 (C=C<sub>Ar</sub>), 1532 (C=N), 1213(C–C). <sup>1</sup>**H-NMR**  $\delta$  ppm: 1.93 (m, 2H, CH<sub>2</sub>), 3.46 (br.m, 2H, CH<sub>2</sub>), 3.64 (br.m, 2H, CH<sub>2</sub>), 7.41–7.97 (m, 8H, CH<sub>Ar</sub>), 8.27 (m, 1H, CH<sub>Ar</sub>), 8.81 (m, 1H, CH=N<sub>Ar</sub>). <sup>13</sup>C-NMR  $\delta$  ppm: 27.66, 37.33, 53.00, 116.26, 117.48, 118.52, 119.19, 124.18, 125.31, 127.04, 130.34, 133.82, 134.24, 147.38, 148.53, 151.59, 153.94, 160.42, 161.53. **EI-MS**, m/z (C<sub>22</sub>H<sub>18</sub>ClN<sub>3</sub>O<sub>3</sub>) calcd, 407.10; found, 407.66[M<sup>+</sup>] = 9.97%, 409.78 [M+2]<sup>+</sup>.

#### N-(4-((7-Chloroquinolin-4-yl) amino) phenyl)-2-oxo-2H-chromene-3-carboxamide (7d)

Brownish yellow solid, yield (0.47 g, 83%) m.p: 258–2606 °C, **FT-IR** (KBr) cm<sup>-1</sup> v: 3390 (NH), 1688 (C–C=O), 1664 (O–C=O), 1609 (C=C<sub>Ar</sub>), 1539 (C=N), 1203 (C–C). <sup>1</sup>**H-NMR**  $\delta$  ppm: 7.39–8.49 (m, 12H, CH<sub>Ar</sub>), 8.53 (m, 1H, CH<sub>Ar</sub>), 8.93 (s, 1H, CH=N<sub>Ar</sub>), 9.62 (br.s, 1H, NH), 10.73 (s, 1H, NH). <sup>13</sup>**C-NMR**  $\delta$  ppm: 116.44, 118.00, 118.63, 120.14, 124.17, 124.76, 125.03, 125.85, 126.58, 130.46, 134.87, 135.86, 136.01, 144.70, 147.50, 147.91, 149.40, 152.72, 154.04, 160.62. **EI-MS**, m/z (C<sub>25</sub>H<sub>16</sub>ClN<sub>3</sub>O<sub>3</sub>) calcd, 441.09; found 441.72 [M<sup>+</sup>] = 20.22%, 443.81 [M+2]<sup>+</sup>.

#### N-(4-(4-((7-Chloroquinolin-4-yl) amino) benzyl) phenyl)-2-oxo-2H-chromene-3-carboxamide (7e)

Yellow solid, yield (0.55 g, 82%) m.p: 251–2566 °C, **FT-IR** (KBr) cm<sup>-1</sup> v: 3202 (NH), 1702 (C–C=O), 1660 (O–C=O), 1597 (C=C<sub>AT</sub>), 1545 (C=N), 1205 (C–C). <sup>1</sup>H-NMR  $\delta$  ppm: 4.02 (m, 2H, CH<sub>2</sub>), 7.35–8.18 (m, 8H, CH<sub>AT</sub>), 8.25 (m, 4H, CH<sub>AT</sub>), 8.50 (m, 4H, CH<sub>AT</sub>), 8.81 (m, 1H, CH<sub>AT</sub>), 8.90 (s, 1H, CH=N<sub>AT</sub>), 10.67 (s,1H, NH), 10.90 (br. s, 1H, NH). <sup>13</sup>C-NMR  $\delta$  ppm: 52.09, 116.20, 120.29, 122.19, 124.41, 125.36, 125.88, 126.00, 127.28, 128.21, 128.88, 129.40, 130.29, 135.53, 140.26, 141.46, 148.90, 152.05. **EI-MS**, m/z (C<sub>32</sub>H<sub>22</sub>ClN<sub>3</sub>O<sub>3</sub>) calcd, 531.13; found 531.94 [M<sup>+</sup>] = 11.32%, 533.68 [M+2]<sup>+</sup>.

