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Original Article

SYNTHESIS AND EVALUATION OF A COUMARIN SCHIFF-BASE FOR *IN VITRO* **ANTIBACTERIAL ACTIVITY AGAINST** *STAPHYLOCOCCUS AUREUS*

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ABSTRACT

Objective: 44 novel Schiff bases of aminated 4-methylumbelliferones were designed and subjected to *in silico* evaluation of activity against *S. aureus,* with Dihydrofolate Reductase (DHFR) as the target. The top-scoring compounds (as per binding affinities) were subjected to drug-likeness and ADMET evaluation. Overall assessment of the binding affinities, drug-likeness and ADMET profile (especially toxicity) suggested that the derivative, BVSSS22 was found to be the most promising Schiff base (even when compared to the standard, Trimethoprim). Hence, the objective was to synthesize BVSSS22 and evaluate it for *in vitro* activity against *S. aureus*.

Methods: BVSSS22 was synthesized, characterized *via* melting point, TLC, and spectral data acquisition (ATR-IR, NMR, and HRMS), and evaluated for *in vitro* antibacterial activity against *S. aureus* using the agar-well diffusion method, with Trimethoprim as the standard (n=3).

Results: BVSSS22 was successfully characterized, and the *in vitro* antibacterial assay showed that BVSSS22 possessed zones of inhibition, where at 400 µg/ml, the zone of inhibition was slightly less than that of trimethoprim (18.33±0.57 mm v/s 17.33±1.15 mm).

Conclusion: The results show that BVSSS22 is a potent and safe drug candidate for anti-*S. aureus* action. However, it can be evaluated at a concentration higher than 400 µg/ml or undergo further structural optimization to enhance its *in vitro* potency to surpass that of Trimethoprim.

Keywords: 4-methylumbelliferone, Schiff base, *S. aureus*, Synthesis

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INTRODUCTION

Antibiotics have been a boon to humankind since the release of penicillin to the public in 1942. However, with prescription errors, misuse of antibiotics, and non-adherence to the dosage regimen, bacteria have become more resistant to antibiotics, considerably challenging the once-effective antibacterials' efficacy [1]. Thus, due to Antimicrobial Resistance (AMR), around 7.7 million individuals succumb to bacterial infections every year, and sadly, this number is expected to grow to 10 million annual AMR-related deaths by 2050 if AMR remains unchecked [2, 3].

Coumarins have been found to possess various biological activities, and 4-methylumbelliferone is one of those coumarins [3]. Schiff-base synthesis has been recommended in drug design and development due to its ease of synthesis. Schiff bases of coumarins have been designed and evaluated for various biological activities, including antibacterial activity. However, these carbonyl compounds are more elaborate or complex in structure. Hence, the number of steps required to synthesize the Schiff base becomes unnecessarily large [4, 5]. Till date, there are no Schiff bases synthesized from simple carbonyl compounds and 4-methylumbelliferone and so, this study aims to synthesize these Schiff bases.

We had conducted *in silico* work for 44 designed Schiff base derivatives, taking the target, *Staphylococcus aureus* Dihydrofolate Reductase (DHFR) (PDB ID: 6PBO). Molecular docking, drug-likeness evaluation and ADMET evaluation were already carried out, in which the designed derivative, BVSSS22 (fig. 1), was found to possess a binding score that was more than that of the standard, Trimethoprim and the best ADMET characteristics, especially in terms of absorption and toxicity profile. Hence, this compound was shortlisted for synthesis, characterization and evaluation of *in vitro* biological activity against *S. aureus*.

Fig. 1: Chemical structure of BVSSS22

MATERIALS AND METHODS

Synthesis

Materials

Resorcinol, ethyl acetoacetate (Thermo Fischer Scientific, Bengaluru, India) Silica gel G (Avra Synthesis Pvt. Ltd, India), phydroxyacetophenone, dichloromethane (SDFCL, India), iron powder (BFCLAB, India), concentrated sulphuric acid, n-hexane, concentrated nitric acid, ethyl acetate, concentrated hydrochloric acid (SDFCL, India), Ethanol (Labogens Fine Chem Industry, India), digital melting point apparatus (Inlab, India).

Spectral data acquisition

ATR-IR (Bruker Alpha II ECO-ATR), NMR (Agilent 400MHz NMR, solvent DMSO) MS (Waters Xevo G2-Xs QTof with electrospray ionization as the ionization technique, TOF as the mass analyzer and calibration set to M+1).

Procedure

The scheme followed or the synthesis is given in fig. 2

The steps followed were:

Fig. 2: Synthetic scheme for BVSSS22

Synthesis of 4-methylumbelliferone

98 ml of conc. H2SO⁴ was taken in a round-bottomed flask and cooled to 0-5 °C. In the meantime, a solution of 22 g resorcinol and 27 ml ethyl acetoacetate was prepared and poured into the conc. H2SO⁴ while maintaining the temperature at 0-5 °C. On complete addition, the reaction mixture was stirred at room temperature for 6 h. Then, the mixture was poured gradually into crushed ice while stirring vigorously. The crude product was filtered out, washed with water, and dried. The crude product was then recrystallized with sufficient quantity of 95% ethanol and a pinch of activated charcoal for decolorization [6].

