New Insights into cytotoxic metabolites produced by *Streptomyces griseus* isolate KJ623766: Large scale production, structure elucidation and biological activities

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Abstract

Natural products particularly microbial metabolites have been the mainstay of cancer chemotherapy and are likely to provide many of the lead structures and derivatives with new biological activities. In this research, the production of some potential cytotoxic metabolites from *Streptomyces* (S.) *griseus* isolate KJ623766 was carried out in 14 L laboratory fermenter under specified optimum conditions (28°C temperature, 200 RPM rotation speed, uncontrolled PH, 3 vvm aeration and 2 bar airflow pressure). After 72 hrs of incubation, the cell free culture supernatant (CFCs) was collected and extracted using ethyl acetate (1:1, v/v) at pH 7.0. Using 3-(4,5-dimethylthazol-2-yl)-2,5-diphenyl tetrazolium-bromide (MTT) assay, the cytotoxic activity of the ethyl acetate extract against Caco2 and Hela cancer cell lines was determined with CD50 of 14 µg/ml and 20 µg/ml, respectively. Bioassay guided fractionation of the ethyl acetate extract using different chromatographic techniques had led to the purification of the cytotoxic metabolites coded W1, R1 and R2 with reproducible amounts of 20, 5, and 1.5 mg/l, respectively. The structures of respective metabolites were determined based on the mass, 1D and 2D NMR spectroscopic analysis and were identified as genistein, β-rhodomyacinone and γ-rhodomyacinone, respectively. Accordingly, *S. griseus* isolate KJ623766 can be used as a potential industrial strain for the commercial production of the isoflavonoid genistein, as well as for the production of β- and γ-rhodomyacinone to be used for the construction of new derivatives with more potent cytotoxic activities of the anthracycline family. To the best of our knowledge, this is the first report about the production of the isoflavonoid genistein by *S. griseus* KJ623766.

Introduction

Cancer diseases are a major health problem in both developed and developing countries. Therefore, combating against cancer requires integration between primary, secondary, and tertiary medical care in any nation [1]. In Middle Eastern countries, liver cancer is a major concern among men, especially in certain countries such as Egypt and Saudi Arabia, and to a lesser extent in other countries of this region [2]. While colorectal cancer is the second highest cause of malignancy and death in both sex in the USA [3]. On the other hand, breast cancer is the most frequently diagnosed cancer and the leading causing of cancer death in females, worldwide [4]. Moreover, cervical cancer is the third leading cause of cancer death among females in the less developed countries [5,6].

Natural products particularly microbial metabolites have been the mainstay of cancer chemotherapy and are likely to provide many of the lead structures, that can be used as models for the discovering of new derivatives with improved biological activities [7]. These chemotherapeutic agents isolated from plant, marine and bacterial sources are commonly used in chemotherapy as well as to treat lupus and juvenile rheumatoid arthritis [8]. Various members of *Streptomyces* have made imperative contributions to human with their capabilities to produce various important active metabolites including antimicrobial, antioxidant, anticancer agents [8-10]. Many drugs such as bleomycins, dactinomycin, mitomycin C, daunomycin and doxorubicin originating from Streptomycetes are currently used in the treatment of cancer and most of them were introduced into clinic before discovering their modes of actions [7]. Therefore, this study aimed at production, purification, structural elucidation and determining the biological activities of certain metabolites produced by *S. griseus* KJ623766, a recovered soil isolate previously screened in our lab to have a promising cytotoxic activities against various cancer cell lines [11].

Materials And Methods

Microorganisms

Locally isolated *Streptomyces* strain from Egyptian soil sample, *S. griseus* KJ623766, identified using 16S ribosomal RNA gene sequences (NCBI GenBank access code, KJ623766) [11]. The isolate was preserved onto starch nitrate agar slants slant (soluble starch 10 gm, KNO₃ 2 gm, K₂HPO₄ 1 gm, NaCl 0.5 gm, MgSO₄.7H₂O 0.5 gm, CaCO₃ 3 gm, agar 20 gm, FeSO₄.7H₂O 0.1 gm, MnCl₂ 4H₂O 0.1 gm, ZnSO₄ 7H₂O 0.1 gm per 1liter of distilled water) and sub cultured every month. While slant-medium (50:50) (tryptone 10 gm, yeast extract 5 gm, glycerine 500 ml and distilled H₂O to 1L) was used for long term preservation.