#### *N-(2-(Acridin-9-ylamino)ethyl)-2-oxo-2H-chromene-3-carboxamide (10b)*

Brownish yellow solid, yield (0.4 g, 77%) m.p: <sup>3</sup>300, **FT-IR (KBr)** cm<sup>-1</sup> v: 3363(NH), 3390 (NH), 2918(CH), 1698 (C–C=O), 1659 (O–C=O), 1611 (C=C<sub>Ar</sub>), 1519 (C=N), 1238 (C–C). <sup>1</sup>H-NMR  $\delta$  ppm: 3.71 (m, 2H, CH<sub>2</sub>), 4.03 (m, 2H, CH<sub>2</sub>), 7.19–8.28 (m, 12H, CH<sub>Ar</sub>), 8.85 (s, 1H, CH<sub>Ar</sub>), 9.19 (s, 1H, NH) 11.75 (s, 1H, NH). <sup>13</sup>C-NMR  $\delta$  ppm: 50.72, 56.01, 116.08, 117.32, 118.44, 118.66, 120.47, 120.97, 122.54, 125.09, 125.99, 130.27, 133.42, 134.07, 140.88, 147.58, 153.87, 160.33, 161.37. **EI-MS**, m/z (C<sub>25</sub>H<sub>19</sub>N<sub>3</sub>O<sub>3</sub>) calcd, 406.14; found 406.01[M<sup>+</sup>] = 10.7%.

#### *N*-(3-(Acridin-9-ylamino) propyl)-2-oxo-2H-chromene-3-carboxamide (10c)

Brownish yellow, yield (0.44 g, 82%) m.p: 261–263 °C, **FT-IR** (KBr) cm<sup>-1</sup> v: 3235 (NH), 2922 (CH),1679 (C–C=O), 1596 (C=C<sub>Ar</sub>), 1531 (C=N), 1260 (C–C). <sup>1</sup>**H-NMR**  $\delta$  ppm: 1.21 (br.m, 2H, CH<sub>2</sub>), 4.02 (m, 2H, CH<sub>2</sub>), 7.24–7.94 (m, 10H, CH<sub>Ar</sub>), 8.23 (br.m, 2H, CH<sub>Ar</sub>), 8.83 (m, 1H, CH<sub>Ar</sub>), 11.78 (s, 1H, NH). <sup>13</sup>**C-NMR**  $\delta$  ppm: 29.07, 37.01, 116.08, 117.34, 118.43, 118.79, 120.48, 120.99, 122.11, 125.07, 126.00, 130.18, 133.44, 134.07, 140.90, 147.22, 153.82, 161.25, 176.77. **EI-MS**, m/z (C<sub>26</sub>H<sub>21</sub>N<sub>3</sub>O<sub>3</sub>) calcd, 423.16; found 423.10 [M<sup>+</sup>] = 15.84%.

*N*-(2-((5-*Methyl*-5*H*-indolo[2,3, *b*] quinolin-11-*y*]) amino) ethyl)-2-oxo-2*H*-chromene -3-carboxamide (13b) Brown solid, yield (0.5 g, 85%) m.p: 158–160, **FT-IR (KBr) cm**<sup>-1</sup> **v**: 3202(NH), 3327 (NH),1705 (C–C=O), 1659 (O–C=O), 1611 (C=C<sub>Ar</sub>), 1519 (C=N), 1238 (C–C). <sup>1</sup>**H**-NMR δ ppm: 3.65 (br.m, 2H, CH<sub>2</sub>), 4.05 (s, 3H, N-CH<sub>3</sub>), 4.16 (br.m, 2H, CH<sub>2</sub>), 7.06–7.92 (m, 12H, CH<sub>Ar</sub>), 8.51 (m, 1H, CH<sub>Ar</sub>), 12.65 (br. s, 1H, NH). <sup>13</sup>**C**-NMR δ ppm: 32.15, 47.71, 115.72, 116.03, 116.39, 118.08, 118.28, 120.56, 122.21, 123.88, 124.79, 125.09, 130.22, 130.48, 134.10, 137.16, 147.20, 148.65, 152.13, 153.73, 156.29, 159.82, 161.54. **EI-MS**, m/z (C<sub>28</sub>H<sub>22</sub>N<sub>4</sub>O<sub>3</sub>) calcd, 462.17; found 462.53 [M]<sup>+</sup> = 39.6%.

