

Evaluation of antibiotics as potential antiproliferative agent against triple-negative breast cancer: a cell-based repurposing study

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Received: 2 October 2024

Revised: 31 October 2024

Accepted: 3 November 2024

Published: 27 June 2025

**Journal of The Arab Society for Medical
Research** 2025, 20:26–33

Background/aim

Drug repurposing is a well-known strategy that involves identifying new therapeutic applications for existing medications. Given the global burden of breast cancer, this study aims to investigate the potential of repurposing antibiotics as a potential treatment option for this disease.

Patients and methods

Cytotoxic activity of about 50 antibiotics against triple-negative breast cancer cell line (MDA-MB231) was investigated using a 3-(4, 5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay. The potency of these antibiotics was compared with doxorubicin as positive control while their specificity was assessed using the noncancerous cell line (BJ-1). A three-dimensional model was also used for determining the penetration power of the strongest antibiotics and an acid phosphatase assay was used to evaluate their cytotoxic effects. Moreover, gene expression analysis of P53, Bcl-2, and Bax was performed using real-time PCR.

Results

A screening assay was conducted to evaluate the cytotoxic effects of 50 antibiotics against MDA-MB-231 breast cancer cells. At a concentration of 100 μ M, 19 antibiotics demonstrated significant cytotoxicity reducing cell viability by more than 50% after 48 h and nonsignificant affecting on normal cells. Additionally, two of these antibiotics coded with S44 and S49, exhibited potent anticancer activity with high penetration power, as evidenced by their ability to shrink three-dimensional spheroids of MDA-MB-231 cells with diameters less than untreated cells. The real-time PCR showed a significant upregulation ($P < 0.5$) in the P53 and Bax levels, while Bcl-2 expression was down regulated significantly ($P < 0.5$) by S44 and S49.

Conclusion

The findings suggest that repurposing antibiotics may offer promising therapeutic avenues for triple-negative breast cancer treatment. Further investigations are warranted to elucidate the underlying mechanisms of action and evaluate the clinical potential of these compounds.

Keywords:

antibiotics, breast cancer, human cancer cell lines, repurposing of drugs

J Arab Soc Med Res 20:26–33

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1687-4293

Introduction

Drug repurposing, which involves identifying new uses for existing medications, offers a promising viable strategy in cancer drug discovery and antibiotic development. It potentially accelerates drug development and reduces costs compared with traditional methods of creating new drugs [1]. It has already been found to be safe in preclinical experiments and in humans [2]. Since most preclinical testing, safety evaluations, and, in certain situations, formulation development, will already have been finished, the time required for drug development can be shortened [3]. These advantages of reduced risk, accelerated development, and potential cost savings can lead to a more efficient return on investment in drug repurposing [4].

The normal breast contains, besides fat cells, a network of lobes that are made up of tiny, tube-like structures called lobules that contain milk glands. Tiny ducts connect the glands, lobules, and lobes, and carry milk from the lobes to the nipple. Most breast cancers begin in the cells lining the milk ducts and are called ductal carcinomas [5–7].

Lobular carcinoma is the second most commonly seen form that begins in the lobules. The cancer's location, growth rate, and the probability that it has spread are all

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indicated by the stage. There are generally 5 stages for breast cancer: stage 0 (zero), which is also known as noninvasive cancer or ductal carcinoma in situ (DCIS), and stages I through IV (1 through 4). Every person's breast cancer is distinctive, and the biology and behaviors associated with the disease determine the treatment plan [8].

The initial course of treatment for cancer in its earlier stages is typically surgery to remove the tumor and any surrounding lymph nodes. After surgery, additional treatment is typically administered to reduce the chance of the cancer returning. This treatment may include chemotherapy, radiation therapy, hormone therapy, or targeted therapy. To reduce the growth of the tumor, these treatments may also be administered before surgery [9].

Several noncancer drugs, including antibiotics, have shown potential anticancer properties by targeting common molecular pathways linking diseases. The most successful example of drug repurposing are repurposing of Raloxifene as an anticancer (Breast cancer) while its original indication was Osteoporosis. Similarly, aspirin, which has long been used to treat pain, has shown promise as an anticancer agent, particularly for colorectal cancer. Other notable examples of drug repurposing include sildenafil

(Viagra), originally developed for erectile dysfunction, which has also been found effective for pulmonary hypertension, and thalidomide, which was initially used for leprosy but is now approved for multiple myeloma [10]. Therefore, the aim of the present work is the repurposing of different widely used antibiotics for their anticancer effect against triple-negative of breast cancer cell line MDA-123).