Cell lines

Three cell lines including, kidney epithelial cells derived from the African green monkey (*Vero* cell line, ATCC No.CCL-81), colorectal adenocarcinoma derived from human colon (*Caco-2* cell line) and epithelioid carcinoma derived from human cervix (*HeLa* cell line) were obtained from VACSERA, Egypt. The *Caco-2* and *HeLa* cell lines were used to study the cytotoxic activities of the crude extract and isolated metabolites; stock culture of these cell lines was grown in T-75 tissue culture flasks containing 20 ml of RPMI-1640 medium with 1% antibiotic antimycotic solution and 10% fetal bovine serum. The medium was changed at 48 h intervals and cells dissociated with trypsin solution (0.25% in phosphate buffer saline). *Vero* cell line is a continuous non-tumourigenic when a cell passage was not prolonged [12] and was used to analyze the cytotoxic activities of isolated metabolites. It was propagated in Eagle minimum essential medium (EMEM) with Hank's balanced salt solution (HBSS), supplemented with 10% Fetal bovine serum (FBS) and antibiotics (100 IU penicillin and 100 IU streptomycin/ml) solution, and maintained in EMEM with Earl's balanced salt solution (EBSS) supplemented with 2% FBS and antibiotics solution[12].

Production of cytotoxic agent(s) by *S. griseus* KJ623766

Preparation of seed culture

This was carried out according to Yin et al. [13] with some modifications as follows; surface inoculation of the tested isolate onto starch nitrate agar slant was carried out. After 7 days of incubation at 28°C, the formed spores were scalped from the agar surface and suspended in 3 ml distilled water. An aliquot (200 µl) of the resulting spore bacterial suspension was used to inoculate 10 ml of the soybean meal medium (soybean 15 gm, glucose 15 gm, NaCl 5gm, CaCO₃ 1gm per 1L of distilled water in 100 ml flask) and incubated at 28°C with shaking 200 RPM for 72 hrs in C5KC shaking incubator (New Brunswick scientific, Edison, New jersey, USA). The culture obtained was used to inoculate the main culture.

Fermentation in a laboratory fermenter

This was done according to Radwan et al. [14] with some modifications, where the fermentation process was conducted in 14 L CelliGen 310 bioreactor (New Brunswick Scientific, Edison, NJ, USA), with its Reactor Process Control (RPC) software. Fermentation process was carried out using 5 liters working volume of soybean meal medium under a condition of 28°C incubation temperature; 200 RPM agitation speed, 3 vvm i.e 15 SLPM aeration rate and 2 bar airflow pressure. The dissolved oxygen concentration was adjusted to obtain 100% saturation at the beginning of the run and DO percentage was sensed by the DO probe and monitored during the fermentation process. PH was adjusted at PH 7 at the beginning of the run, remains uncontrolled along the fermentation process (72 h). After inoculation with 3 days age culture of the test isolate (5% v/v) under aseptic conditions, the run started under the previously stated conditions. Fermentation process was left for 72 h during which foam was suppressed using silicon oil. The culture obtained was centrifuged at 6000 RPM for 10 min using EBA20 Centrifuge (Hettich, Germany) and the cell free culture supernatant (CFCS) obtained was collected.