$$\begin{split} &N-(3-((5-methyl-5H-indolo[2,3-b]~quinolin-11-yl)~amino)~propyl)-2-oxo-2H-chromene-3-carboxamide~(13c) \\ Brown solid, yield~(0.48~g, 79\%)~m.p:~158-160~°C,$$
**FT-IR** $(KBr) cm<sup>-1</sup> v:~3202(NH), 3327 (NH), 2967 (CH), 1705 (C-C=O), 1659 (O-C=O), 1611 (C=C_{Ar}), 1519 (C=N), 1238 (C-C). <sup>1</sup>H-NMR <math>\delta$  ppm: 1.97(m, 2H, CH<sub>2</sub>), 3.92 (br.m, 2H, CH<sub>2</sub>), 4.16(m, 3H, CH<sub>3</sub>), 7.06-7.92 (m, 12H, CH<sub>Ar</sub>), 8.65 (m, 1H, CH<sub>Ar</sub>). <sup>13</sup>C-NMR  $\delta$  ppm: 32.46, 36.71, 45.34, 115.72, 115,98, 116.09, 118.33, 118.43, 118.58, 120.83, 122.13, 123.72, 123.92, 124.59, 125.10, 130.24, 130.71, 134.03, 137.27, 147.15, 148.46, 153.80, 160.06, 161.20. **EI-MS,** m/z (C<sub>29</sub>H<sub>24</sub>N<sub>4</sub>O<sub>3</sub>) calcd, 476.18; found 476.40 [M<sup>+</sup>] = 24%. \end{split}

#### **Bio-evaluation assays**

#### Safety assays and anti-proliferative activities of Coumarin hybrids

The safety assays of coumarin hybrids on noncancerous cell lines and their anticancer effects on cancerous cells were gauged utilizing MTT-assay (*Promega*) in accordance with the instruction protocol. Starting from 60 to 200  $\mu$ M, serial dilutions of coumarin hybrids were prepared in sterile DMSO to be treated cell lines at final concentrations 30 to 100  $\mu$ M. The treated cells were incubated for 2 days, and the cellular cytotoxicity was detected by quantifying the solubilized formazan in DMSO at 570 nm. The inhibition concentration of 50 (IC<sub>50</sub>) was calculated from the cytotoxicity% curve using *GraphPad prism 9*.

#### Selectivity index of Coumarin hybrids

Cancer cell selectivity index of Coumarin hybrids samples were measured as explained by <sup>45</sup>, with a minor modification; (SI =  $IC_{50}nc/IC_{50}cc$ ), where  $IC_{50}nc$  refers to the value of  $IC_{50}$  of the coumarin hybrids compounds on normal cells, while  $IC_{50}cc$  refers to the  $IC_{50}$  of the coumarin hybrids compounds on cancer cell line.

#### Antimicrobial activities of coumarin hybrid

The antimicrobial activity of coumarin hybrids samples were checked against different multiple drug resistant microorganisms (*Escherichia coli, pseudomonas aeruginosa, Staphylococcus aureus* and *Streptococcus mutans* and *Candida albicans*) using various concentrations (30 to 100  $\mu$ M). An aliquot of 100.0  $\mu$ l of each sample concentration was added to an equal volume of each microbial growth dilution (about 10<sup>6</sup> CFU/ml) and inoculated into 96 well plate. Furthermore, 100.0  $\mu$ l of LB media was added to 100.0  $\mu$ l of microbial growth to set as the negative control group. After that, the inoculated plates were incubated overnight at 37 °C then, the microbial turbidity was measured using automated ELIZA microplate reader (BINDER BIOTECK E LX 800) adjusted at 620 nm. The microbial inhibition percentage after treatments were quantified using the following equation:

Inhibition percentage = 
$$(A - A1/A0) \times 100.$$
 (1)

where, A: the treatment group absorbance, A1: the blank absorbance, and A0: the control group absorbance. The Minimal Inhibitory Concentrations (MIC) were estimated and expressed as the lowest concentration of the tested samples which resulted in microbial growth inhibition.

#### **Ethical statement**

All research studies followed the Helsinki World Medical Association's Declaration: Ethical Medical Research Principles Involving Human Subjects and were approved by the ethics committee at Menoufia University in Egypt, Faculty of Science. Where, no animal experiments were conducted in this research, while all used cell lines were obtained from ATCC with original and recognized serial number of each cell line as aforementioned in section of cell line and microbial strains.

### Results and discussion

#### General synthesis of coumarin- N-heterocyclic hybrids

Synthesis of coumarin-3-carbonyl chloride

The key intermediate acid chloride **3** was prepared in good yield as pale-yellow crystals according to published method<sup>21,46,47</sup>, by the reaction of coumarin-*3*-carboxylic acid **1** and thionyl chloride (SOCl<sub>2</sub>) refluxed for 2 h as depicted in Scheme 1.

#### Synthesis of 4 bis-aminosubstituted heterocycles

The reaction of 4,7-dichloroquinoline **4** with hydrazine, aliphatic and aromatic diamines **5a**–**e** in presence of triethyl amine as a base catalyst afforded **6a**-**e** in good yields via nucleophilic aromatic substitution ( $S_{NAr}$ ). The synthesized compounds showed analytical data consistent with previously published results<sup>4,48</sup>.