Patient and methods

Sampling and study design

Fifty antibiotics were randomly purchased from commercial drug stores in Cairo, Egypt, as listed in Table 1.

Ethical approval

The present study was conducted with the Code of Ethics of the World Medical Association, according to the principles expressed in the Declaration of Helsinki. This study has been approved by the Ethical Committee of the National Research Centre with approval number 19/289.

Methods

Cell lines

The estrogen-negative MDA-MB-231 human cancer cell line and Skin Fibroblast (BJ-1) normal cells were

Table 1 Antibiotics drugs used under investigation

Type	Drug name	Code	Type	Drug name	Code
Coated tablet	Cephalexin	S26	Ampoule	Ampicillin	S1
Tablet	Amoxicillin and flucloxacillin	S27	Ampoule	Cephadrine	S2
Coated tablet	Levofloxacin	S28	Ampoule	Ceftriaxone	S3
Coated tablet	Cephalexin	S29	Ampoule	Ceftazidime	S4
Ampoule	Ceftriaxone	S30	Ampoule	Benzathinbenzy lpenicilline	S5
Ampoule	Cephadrine	S31	Ampoule	Cefoperazone	S6
Ampoule	Sulbactam and Ampicillin	S32	Ampoule	Cefotaxime sodium	S7
Ampoule	Amoxycillin and Flucloxacillin	S33	Ampoule	Sulbactam/Ampicillin	S8
Capsules	Chloramphenicol	S34	Ampoule	Clindamycin	S9
Coated tablet	Gemifloxacin	S35	Ampoule	Ciprofloxacin	S10
Coated tablet	Clarithromycin	S36	Ampoule	Gentamicin	S11
Coated tablet	Cefadroxil monohydrate	S37	Capsules	Cefdinir	S12
Ampoule	Benzathinebenzylpenicillin	S38	Coated tablet	Zithromycin	S13
Ampoule	Ceftazidime	S39	Tablet	Ofloxacin	S14
Ampoule	Clindamycin	S40	Coated tablet	Spiramycin	S15
Coated tablet	Linezolid	S41	Tablet	Neomycin sulphat	S16
Coated tablet	Moxifloxacin	S42	Coated tablet	Amoxicillin and Clavulanic acid	S17
Coated tablet	Rifaximin	S43	Coated tablet	Moxifloxacin	S18
Capsules	Doxycycline	S44	Capsules	Chloramphenicol	S19
Ampoule	Cefoperazone	S45	Coated tablet	Clarithromycin	S20
Tablet	Neomycin sulphate	S46	Coated tablet	Rifaximin	S21
Ampoule	Cefotaxime Sodium	S47	Capsules	Doxycycline	S22
Bottle	Levofloxacin	S48	Coated tablet	Linezolid	S23
Coated tablet	Nitazoxanide	S49	Coated tablet	Cefadroxil monohydrate	S24
Coated tablet	Tedizolid phosphate	S50	Coated tablet	Gemifloxacin	S25

obtained from the American Type Culture Collection (USA). Cells were cultured in DMEM-12 medium supplemented with fetal bovine serum, penicillin, streptomycin, L-glutamine, and nonessential amino acids. The cultures were maintained at 37°C in a humidified incubator with 5% CO₂ [11].

Cytotoxicity of antibiotics on MDA-MB231 and BJ-1 monolayers

The anticancer activity of the selected drugs was evaluated against MDA-MB231 human breast carcinoma cells and BJ-1 normal foreskin fibroblast cells.

MTT assay [3-(4, 5- dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide]

MDA-MB231 and BJ-1 cells were seeded into 96-well plates at a density of 10×10^3 and 50×10^3 cells per well, respectively. After 24 h, the cells were treated with a final concentration of 100 ppm of the tested drugs in triplicate. The cells were incubated for 48 h, with 1 µM doxorubicin as a positive control and 0.5% DMSO as a negative control. Cytotoxicity was assessed using the 3-(4, 5- dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay as described and was calculated according to the following formula: $[1 - (av(x))/(av(NC))] \times 100$. Where: Av: average, X: absorbance of sample, NC: absorbance of negative control. Absorbance was measured at 595 nm with reference 690 nm [10].