Extraction of cytotoxic metabolite(s)

1 L of CFCS was extracted with ethyl acetate at the level of 1:1 (v/v) in subsequent manner [15,16] and the collected organic layers were evaporated using rotavapour (Heidolph instruments GmbH and Co. Schwabach, Germany) under vacuum at 45°C. After complete evaporation, sample of the fraction residue was redissolved in 1.25% DMSO in tissue culture medium. The cytotoxic activity of the redissolved fraction was evaluated against *Caco2* and *Hela* cancer cell lines using MTT assay. According to the screening program of American National Cancer Institute (NCI), a crude extract is generally considered to have in *vitro* cytotoxic activity if the CD_{50} value is $\leq 30\mu\text{g}/\text{m}$

Isolation of the cytotoxic metabolite(s)

Medium pressure chromatographic separations were carried out with a PuriFlash 4100 system (Interchim; Montluçon, France) consisting of a mixing HPLC quaternary pump, a PDA-UV-Vis detector 190-840nm, a fraction collector, and a sample loading module. For system controlling and process monitoring, Interchim Software 5.0 was used. 360 mg sample was dissolved in 50 mL of methanol then introduced into the column via dry load using 4 gm silica. Elution started with hexane:ethyl acetate (100%:0%) followed by ethyl acetate:methanol (100%:0%) gradient to yield 117 fractions. Elution was monitored using TLC (normal phase silica gel precoated plates F254, Merck, Germany) with UV-detection and spraying with vanillin sulphuric acid spray reagent to yield 12 major fractions (SG1-SG12); these fractions were subjected to MTT assay to measure cytotoxic activity of each against *Caco-2* and *Hela* cell lines. Final purification steps were performed using preparative HPLC (Knauer, Germany) on Kromasil ODS preparative column (10 mm× 250 mm) at flow rates 4 ml/min and UV detection at 254, yielding metabolites W1, R1 and R2. MTT assay was carried out to measure cytotoxic activity of each against *Vero*, *Caco-2* and *Hela* cell lines.

Spectroscopic analysis of the recovered cytotoxic metabolites

Structures of the isolated metabolites were determined based on LCESIMS analysis performed using waters@xevo-tqd® (USA) and NMR spectroscopic analysis which recorded on a Bruker AVANCE HD III 400 MHz spectrometer (Switzerland). NMR samples were dissolved in methanol-d₄ or chloroform-d₃ (Sigma Aldrich, Germany) and transferred to 3 mm NMR tubes (Bruker).

Cytotoxicity assay using MTT method

MTT assay was carried out as described by Saliba et al. [17] with some modifications; 100 μl of the tested metabolite (1mg of the metabolite dissolved in 5% DMSO and tissue culture medium) was added to the well that contain 100 μl of tissue culture medium, followed by two fold serial dilution. A total of 12 dilutions were used for each metabolite to calculate CD_{50} for each. Control wells contained two aliquots of 100 μl of ethyl acetate extract of soybean meal medium (1 mg dissolved in 5% DMSO and tissue culture medium) and 100 μl of tissue culture medium, followed by two fold serial dilutions.

After 24 hrs incubation period at 37 °C in CO₂ incubator, wells were washed with PBS, followed by incubation with 100 μl MTT solution (1 mg/ml) per each well for 1 hr at 37°C. Supernatants were obtained by decantation and the cells were mixed with 100 μl DMSO per each well to dissolve formazan particles. Elutes of the 12 wells of each tested metabolite were obtained and measured spectroscopically at 540 nm using differential wave length of 630 nm in Platos R496 Microplate reader AMD diagnostics, Graz, Austria. Control wells were similarly handled and the percentage cytotoxicity was calculated by the following formula [16].

Cytotoxicity% = 1 - {A₅₄₀ of test culture/A₅₄₀ of control culture} ×100. [16].

Results

Cytotoxic activities of the CFCS ethyl acetate extract

The results showed that the CFCS ethyl acetate extract has potential cytotoxic activities against both cell lines with higher activity against *Caco2* (CD₅₀ 14 µg/ml) than *Hela* cell line (CD₅₀ 20 µg/ml), which indicates the presence of metabolites with higher selectivity to human colorectal adenocarcinoma.