The synthetic pathway for formation of coumarin-quinoline hybrids 7a-e was achieved by the reaction of **3** with diamines **6a-e**, in equimolar ratio in presence of triethyl amine as a base to afford the corresponding hybrids **7a-e**, in good to excellent yields (80–92%) as given in Scheme 1. Additionally, the target hybrids **10b,c** were synthesized by the condensation of 9-chloroacridine **8** with amines **5b**, **5c** afforded the **9b** and **9c**, according to the reported method.<sup>49,50</sup> Further reactions of **9b,c** with **3** in equimolar ratio in presence of excess of triethyl amine in good yield (77,82%) as depicted in Scheme 2.

Furthermore, the condensation of 11-chloroneocryptolpine 11 with 5b and 5c yielding 12b,c in good yield (79%) according to the reported methods<sup>51,52</sup>. While, the reaction of 12b, 12c with 3 in equimolar ratio in presence of excess of triethyl amine afforded the corresponding coumarin-neocryptolepine hybrids 13b,c in good yields (79,85%), as shown in scheme 2. The progress of reactions was easily monitored by TLC using benzene and ethanol as eluent mixture (3:1), till starting materials are consumed.

#### Chemical structure verification by FTIR and NMR analysis

*FTIR analysis of coumarin hybrids* In structure elucidation of 7 **a-e**, **10b,c** and **13b,c**, FT-IR spectra show a strong absorption band for  $v_{(NH)}$  at: 3265, 3409, 3331, 3390 and 3390 cm<sup>-1</sup> for **7a-e**; respectively, 3235,3390 cm<sup>-1</sup> for **10b,c**, 3202, 3327 and 3372 cm<sup>-1</sup> for **13b,c**. Moreover, the absorption band for v (C–C=O) at 1700, 1685, 1710, 1688 and 1702 cm<sup>-1</sup> for **7a-e** respectively, 1679,1698 cm<sup>-1</sup> for **10b,c**, and 1705 cm<sup>-1</sup> for **13b,c** which confirmed the presence of carbonyl group. On the other hand, the absorption band for v (O–C=O) at v: 1655, 1648, 1640,



Scheme 1. Synthetic pathway of coumarin-3-carbonyl chloride; synthesis of 4-bis amino quinoline **6a–e** and synthesis of coumarin hybrids **7a–e**.



Scheme 2. Synthesis of coumarin-acridine hybrid 10, and synthesis of coumarin-neocryptolepine hybrid 13.

1664 and 1660 cm<sup>-1</sup> for **7 a-e**; respectively, 1659 cm<sup>-1</sup> for **10b**, 1659 and 1650 cm<sup>-1</sup> for **13b**,**c** which confirmed the presence of lactone. Moreover, the absorption band for  $\upsilon$  (C=C<sub>Ar</sub>) at 1603, 1610, 1609, 1609, and 1597 cm<sup>-1</sup> for **7a-e**; respectively,1596, 1611 cm<sup>-1</sup> for **10b**,**c** 1611 and 1609 cm<sup>-1</sup> for **13b**,**c**. On the other hand, the absorption band for  $\upsilon$  (C=N) at: 1532, 1538, 1532, 1539 and 1545 cm<sup>-1</sup> for **7a-e**; respectively, 1531,1568 cm<sup>-1</sup> for **10b**,**c**, 1519 and 1538 cm<sup>-1</sup> for **13b**,**c**. Furthermore, the absorption band for  $\upsilon$  (C–C) at: 1202, 1240, 1213, 1203, and 1205 for **7a-e**; respectively, 1238 cm<sup>-1</sup> for **10b**,**c**, 1238 and 1245 cm<sup>-1</sup> for **13b**,**c**.

*NMR analysis of coumarin hybrids* <sup>1</sup>H-NMR spectrum shows multiple  $\delta$  3.46 and 3.89 ppm which correspond to the aliphatic spacers in **7b**. Moreover, the spectra showed multiple with  $\delta$ : 1.93 ppm, broad multiple with  $\delta$ : 3.46 and 3.64 ppm, that confirmed the presence of aliphatic spacer in **7c**. On the other hand, spectra showed multiple with  $\delta$ : 4.02 ppm which confirmed the presence of (Ph-CH<sub>2</sub>-Ph) for **7e**, showed broad multiple with  $\delta$ : 4.16, 4.05 for which confirmed the presence of (N-CH<sub>3</sub>) for **13b,c**. On the other hand, coumarin (CH<sub>Ar</sub>) appear with  $\delta$ : 8.42, 8.54, 8.26, 8.53, 8.81 ppm for **7a–e**; respectively, 8.83, 8.85 ppm for **10b,c**, 8.65 and 8.51 ppm for **13b,c**. Furthermore, (HC=N<sub>Ar</sub>) appear with  $\delta$ : 8.89, 8.91, 8.81, 8.93 and 8.90 ppm for **7a–e** respectively. Moreover, the spectra showed broad singlet with  $\delta$ : (NH) 10.82, 14.67, 13.32, 9.62, 10.73, 10.67, 10.90 ppm for **7a–e**, and 9.19, 11.75, and 11.78 ppm for **10b,c** and 12.65 ppm for **13b**.