Determination of IC₅₀ values

Drugs demonstrating at least 50% cytotoxicity against breast cancer cells were selected for further analysis. Dose-response studies were conducted at various concentrations and the resulting data was used to calculate IC₅₀ values using probit analysis with SPSS software.

Cytotoxicity bioassay on the estrogen negative MDA-MB-231 multi-cellular spheroids

Three-dimensional (3D) multicellular tumor spheroids (MCTs) are becoming a valuable tool in cancer research due to their ability to better replicate the *in vivo* tumor microenvironment compared with traditional 2D cell cultures. MCTs exhibit growth dynamics and cell-cell interactions that closely resemble those of *in-vivo* tumors, providing a promising model for studying tumor biology and evaluating therapeutic efficacy. Drugs that demonstrated significant cytotoxicity ($\geq 60\%$) against MDA-MB-231 breast cancer cells in 2D culture were further evaluated in 3D MCTs. Selected drugs were

added to triplicate wells of 3D spheroids at a final concentration of 100 µg/ml and incubated for 7 days. Cisplatin and DMSO served as positive and negative controls, respectively. Cytotoxicity was measured using the acid phosphatase assay. Treated spheroids were washed, lysed, and incubated with p-nitrophenyl phosphate. The reaction was terminated, and absorbance was measured at 405 nm. Cytotoxicity was calculated using the formula: % cytotoxicity = $[1 - (\text{average sample absorbance}) / (\text{average negative control absorbance})] \times 100$. Drugs exhibiting significant activity on 3D spheroids at 100 µg/ml underwent dose-response studies at various concentrations (100, 50, 25, and 12.5 µg/ml) [12].

Phase contrast microscopy and image analysis

Images of all spheroids were captured on day 5 (start of treatment) and day 7 (after treatment) using an Olympus inverted microscope equipped with an Olympus SC100 camera (Germany) and cellSens software. The images were used to evaluate spheroid growth.

Quantitative real time PCR for P53, Bcl-2, and Bax

MDA-MB-231 cells were treated with their respective IC₅₀ concentrations of promising antibiotics for 48 h. Gene expression analysis of P53, Bcl-2, and Bax was performed using real-time PCR.

RNA isolation

The assay was carried out according to the methodology described by Oliveira *et al.* [12] In brief, MDA-MB-231 cells were seeded into a six-well plate at a density of 3×10^5 cells per well and incubated for 24 h. The IC₅₀ dose of promising antibiotics was then added to each well, except for the control well. Total RNA was isolated using the Trizol reagent protocol. The cells were collected and centrifuged, followed by RNA isolation using the RNeasy extraction kit and cDNA synthesis using the Transcriba First Strand cDNA Synthesis kit.

Reverse transcription (RT) reaction

RT-PCR analysis was carried out using specific primers for the target genes. Gene expression levels were normalized to beta-actin as a housekeeping gene. The RT reaction involved an initial incubation at 50°C for 45 min to synthesize cDNA, followed by real-time PCR amplification cycles (95°C for 10 s and 60°C for 60 s) on a Rotor-Gene 3000 system. A negative control was included in each run to assess primer specificity and contamination, using primer sequences as in Table 2 [13].

Table 2 Primers for the target genes P53, Bcl-2, and Bax

p53 F	5'-CCCCTCCTGGCCCCTGTCATCTTC-3'
p53 R	5'-GCAGCGCCTCACAACCTCCGTCAT-3'
Bcl-2 F	5'-CCTGTGGATGACTGAGTACC-3'
Bcl-2 R	5'-GAGACA GCC AGG AGA AAT CA-3'
Bax F	5'-GTTTCA TCC AGG ATC GAG CAG-3'
Bax R	5'-CATCTT CTT CCA GAT GGT GA-3'
β -actin F	5'-GTGACATCCACACCCAGAGG-3'
β -actin R	5'-ACAGGATGTCAAACTGCCC-3'

Statistical analysis

Statistical analysis and graph plotting for in-vitro cell line studies were performed using Microsoft Excel and GraphPad Prism 8 software (GraphPad Software, San Diego, California, USA). All data were collected in triplicate and expressed as the mean $\pm$ standard error (SE) from three independent experiments. The IC_{50} was calculated by linear regression analysis using the SPSS program. ANOVA test was used to compare between more than two groups with parametric data. Statistical significant was considered at P value less than 0.05.