Isolation and purification of cytotoxic metabolite(s)

As determined by TLC with UV-detection and spraying with vanillin sulphuric acid spray reagent, a total of 12 major fractions (SG1-SG12) were obtained. All the fractions were subjected to MTT assay to measure the cytotoxic activity of each. The fraction SG-3 (70% Hexane) showed potential cytotoxic activity against *Caco-2* and *Hela* cell lines with CD₅₀ equals to 9.4 µg/ml and 12.2 µg/ml, respectively. Results of TLC indicate the presence of two major metabolites and one minor metabolite. So further purification of the active fraction SG-3 was carried out using semi-preparative HPLC, by which three pure metabolites namely, W1, R1 and R2 were recovered (Fig. 1).

Cytotoxic activities of the recovered metabolites (W1, R1 and R2)

The three isolated metabolites W1, R1 and R2 showed potential cytotoxic activities against *Caco2* and *Hela* cell lines and lower cytotoxic activities were observed against *Vero* cell line (Table 1). Being cancer cell lines, *Caco2* and *Hela* cell lines have lost their normal regulation of growth or cell death and were more sensitive to the effect of cytotoxic agents than *Vero* cell line.

Table 1 Cytotoxic activities of ethyle acetate extract and isolated metabolites against different cell lines.

Tested metabolite	Cytotoxic activity (CD ₅₀) against different cell lines (µg/ml)		
	<i>Caco2</i>	<i>Hela</i>	<i>Vero</i>
Ethyl acetate extract	14	20	
Fraction SG-3	9.4	12.2	
W1	4.05	5.4	43
R1	6.3	9.45	64.8
R2	7.11	9.35	67.3

Identification and cytotoxic activities of the recovered metabolites

Metabolite W1

Metabolite W1 (20 mg) was isolated as white amorphous powder. ESI-MS revealed pseudomolecular ion peak at m/z 268.99 (M-H)⁻ (Fig. 2) corresponding to molecular formula C₁₅H₁₀O₅. Interpretation of the ¹H NMR spectrum of W1 (Fig. 3) revealed typical pattern of isoflavonoid aglycone derivatives with distinct singlet signal at δH 8.07 (s, 1H) characteristic for H-2 of the isoflavonoids, in addition to the presence of the para-disubstituted aromatic

system with two sets of ortho coupled protons at δ H 7.39 (d, 8.1, 2H) and 6.86 (d, 8.1, 2H) corresponding to H-2', 6' and H-3', 5', respectively. Finally the presence of two Meta coupled protons at δ H 6.35 (d, 2.1, 1H) corresponding to H-6 and δ H 6.23 (d, 2.1, 1H) corresponding to H-8. The APT spectrum of W1 revealed the presence of thirteen carbon signals classified as five methine signals and eight quaternary signals. The 1 H and APT spectra of W1 (Table 2) were identical to those of the isoflavonoid aglycone genistein, thus metabolite W1 was identified as genistein (Supplementary Fig. 1S)

Table 2 NMR data of metabolite W1, Chemical shifts are expressed in δ H values (ppm) from internal TMS. Coupling constants in parentheses are in Hz.

Position	δ _H (DMSO- <i>d</i> 6, 400 MHz, <i>J</i> in Hz)	δ _C (DMSO- <i>d</i> 6, 100 MHz)
2	8.29 (s, 1H)	154.3 (CH)
3		121.7 (C)
4		180.1 (C)
4a		104.7 (C)
5		162.4 (C)
6	12.95 (s, 5-OH)	99.5 (CH)
7	6.21 (<i>d</i> , 2.1, 1H)	165.1 (C)
8	6.36 (<i>d</i> , 2.1, 1H)	94.17 (CH)
8a		158.0 (C)
1'		122.7 (C)
2',6'	7.37 (<i>d</i> , 8.6, 2H)	130.6 (CH)
3',5'	6.82 (<i>d</i> , 8.6, 2H)	115.5 (CH)
4'		157.8 (C)