While <sup>13</sup>C-NMR spectra showed Peak of (*N*-CH<sub>3</sub>) for **13b,c** at 32,46 and 32.15 ppm. On the other hand, the coumarin (HC=C) appears at:  $\delta = 119.45$ , 118.75, 119.19, 120.14, 120.29 ppm respectively for **7a–e**, 120,99 120.47 ppm for **10** and 120,83, 120.56 ppm for **13**. Moreover, lactone (O–C=O) appear at  $\delta$ : 151.97, 160.19, 160.42, 154.04, 152.05 ppm; respectively for **7a–e**, 161.25, 160.33 ppm for **10b,c** and 159.82,160.06 ppm for **13b,c**. Furthermore, the carbonyl group (C–C=O) appears at  $\delta$ : 153.84, 161.70, 161.53, 160.62, 152.05 ppm for **7a–e**; respectively, 176.77,161.37 ppm for **10b,c** and 161,20, 161.54 ppm for **13b,c**. on the other hand, for the mass spectra for synthesized hybrids showed coincidence of the molecular ion peak with their expected molecular weight.

#### **Bioevaluation assessment**

#### Safety assays and anticancer activities of coumarin hybrids compounds

The safety profiles on *WISH* cells and the anticancer potentialities of coumarin hybrids compounds against *MDA-MB-231*, *CaCo-2*, *A549* and *HepG-2* cell lines were measured utilizing MTT-assay (Fig. 2A,B). The concluded data indicated that the compounds **7c**, **7d** and **13c** were the safest compounds on *WISH* cells with  $IC_{50}$  values of 112.39, 138.39 and 95.88; respectively (Table 1). While, among the control samples, **1** and coumarin (**Ct**) samples were safer than the other coumarin hybrids compounds with  $IC_{50}$  values of 333.57 and 176.36 µM; respectively (Fig. 2A and Table 1).

On the other hand, the most active compounds showed different anti-proliferative profiles on different cancerous cell lines. For example, **7c** and **7d** were the most effective compounds against *CaCO-2* cell line with  $IC_{50}$ values of 57.29 and 51.59  $\mu$ M, with anticancer selectivity index 1.9 and 2.6; respectively (Fig. 2A). On *A549* cell lines, the treatments **7a**, **7b** and **13c** were the most significant treatments with selectivity index of 1.5, 1.6 and 1.4; respectively (Table 2). Also, **7c** was the most potent treatment against *MDA-MB-231* cell line with SI 2.2



**Figure 2.**  $IC_{50}(A)$  and SI (**B**) values of tested synthesized coumarin hybrid compounds.

Cell line	Ct	1	7a	7b	7c	7d	7e	10	13	DOX
WISH	176.36	333.575	79.2825	92.92	112.39	138.39	93.065	56.744	95.88	1.7635
CaCo-2	99.46	1111.5	57.1	52.785	57.29	51.95	86.31	144.775	84.755	2.91
A549	259.67	351.475	51.72	54.815	97.395	203.99	151.87	130.097	65	3.777
MDA-MB-231	111.19	132.32	211.315	154.04	50.965	102.625	70.54	108.947	89.94	0.655
HepG-2	231.67	924.325	75.42	88.69	133.11	82.835	350.645	107.197	70.95	1.175

Table 1. IC<sub>50</sub> values of measured coumarin hybrid compounds compared to DOX as reference drug.