Results

Cytotoxicity of antibiotics on MDA-MB231 monolayers

Nineteen of the tested antibiotics exhibited cytotoxicity rates exceeding 50% against MDA-MB231 breast cancer cells at a concentration of 100 μ m after 48 h. The remaining antibiotics demonstrated low cytotoxic activity against this cell line, as shown in Table 3. These

promising antibiotics were introduced to further investigation at different concentrations to calculate their IC_{50} .

Determination of IC_{50} values

The results in Table 4 showed that seven antibiotics only from the nineteen promising antibiotics demonstrated less activity against normal cells and have a significant selectivity index. These antibiotics were further tested in 3D culture systems.

Cytotoxicity of promising antibiotics on 3D (spheroid model) for MDA-MB231 and BJ-1

The results represented in Table 5 summarize the effects of seven promising antibiotics on 3D model of both MDA-MB231 and BJ-1. Only two of the seven antibiotics tested (S44 and S49) demonstrated significant effectiveness against cancer cells while remaining safe for normal cells. Moreover, the graph of the 3D model of the MDA-MB231 cell line showed that only S44 and S49 have diameter around 170 to 190 μ m otherwise the control ranged from 270 to 290 μ m in diameter. Moreover, only S44 and S49 showed cytotoxic effects more than Cisplatin which is the positive control (Fig. 1).

Comet assay

The DNA damage was assessed by performing a comet assay on MDA-MB-231 cell line before and after applying antibiotics coded with S44, S49 and doxorubicin (positive control). The results showed

Table 3 Percentage inhibition of Antibiotic drugs against Human breast cancer cell line (MDA-MB231) at 100 μ m after 48 h

Sample code	⁺ Cytotoxicity at 100 μ m against MDA-MB231 after 48h	Sample code	⁺ Cytotoxicity at 100 μ m against MDA-MB231 after 48 h	Sample code	⁺ Cytotoxicity at 100 μ m against MDA-MB231 after 48 h
S1	25.5 $\pm$ 3.9	S18*	69.9 $\pm$ 5.14	S35*	87.9 $\pm$ 7.86
S2	18.3 $\pm$ 2.1	S19	23.2 $\pm$ 1.82	S36*	59.2 $\pm$ 5.23
S3	8.2 $\pm$ 0.79	S20*	78.6 $\pm$ 6.96	S37	12.2 $\pm$ 0.94
S4	28.1 $\pm$ 2.3	S21	20.5 $\pm$ 1.69	S38*	73.9 $\pm$ 6.43
S5*	90.3 $\pm$ 8.7	S22*	87.4 $\pm$ 8.16	S39	5.8 $\pm$ 0.63
S6	0 $\pm$ 0.0	S23	36.7 $\pm$ 3.79	S40	46.5 $\pm$ 5.62
S7*	92.1 $\pm$ 0.82	S24	12.3 $\pm$ 0.98	S41*	57.1 $\pm$ 4.79
S8	39.3 $\pm$ 4.2	S25*	59.7 $\pm$ 5.88	S42	35.5 $\pm$ 3.11
S9	15 $\pm$ 0.9	S26	26.8 $\pm$ 2.79	S43	41.3 $\pm$ 3.87
S10	0 $\pm$ 0.0	S27*	56.8 $\pm$ 5.43	S44*	92.7 $\pm$ 6.98
S11*	61.2 $\pm$ 5.1	S28*	54.2 $\pm$ 4.89	S45	15.8 $\pm$ 1.21
S12	29.2 $\pm$ 2.5	S29*	72.6 $\pm$ 6.99	S46	10.8 $\pm$ 0.87
S13	20.2 $\pm$ 1.87	S30	8.11 $\pm$ 0.83	S47	6.76 $\pm$ 5.14
S14	25.9 $\pm$ 2.18	S31	5.8 $\pm$ 0.47	S48	13.5 $\pm$ 0.98
S15*	78.9 $\pm$ 4.21	S32	9.72 $\pm$ 0.86	S49*	69.6 $\pm$ 3.99
S16	35.8 $\pm$ 2.99	S33	6.18 $\pm$ 0.57	S50*	54.9 $\pm$ 2.76
S17	21.8 $\pm$ 1.89	S34*	59.9 $\pm$ 4.89		