Metabolite R1

Metabolite R1 (5mg) was isolated as reddish powder, ESI-MS revealed pseudomolecular ion peak at *m/z* 385.05 (M-H)⁻ (Fig. 4) corresponding to molecular formula C₂₀H₁₈O₈. The 1 H NMR of R1 (Fig. 5) revealed typical pattern of anthracycline derivatives, showing three chelating hydroxyl groups at δ H 12.15, 12.90, 13.61 corresponding to 4-OH, 6-OH and 11-OH, respectively. Moreover, the spectrum revealed the presence of three adjacent aromatic protons at δ H 7.93 (dd, 7.4, 1.2, 1H), δ H 7.76 (m, 1H) and δ H 7.37 (dd, 8.4, 1.2, 1H), corresponding to H-1, H-2 and H-3, respectively. The presence of two deshielded methine protons at δ H 5.25 (m, 1H) and δ H 4.91 (brs, 1H) suggested the presence of two tertiary alcohol methine protons at H-7 and H-10 respectively.

This is confirmed by COSY correlations where H-7 showed strong correlations with the methylene protons H-8a and H-8b at δ H 2.19 (m, 1H) and δ H 2.22 (m, 1H). The comparison of 1 H NMR spectrum of R1 (Table 3) with that of known anthracyclins in the literature showed strong similarity to the known metabolite β -rhodomycinone (Supplementary Fig. 2S).

Table 3 NMR data of metabolites R1, R2 and γ - rhodomycinone [18], Chemical shifts are expressed in δ H values (ppm) from internal TMS. Coupling constants in parentheses are given in J in Hz.

Position	δ_H (CDCl ₃ , 400 MHz, <i>J</i> in Hz), Metabolite R1	δ_H (CDCl ₃ , 400 MHz, <i>J</i> in Hz), Metabolite R2	δ_H (CDCl ₃) γ -rhodomycinone
1	7.93 (<i>dd</i> , 7.4, 1.2, 1H)	7.91 (<i>dd</i> , 7.8, 1.0, 1H)	7.91 (<i>dd</i> , 7.8, 1.1, 1H)
2	7.76 (<i>t</i> , 8.0, 1H)	7.73 (<i>t</i> , 8.0, 1H)	7.75 (<i>t</i> , 7.9, 1H)
3	7.37 (<i>dd</i> , 8.4, 1.2, 1H)	7.35 (<i>dd</i> , 8.2, 1.0, 1H)	7.41 (<i>dd</i> , 7.8, 1.1, 1H)
7	5.26 (<i>m</i> , 1H)	7a: 2.98 (<i>m</i> , 1H), 7b: 2.90 (<i>m</i> , 1H)	2.98-2.82 (<i>m</i> , 2H)
8a	2.19 (<i>m</i> , 1H)	2.10 (<i>m</i> , 1H)	2.29-2.14 (<i>m</i> , 2H)
8b	2.22 (<i>m</i> , 1H)	2.35 (<i>m</i> , 1H)	
10	4.91 (<i>s</i> , 1H)	4.82 (<i>s</i> , 1H)	4.81 (<i>s</i> , 1H)
4-OH	12.15 (<i>s</i> , 1H)	12.27 (<i>s</i> , 1H)	12.20 (<i>s</i> , 1H)
6-OH	12.91 (<i>s</i> , 1H)	12.77 (<i>s</i> , 1H)	12.90 (<i>s</i> , 1H)
11-OH	13.61 (<i>s</i> , 1H)	13.87 (<i>s</i> , 1H)	13.60 (<i>s</i> , 1H)
1'a	1.81 (<i>m</i> , 1H)	1.95 (<i>m</i> , 1H)	1.9-1.7 (<i>m</i> , 2H)
1'b	1.92 (<i>m</i> , 1H)	2.00 (<i>m</i> , 1H)	
2'	1.15 (<i>t</i> , 7.5, 3H)	1.13 (<i>t</i> , 7.5, 3H)	1.03 (<i>t</i> , 7.5, 3H)

β - rhodomycinone showed potent cytotoxic activity against *Caco2* and *HeLa* cell lines, the CD₅₀ were 6.3 μ g/ml and 9.45 μ g/ml, respectively. While lower cytotoxic activity was shown against *Vero* cell line (CD₅₀ = 64.8 μ g/ml).