Cell line	Ct	1	7a	7b	7c	7d	7e	10	13	DOX
CaCo-2	1.7005019	0.30274004	1.38978276	1.76403734	1.96871657	2.66600438	1.07865591	0.77853734	1.13263098	0.61475703
A549	0.69149983	0.94968866	1.53332834	1.69539443	1.15483508	0.67892756	0.61291219	0.153999	1.47786267	0.76533619
MDA- MB-231	1.59929755	2.52101246	0.37518942	0.6037819	2.21008859	1.35138147	1.32002314	0.84618234	1.07704405	1.95350877
HepG-2	0.76134548	0.36167827	1.05121285	1.04899673	0.84562991	1.67133135	0.26583552	0.55325474	1.37354783	1.65253623

**Table 2.** Selective index values of measured coumarin hybrid compounds.

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(Fig. 2B). Furthermore, 7d was the most potent compound against HepG-2 cell line with IC<sub>50</sub> 82.83  $\mu$ M and SI 1.67 (Fig. 2A).

#### Antimicrobial activities of coumarin hybrids

The antimicrobial potency of coumarin hybrids was tested against different multiple drug resistant strains using microplate assay method. The results indicated that all investigated hybrids showed positive effects against all investigated microbes.

Furthermore, samples **10b** and **10c** considered as the most efficient treatment against *p. aeruginosa* with MIC value of 14.8 and 15.4  $\mu$ M, respectively (Table 3). Also, against both *C. albicans*, *S. aureus* samples **10c** and **13c** were the most potent treatments with MIC values of 0.0011 and 0.0013  $\mu$ M, and 3.2 and 7.2  $\mu$ M, respectively (Fig. 3). Concerning the antimicrobial activity against *E. Coli*, **10b**, **13b** and **13c** were the most potent compounds with MIC values of 9.5, 9.8 and 9.4, respectively (Fig. 4). Finally, **10c**, **13c** and **13b** were the most effective against *Strep. mutans* with a MIC value of 5.2, 14.8 and 28.5  $\mu$ M, respectively (Table 3).

#### Molecular docking studies

The anticancer effect of coumarins is assumed to be attributed to a variety of mechanisms, such as regulation of the estrogen receptor, activation of cell death, blockage of the cell cycle, and inhibition of DNA-associated enzymes including telomerase and topoisomerase (TOP).<sup>36,53,54</sup> A significant portion of fundamental cellular biology involves DNA topoisomerases, which are also molecular targets for several medications, including antibiotics, antibacterials, and anticancer medications. They work by preventing the topoisomerase molecule from relegating DNA strands upon cleavage, so changing it into an agent that damages DNA<sup>55,56</sup>. In the field of medicinal chemistry and drug development, hybrid molecules which are created by combining two or more pharmacophores are a relatively new idea that has gained a lot of attention lately. Therefore, hybridization of coumarin with other anticancer pharmacophores may provide novel bioactive candidates with potential activity as well as low cytotoxicity. The design approach involves the coupling of coumarins with other bioactive molecules as neocryptolepine a quinoline based compound- aiming to afford anticancer compounds that might target topoisomerase.<sup>57-60</sup>.

*In-silico* screening of the quinoline based compounds revealed their promising affinity to Topoisomerase II which is consistent with studies done on similar compounds as acridine and neocryptolipines<sup>61</sup>. The proposed binding mode of the investigated compounds showed marked affinity values ranging from a maximum value of -6.5392 kcal/mol recorded by 7e and a minimum value of -5.5638 kcal/mol recorded by 7c. Docking results disclosed that the planar di or tri ring aromatic system showed *pi*-hydrophobic or  $\pi$ - $\pi$  interactions mostly with the amino acids residues *Asp78, Asp83, Ser79* while, DG1, DC4, and DG5 were the residues for interaction at the DNA-minor groove, (Fig. 5). These interaction are consistent with the reported interactions by the co-crystallized ligand "7-[(**3R**)-**3-aminopyrrolidin-1-yl]-8-chloro-1-cyclopropyl-6-fluoro-4-oxo-1,4-dihydro quinoline-3-carboxylic acid**" in the key residues *Ser79* as well as DG1 in the DNA groove.<sup>44</sup> The synthetic compounds' docking results include the binding affinity Score and Root Mean Square Deviation (RMSD). The following table lists the ligand interactions with the active site residues, including hydrogen bonding and hydro-phobic interactions (Table 4).

#### In-silico ADME assessment

The synthesized compounds were put through an in-silico investigation to determine their pharmacokinetic characteristics using Swiss ADME<sup>62</sup>. The investigated compounds demonstrated high GI absorption ability except for compound **7e**. Moreover, the tested hybrids show low Blood Brain Barrier (BBB) permeability (Table 5). Cell membrane penetrability reflected by the log *P* values, range from 2.77 to 3.91, these values are < 5 which reflects marked cell membranes tolerability<sup>63</sup>. Following "rule of five" for Lipinski, all compounds' molecular weights were less than 500 except for compound **7e** which was slightly higher than 500 recording value of 531.99. In addition, Hydrogen-Bond Donors (HBD) were 2 and Hydrogen-Bond Acceptors (HBA) were 4 for all tested compounds, with rotatable bonds ranging from 4 to 7. The bioavailability score was 0.55 for all compounds except **7e** which recorded the lowest bioavailability score 0.17, this might be attributed to its high molecular weight. All screened compounds had promising ADME data that revealed compliance with the Lipinski's rule except for compound **7e** that had two *Lipinski's* violations with high molecular weight and low bioavailability score which affected its GI absorption probabilities (Table 6).