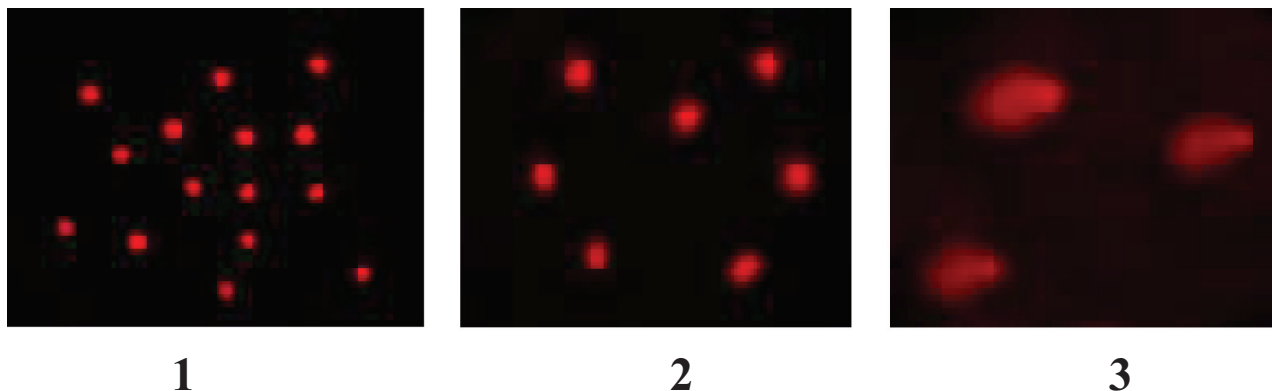
⁺: Data are represented as mean $\pm$ SE (Each sample was done in triplicate). *: Antibiotics within the same column exhibited more potent cytotoxic effects, with inhibition rates exceeding 50%.

Table 4 IC₅₀ of nineteen antibiotics against MDA-MB231 and percentage inhibition of the antibiotics on BJ1 after 48h

Sample code	⁺ IC ₅₀ of antibiotics against MDA-MB231 after 48h	⁺ Cytotoxicity of antibiotics against BJ1 after 48h	Selectivity index
S5*	37.5±2.6	3.2±0.1	2.7
S7*	29.9±1.9	2.3±0.2	3.3
S11	86.1±5.6	6.5±0.6	1.16
S15	70.7±6.7	7.1±0.4	1.4
S18	77.7±4.5	19.2±1.2	1.3
S20	71.0±8.1	31.9±2.6	1.4
S22*	13.7±0.9	12.9±1.5	7.3
S25	87.0±8.1	25.1±2.6	1.14
S27	90.2±6.4	15.9±2.4	1.2
S28	92.1±5.7	11.8±1.8	1.1
S29 *	8.1±0.7	2.4±0.3	12.3
S34	86.9±7.2	13.5±1.2	1.2
S35*	14.3±1.1	12.8±2.1	7
S36	88.1±3.7	30.8±2.8	1.13
S38	61.1±4.2	14.7±0.9	1.63
S41	89.4±6.2	34.2±2.4	1.1
S44*	14.4±0.4	18.8±0.9	6.9
S49 *	11.3±5.3	2.1±0.5	8.8
S50	90.1±8.2	3.7±0.3	1.1

*: Data are represented as mean±SE. The IC₅₀ were calculated by linear regression analysis (SPSS program). *: Data within the same column represent the antibiotics with a greater ability to selectively target cancer cells.

that the negative control, which was MDA-MB-231 cell before applying any drug, had a significant decrease ($P<0.05$) regarding DNA damage. Its value was 10.25 ± 1.31 . Nevertheless, MDA-MB-231 cells after treating with S44 revealed a significant increase ($P<0.05$) with value of 21.50 ± 1.04 and also it exceeded the value of MDA-MB-231 cells after treating with doxorubicin (positive control). Regarding to S49 its effect less than the effect of the positive control but still a significant increase ($P<0.05$), as expressed in Table 6 and shown in Fig. 2.

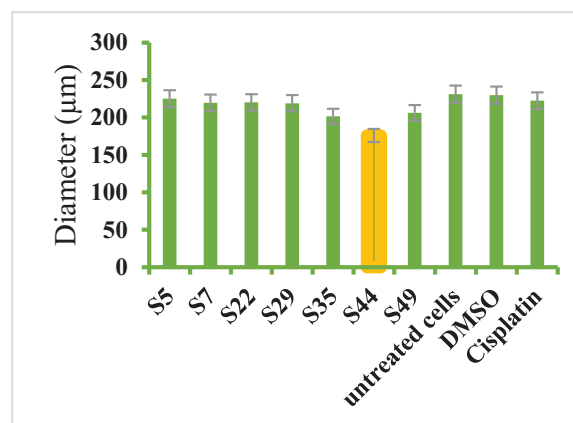
Figure 2

Visual score of normal DNA: 1(class 0), 2 (class 1 and 2) and 3 (class 3) using comet assays.