Metabolite R2

The metabolite R2 (1.5 mg), red colored metabolite, showed identical ¹HNMR spectrum to R1 except in the replacement of H-7 at δ_H 5.25 (*m*, 1H) in R1 with the methylene protons δ_H 2.98 (*m*, 1H) and 2.90 (*m*, 1H) corresponding to H-7a and H-7b in R2 (Fig. 6). Comparison of ¹HNMR spectrum of R2 (Table 3) with that of known anthracyclins in the literature showed strong similarity to the known metabolite γ - rhodomycinone (Supplementary Fig. 3S). This was confirmed by ESI-MS which revealed pseudomolecular ion peak at *m/z* 369.09 (*M*-H)⁺ corresponding to molecular formula C₂₀H₁₈O₇ (Fig. 7).

The same as β - rhodomycinone, γ - rhodomycinone showed potent cytotoxic activity against *Caco2* and *HeLa* cell lines, the CD₅₀ were 7.11 μ g/ml and 9.35 μ g/ml respectively. While lower cytotoxic activity was shown against *Vero*

cell line ($CD_{50} = 67.3 \mu\text{g/ml}$).

Discussion

In this study, we aimed at identifying the major active metabolites produced by *S. griseus* KJ623766, a local isolate previously recovered from a soil sample in our lab and showed a promising cytotoxic activities against various cancer cell lines[11]. The production was carried out in 14 L laboratory fermentor under a previously determined optimum conditions [11] in order to recover sufficient pure quantities of the active metabolites as an attempt for structural elucidation. Bioassay guided fractionation using different chromatographic techniques had led to the purification of cytotoxic metabolites coded W1, R1 and R2 in amounts enough to identify the chemical structures of the respective metabolites using various techniques such as mass, 1D and 2D NMR spectroscopic analysis. According, to the obtained results, the structures of respective metabolites were determined and identified as genistein, β -rhodomycinone and γ - rhodomycinone, respectively.

For the metabolite W1, the ^1H MNR, ESI-MS and APT spectra were identical to those of the isoflavonoid aglycone genistein in the literature [18], thus metabolite W1 was unambiguously identified as genistein. However, till now it was not reported in literature that genistein is produced by *Streptomyces* sp. including *S. griseus*. Accordingly, this is the first report of the production of genistein by our studied isolate *S. griseus* KJ623766. Genistein is an isoflavone metabolite found in a number of plants such as [lupin](#), [soybeans](#), [kudzu](#), [fava beans](#) and [psoralea](#) being the principal food source[19,20] as well as in some [medicinal plants](#), including *Flemingia vestita*[21] and *F. macrophylla*[22,23], [coffee](#)[24] and in *Maackia amurensis* cell culture[25]. Genistein showed potent cytotoxic activity against *Caco2* cell line, where CD_{50} against *Caco2*, *Hela* and *Vero* cell lines are $4.05 \mu\text{g/ml}$, $5.4 \mu\text{g/ml}$ and $43 \mu\text{g/ml}$ respectively. These results matched with that reported by Ganai and H. Farooqi [18] and Russo *et al.* [26] which showed the anticancer activity of genistein against human colon cancer. The cytotoxic activity of genistein has been shown to be due to targeting the $\text{PI}_3\text{K}/\text{Akt}$ pathway. Downstream, genistein inhibits EGF induced FOXO3 disassociation from p53 (mut), which further promotes FOXO3 activity and leads to increased expression of the p27kip1 cell cycle inhibitor, which inhibits proliferation in colon cancer cells [27]. The activity of genistein against *Hela* cell line may be due to inhibition of the expression of VEGF and VEGF receptors and Down-regulation of the expression of miR-27a which lead to induction of cell cycle arrest [26]. Genistein was known to have multiple molecular effects, for instance, lipid and glucose metabolism-promoting effects and activities against lipid peroxidation, inflammation, fibrosis, and various tumours [28-30] as well as its antiviral activity against Herpes B Virus [31] and chondroprotective Effects against osteoarthritis [32]