	Ct		Ct 1		1		7a		7b 7c			7d		7e		10b		10c		13b		13c	
Staph. aureus	62	63	8	8.5	30	31	20	22	29	28	29	30	25	24	3.2	3.2	20	19	7	7.4	65	66	
C. albicans	30	29	10	9.8	62	61	28	27	30	31	62	60	25	26	0.12	0.13	27	27.5	0.0011	0.0014	29	29.5	
E. coli	59	58	20	21.76	29	28	45	43	126	125	46	45	19	16	10	9.7	10	9	7	7.88	10	9	
S. mutans	240	241	17	16.65	248	246	100	103	150	152	130	132	160	162	5	5.5	39	35	15	14.67	29	28	

**Table 3.** MIC values of the synthesized coumarin hybrids against Staphylococcus aureus, Candida albicans,E. coli, Streptococcus mutans and Pseudomonas aeruginosa.

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MIC S. aerous





**Figure 3.** MIC values of the synthesized coumarin hybrid compounds against *Staphylococcus aureus* and *Candida albicans*.

*Structural activity relationship* The main objective of the installation of new hybrids containing coumarin and heteroarenes bearing two, three





MIC S. mutans

В

C MIC P. aeruginosa



Figure 4. MIC values of the synthesized coumarin hybrid compounds against *Pseudomonas aeruginosa Streptococcus mutans* and *E. coli*.

and four fused cyclic rings is to optimize and reach the best activity depending on the synergistic effect of the utilized precursors. Notably, the acridine and neocryptolepine pharmaceutical cores have quinoline motif fused with benzyl as showed in acridine or indole in case of neocryptolepine. Relying on the and from the reported IC<sub>50</sub> values in Table 2, structure activity relationship of the synthesized hybrids was studied. Table 1 emphasized the presence of anticancer activity of these hybrids comparing to their starting core used as positive control and revealed the synergistic effect of these hybrids. In light of the afforded results of the synthesized hybrids against MDA-MBA-231 cancer cell line, it is noteworthy that hybrid 7c containing coumarin and quinoline cores as two fused rings with presence of three carbon spacers of quinolone core showed the best activity with  $IC_{50}$ :50.96  $\mu$ M and SI:2.2 folds among of their relatives presenting in hybrids 7a-e. Furthermore, it is noteworthy that hybrid 7c containing two fused rings with propyl spacer at C-4 of quinoline core had higher activity than tetra- and tri cyclic ring belonging to neocryptolepine and acridine cores with the same spacer attached on C-11 and C-9 resembled to 13c and 10c respectively. On the other hand, hybrid 13c showed antiproliferative activity against HepG-2 cell line compared to their relative hybrids **10c** and **7c**. In addition to, IC<sub>50</sub> of A549 cell line illustrated that 13c have the highest potency than 7c and 10c. Furthermore, hybrid 7d showed the best activity against CaCO-2 cell line which have phenylene diamine as spacer with two fused rings. That illustrated that length of spacer influences the antiproliferative activity. In addition 10b and 13b, which contained two carbons spacer with tri and tetra fused rings, showed higher antibacterial and antifungal activity against S.aureus, s.mutans, E.coli and C.albicans.

#### Conclusions

In the current study, hybrids of coumarin-quinolines **7a–e**, coumarin-acridines **10b,c** and coumarin- neocryptolepines **13b,c** were synthesized and evaluated for their antiproliferative and antimicrobial activities. Hybrids **7c** and **7d** were proved to be the most potent as an antiproliferative agents, showing inhibitory activity against *MDA-MB-231* and *CaCo-2* with IC<sub>50</sub> of 50.96 and 51.95  $\mu$ M respectively. Furthermore, Hybrids **10b** and **13b** showed higher antibacterial activity against *S.aureus*, *S.mutans and E. coli* with MIC from 3.2 to 15  $\mu$ M compared to their corresponding derivatives **7a–e** and **10c** and **13c**. as well as potent inhibitory activity against Candida albicans with MIC 0.0011 to 0.12  $\mu$ M. Molecular docking study disclosed that the planar di or tri ring aromatic system showed hydrophobic interactions at the binding site of action mostly with the amino acids residues *Asp78*, *Asp83*, *Ser79* and DG1, DC4, and DG5 residues at the DNA-minor groove. These interactions are consistent with the reported interactions by the co-crystallized ligand of topoisomerase protein in the key residue *Ser79* and in