Table 5 Effect of most promising antibiotic on three-dimensional (spheroid model) of both MDA-MB231 and BJ-1

Sample code	⁺ MDA-MB231% mortality	BJ1% mortality
S5	15.9±1.5 ^a	12.1±0.9 ^a
S7	23.7±1.8 ^a	13.7±1.1 ^a
S22	37.2±2.1 ^a	11.7±0.7 ^a
S29	42.1±1.9 ^a	9.8±0.5 ^a
S35	40.3±2.1 ^a	10.3±0.8 ^a
S44	73.1±3.7 ^b	2.4±0.2 ^b
S49	53.9±3.4 ^b	1.9±0.1 ^b

*: Data are represented as mean±SE. Data with superscript letters (a, b) within the same column are significantly different at P greater than 0.05, using ANOVA test. The 'sample code' Column represents the antibiotic tested, while the '% mortality' rows indicate the observed effects. A higher percentage of mortality indicates a greater effect on the cells.

Figure 1

The diameter in μm of MDA-MB-231 cells spheroids after exposure to the most active antibiotic versus cisplatin.

Evaluation of apoptosis-related genes

Quantitative real time polymerase chain reaction for *P53*, *Bcl-2* and *Bax*

The real-time PCR showed a significant upregulation ($P<0.5$) in the *P53* level in MDA-MB-231 cell line

Table 6 Visual score of DNA damage in breast cancer cell lines treated with S44 and S49

Treatment	Number of cells			Class**				Percentage of DNA damaged cells (Mean±SE)
	No of samples	Analyzed *	Comets	0	1	2	3	
S44	4	400	86	314	30	19	37	21.50±1.04 ^a
S49	4	400	69	331	24	15	30	17.25±0.9 ^b
Negative (–ve)	4	400	41	359	26	5	10	10.25±1.31 ^c
Postive (+ve)	4	400	82	318	24	28	30	20.51±0.65 ^a

*: Number of cells examined per a group, **: Class 0= no tail; 1= tail length and lt; diameter of nucleus; 2= tail length between 1X and 2X the diameter of nucleus; and 3= tail length and gt; 2X the diameter of nucleus. Data with superscript letters (a, b, c) within the same Colum are significantly different at *P* greater than 0.05, using ANOVA test.

Table 7 Effect of S44 and S49 antibiotics in genes expression of P53, Bcl-2, and Bax

Code	Fold Change		
	P53	Bcl-2	Bax
S44	5.3±0.41 ^b	0.4±0.05 ^c	5.2±3.5 ^b
S49	4.4±0.26 ^b	1.54±0.1 ^d	4.7±2.7 ^b
Control –ve	1 ^a	1 ^a	1 ^a
Cisplatin	9.0±2.8 ^c	0.81±0.59 ^b	10.9±4.9 ^c

S44: Doxycycline, S49: Nitazoxanide, A value greater than 1 indicates an up regulation, while a value less than 1 indicates down regulation. Data with superscript letters (a, b, c, d) within the same Colum are significantly different at *P* less than 0.05, using ANOVA test.

treated with antibiotic coded with S44 and S49 in comparison with untreated cells with FLD 5.3 and 4.4, respectively. Regarding to Bcl-2 expression, it was found that a significant down regulation (*P*<0.5) by S44 and S49 while both of them are significantly upregulated Bax expression in MDA-MB-231 cells compared with control group (Table 7).

Discussion

Antibiotics play a pivotal role in treating infectious diseases, significantly reducing mortality rates and enhancing overall quality of life [14]. Beyond their traditional use in treating infections, antibiotics have emerged as potential cancer therapies; however, more research is necessary to fully understand their mechanisms of action and evaluate their clinical efficacy [15–17]. To investigate the potential anticancer effects of these antibiotics, we evaluated 50 compounds against the human adenocarcinoma epithelial breast cell from Mammary gland (MDA-MB-231) breast cancer cell line using different assays.