For the metabolite R1, the ^1H NMR analysis showed a typical pattern of anthracycline derivatives and the spectrum revealed the presence of three adjacent aromatic protons corresponding to H-1, H-2 and H-3, respectively. The presence of the two deshielded methine protons at δH 5.25 (m, 1H) and δH 4.91 suggested the presence of two tertiary alcohol methine protons at H-7 and H-10, respectively. In addition, we confirmed its structure by using the COSY correlations where H-7 showed strong correlations with the methylene protons H-8a and H-8b at δH 2.19 (m, 1H) and δH 2.22 (m, 1H) and when we compared the data of ^1H NMR spectrum of R1) with that of known anthracyclins in the literature, we found a strong similarity to the known metabolite β -rhodomycinone [33,34].

For the metabolite R2 (1.5 mg), the ^1H NMR spectrum showed a strong similarity to metabolite R1 except in the replacement of H-7 at δH 5.25 (m, 1H) in R1 with the methylene protons δH 2.98 (m, 1H) and 2.90 (m, 1H) corresponding to H-7a and H-7b in R2. Comparison of ^1H NMR spectrum of R2 with that of known anthracyclins in

the literature showed a strong similarity to the known metabolite γ -rhodomycinone. This also was confirmed by ESI-MS which revealed pseudomolecular ion peak at m/z 369.09 ($M-H$)⁻ corresponding to molecular formula $C_{20}H_{18}O_7$ [33,34].

Both β -rhodomycinone (R1) and γ -rhodomycinone (R2) showed potent cytotoxic activity against *Caco2* and *Hela* cell lines, the CD_{50} were 7.11 μ g/ml and 9.35 μ g/ml respectively. While lower cytotoxic activity was shown against *Vero* cell line (CD_{50} = 67.3 μ g/ml) confirming its applicability to be used as a potential anticancer drug in human.

These results matched with that of Tsuji *et al.* [35] which showed that γ -rhodomycinone, had no differentiation inducing activity, but was cytotoxic however, this cytotoxicity activity against normal cell lines was much lower than cancer cell lines. Supong and his coworkers recently showed that ϵ -rhodomycinone produced by a rare actinomycete *Nonomuraea rhodomycinica* NR4-ASC07 has a potential antimalarial activity [36]. Further studies should be conducted to explore the biosynthetic pathways of the respective metabolites as well as to produce new derivatives of the anthracycline family with higher antitumor activities and lower side effects. The latter can be achieved by structural modifications or microbial transformation of the recovered metabolites.

Conclusions

This study targeting large scale production of cytotoxic metabolites by *S. griseus* isolate KJ623766 on 14L fermentor and studying different recovery methods for downstream processing. The structures of isolated metabolites were determined based on the mass, 1D and 2D NMR spectroscopic analysis and identified as genistein, β -rhodomycinone and γ -rhodomycinone, respectively. The respective metabolites were biologically analyzed using various cancer cell lines for their cytotoxic activities and exhibiting promising cytotoxic activities. Accordingly, *S. griseus* isolate KJ623766 can be used as an industrial strain for the commercial production of the isoflavonoid genistein, as well as the production of β -rhodomycinone and γ -rhodomycinone to be used for production of different derivatives of antitumor antibiotics of the anthracycline family.

Declarations

Ethics approval and consent to participate

not applicable

Consent to publish

not applicable

Availability of data and materials

All data generated or analyzed during this study are included in this published article in the main manuscript and additional supporting file.

Competing interests

The authors declare that they have no competing interests

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Authors' contributions

All authors contributed extensively to the work presented in this paper. ASA and AEA designed, performed the experiments and wrote the first draft of the manuscript. KMA, and MAY designed the experiments, analyzed data and revised the manuscript. NH supervised and approved the manuscript. All authors discussed the results and commented on the manuscript at all stages.

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Figures

Figure 1

HPLC analysis of fraction SG-3

Figure 2

ESI-MS of isolated compound W1

Figure 3

¹H NMR spectra of isolated compound W1

Figure 4

ESI-MS of isolated compound R1.

Figure 6

^1H MNR spectra of isolated compound R2.

Figure 7

ESI-MS of isolated compound R2.

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