**Figure 5.** (A–D) Docking results of the synthesized compounds, best scoring compounds' interactions 7b (in green sticks) and 7e (in cyan sticks) inside the active site of topoisomerase 2 with H-bonds in green, Pi-Pi stacking in pink, pi-H bonds in light pink and pi anion bonds in orange color.

Comp. No	Binding energy Kcal/mol	RMSD (A)	Amino acids involved and type of interactions
7a	- 6.0373	1.9272	Asp78(H-bond), DA5( $\pi$ – $\pi$ )
7b	- 6.4308	1.4711	DG1(H-bond), DG1( <i>π</i> - <i>π</i> ), DC4( <i>π</i> - <i>π</i> ), DA5( <i>π</i> - <i>π</i> )
7c	- 5.5638	2.1466	DG1(π-π)
7d	- 5.8557	1.9477	D Asp78(pi-H)
7e	- 6.5392	1.6699	DG1( $\pi$ - $\pi$ ), DA5( $\pi$ - $\pi$ ), Asp83(H-bond, pi-anion), Ala 84( <i>pi</i> -H)
10	- 5.8972	1.5769	DG1(π-π)
13	- 6.3032	1.2402	DG1(π–π), Asp78( <i>pi</i> -H), Ser79( <i>pi</i> -H)

**Table 4.** Receptor binding affinity, RMSD values and residues involved in the interaction at the receptor active site.

Comp. No.	MW	No. rotatable bonds	No. H-bond acceptors	No. H-bond donors	MR	iLOGP	GI absorption	BBB	Bioavailability score	Lipinski violations
7a	365.77	4	4	2	99.81	2.77	High	Low	0.55	0
7b	393.82	6	4	2	109.42	3.19	High	Low	0.55	0
7c	407.85	7	4	2	114.23	3.01	High	Low	0.55	0
7d	441.87	5	4	2	126.55	3.45	High	Low	0.55	0
7e	531.99	7	4	2	156.01	3.78	Low	Low	0.17	2
10	423.46	7	4	2	126.73	3.18	High	Low	0.55	0
13	476.53	7	4	2	143.49	3.91	High	Low	0.55	0

Table 5. In-silico screening for ADME properties of compounds 7a-d,10 and 13.

	MW	No. rotatable bonds	No. H-bond acceptors	No. H-bond donors	MR	iLOGP	GI absorption	BBB	Bioavailability score	Lipinski violations
7a	365.77	4	4	2	99.81	2.77	High	Low	0.55	0
7b	393.82	6	4	2	109.42	3.19	High	Low	0.55	0
7c	407.85	7	4	2	114.23	3.01	High	Low	0.55	0
7d	441.87	5	4	2	126.55	3.45	High	Low	0.55	0
7e	531.99	7	4	2	156.01	3.78	Low	Low	0.17	2
10c	423.46	7	4	2	126.73	3.18	High	Low	0.55	0
13c	476.53	7	4	2	143.49	3.91	High	Low	0.55	0

Table 6. In silico screening for ADME properties of compounds 7a-d,10c and 13c.

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DG1 of the DNA groove. All screened compounds showed promising ADME data following Lipinski's rule except for compound 7e that had two Lipinski's violations with high molecular weight and low bioavailability score which affected its GI absorption probabilities. Overall, the results of this study support the possibility of using these coumarin hybrids as promising anticancer and antimicrobial agents for further in vivo animal model study.

#### Data availability

All raw data of measurements is available and could be shared when requested, both corresponding authors (I.E.T. and E.A.K.) are fully responsible for providing all data requested.

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#### Author contributions

E.A.: Experiments, gathered data, analyses, wrote/edited original, and reviewed the manuscript. N.E.-D.: Conducted experiments of microbiology and cell culture. M.Z.: Supervision; A.M.H.: Experiments; E.I.E.: Docking study, data analysis, curation, writing and editing. E.A.K. and I.E.S.: Proposed the research idea and concept, devised a research plan, data analysis, interpretation of data, supervision, and writing/editing the original and final manuscript. The final manuscript was written, reviewed, and approved by all authors.

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#### **Competing interests**

The authors declare no competing interests.

#### Additional information

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