Among the tested compounds, S44 (Doxycycline) and S49 [Nitazoxanide (NTZ)] demonstrated significant anticancer activity compared with the standard chemotherapeutic agent, cisplatin. The treatment of 3D spheroids with S44 and S49 resulted in a significant reduction in size, suggesting their potential to inhibit cancer cell growth and proliferation [18]. Our

cytotoxicity assessment, utilizing both 2D and 3D cancer models, aligns with the increasing emphasis on 3D models for drug screening [9,10,19]. While 3D models are often favored for their ability to better represent the tumor microenvironment, our findings demonstrate that 2D models can still be valuable for identifying promising anticancer agents [20].

Doxycycline and NTZ, two U.S. Food and Drug Administration (FDA)-approved drugs, have demonstrated promising anticancer activity in breast cancer treatment due to their ability to target cancer cells in various stages and through different mechanisms. Doxycycline has been shown reduce tumor burden in bone metastasis models [21], inhibit cancer stem cell properties and decrease their frequency and ameliorate paclitaxel-induced stem cell enrichment in triple-negative breast cancer [22], and decrease reactive oxygen species levels [22,23]. These mechanisms contribute to its anticancer effects, making it a potential candidate for breast cancer therapy. While a small study found limited benefits for lymphedema, patients reported improved quality of life [24]. Given its low toxicity and potential to target both cancer stem cells and bulk tumor cells, doxycycline may be a valuable addition to breast cancer treatment regimens[23].

NTZ, has demonstrated potent inhibition of c-Myc activation, a key factor in tumorigenesis [25]. Additionally, it targets dormant cancer cells in hypoxic tumor regions by inhibiting mitochondrial respiration [26]. Moreover, S7 (Cefotaxime sodium), S35 (Gemifloxacin), and S44 (Doxycycline) with IC₅₀ values of 20.029, 13.571, and 6.961 μm, respectively, demonstrated exceptional anticancer activity among the tested compounds. These antibiotics demonstrated enhanced safety against normal BJ1 cells while maintaining potent antiproliferative activity against cancer cell lines. This align with recent studies that have explored the potential of combining these antibiotics with existing cancer treatments to offer improved therapeutic outcomes

and enhance the efficacy of biological cancer therapies [27–29].

The present findings demonstrate that promising antibiotics have an antiproliferative impact on MDA-MB-231 cells, which could be the result of the induction of apoptosis, cell cycle arrest, and/or the inhibition of growth. Since the antiproliferative effect of many naturally occurring cancer chemopreventive agents is tightly linked to their ability to induce apoptosis [11], Therefore, this study investigated whether these promising extracts could induce apoptosis in triple-negative breast cancer cells.

At the apoptotic gene expression level, our findings showed that S44 and S49 have the potential to induce apoptosis in breast cancer cells through a combination of p53 up regulation and Bcl-2 down regulation. These findings align with previous research by Son and colleagues which demonstrated that doxycycline, a tetracycline antibiotic, exhibits cytotoxic effects against pancreatic cancer cells through various mechanisms, including the activation of pro-apoptotic genes and the suppression of anti-apoptotic genes [30]. Additionally, recent studies have explored the potential anticancer properties of NTZ and doxycycline. NTZ, has been shown to target the 20S proteasome, inhibiting multiple catalytic subunits and promoting cell cycle arrest and apoptosis in cancer cells [31]. It also interferes with crucial metabolic and pro-death signaling pathways, including drug detoxification, autophagy, and c-Myc inhibition [32].

Conclusion

In conclusion, several antibiotics exhibited significant anticancer activity against breast cancer cells while maintaining a favorable safety profile. These compounds targeted various cancer cell pathways, including apoptosis. This research contributes to the growing body of evidence supporting the potential of repurposing antibiotics for cancer therapy. Future investigations may lead to the development of novel and effective therapeutic strategies.

Acknowledgments

Author contributions: K.M.: Performing the molecular biology techniques, cell culture methodology and manuscript preparation, S.M.E.-H.: Study design, performing cell culture methodology, and final manuscript proofing. A.A.F.S.: Performing the molecular and cell culture methodology. Z.A.E.: statistical analysis and writing the initial draft of the

manuscript. E.M.Y.: Conducting molecular and cell culture experiments, and contributing to manuscript preparation. All authors revised and approved the final manuscript.

Financial support and sponsorship

This work has been supported by National Research Centre, Cairo, Egypt.

Conflicts of interest

There are no conflicts of interest.

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