Synthesis of 6-nitro-4-methylumbelliferone

12 g of 4-methylumbelliferone was dissolved in 100 ml of conc. H2SO⁴ in a conical flask, and the mixture was stirred till the complete dissolution of 4-methylumbelliferone while the flask was cooled in an ice bath. Meanwhile, a nitrating mixture was prepared (5 ml conc. HNO3+15 ml conc. H2SO4) and cooled in another ice bath. The nitrating mixture was then added slowly to the reaction mixture, stirring it so that the temperature did not rise above 10 °C. Then, the reaction mixture was stirred for 1 h at room temperature, after which the mixture was poured onto crushed ice under vigorous

stirring. The crude product obtained was filtered off, washed several times with cold water till the filtrate was of neutral pH, and dried. The crude product was then boiled with an appropriate quantity of absolute ethanol and filtered. The residue was 6-nitro-4 methylumbelliferone, which was further purified by boiling with sufficient quantity absolute ethanol [7].

Synthesis of 7-hydroxy-6-{(E)-[1-(4-hydroxyphenyl) ethylidene]amino}-4-methyl-2H-1-benzopyran-2-one [BVSSS22]

A mixture of 1.2 g of 6-nitro-4-methylumbelliferone, 1.0 g phydroxyacetophenone and 0.409 g of iron powder was made in a round-bottomed flask.24 ml of ethanol-water mixture (2:1 v/v) was added to this mixture.0.13 ml of conc. HCl was added, and the mixture was immediately kept for reflux at around 75 °C for about 2 h. Then, the reaction mixture was immediately filtered twice. The filtrate was fractionated twice with 20 ml of dichloromethane, after which the dichloromethane fractions were collected. The dichloromethane extract was dried using 5 g of anhydrous MgSO4, after which the extract was filtered and dried until a solid residue was formed. This residue was purified by a sufficient quantity of dichloromethane. Physical data: Crystalline, yellowish brown, yields 64.88%, MP 158.65 °C, TLC Rf0.56 (ethyl acetate: n-hexane, 6:4) [8].

Fig. 3: ATR-IR spectrum of BVSSS22 (labelled as SB 3 for better identification)

In vitro **antibacterial activity**

The compounds, BVSSS22 and Trimethoprim (control) were evaluated for antimicrobial activity by agar well diffusion method, as described by Champa *et al.*, 2023 with slight modifications. BVSSS22 and Trimethoprim were tested against the bacterial strain, gpositive *S. aureus*. The inoculum was adjusted to approximately 5 × 10⁵ CFU/ml with sterile saline solution. Each compound was dissolved in DMSO, and 10 mg/ml solution was made as a stock. From the stock, different solutions of concentrations ranging from 100 µg/ml to 400 µg/ml were prepared. 20 ml Mueller-Hinton agar medium (HiMedia) was poured into each borosilicate Petri plate and allowed to solidify and the inoculum was then added onto the agar.5 mm wells (4 numbers in a plate; 5 numbers including the blank, DMSO (Dimethylsufoxide) were made with an agar borer and

sample solutions of different concentrations were loaded into the respectively labelled wells. All plates were placed in an incubator at 37 °C for 12h. After incubation, the diameter of inhibition zone (mm) was measured in triplicates and the mean zone of inhibition at each concentration with standard deviation was calculated [9].

RESULTS

Spectral data on the compound BVSSS22

ATR-IR (cm-1) = 3198.19, 3154.24, 2959.78-2853.36, 1740.10, 1666.00, 1535.63, 1518.08; 1H-NMR (DMSO) δ (ppm)= 2.25-2.5 (aromatic CH3, 6H), 6-7.75 (aromatic-H, 6H), near 6.25 (α-pyrone-CH=, 1H), 9.5-10 (phenolic-OH, 2H); HRMS (ES+) (calibrated to M+1): M. W of BVSSS22: 309.32 g/mol; found: 310.107. The spectra are given in fig. 3-5.

Fig. 4:1H-NMR (solvent-DMSO) spectrum of BVSSS22

Fig. 5: HRMS spectrum of BVSSS22

In vitro **antibacterial activity**

The results of the assay are shown in fig. 3 and table 1. Antibacterial studies were performed using the agar diffusion

method for BVSSS22 and the standard Trimethoprim. *Staphylococcus aureus* was the test microorganism used. The compounds were tested at various concentrations (100, 200, 300 and $400 \mu g/ml$.

Fig. 6: *In vitro* **antibacterial activity against** *S. aureus***: BVSSS22 (left) and trimethoprim**

Table 1: In vitro antibacterial activity against S. aureus indicated by zones of inhibition for BVSSS22 and Trimethoprim

Reported as mean ±SD n=3

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DISCUSSION

Both BVSSS22 and Trimethoprim showed *in vitro* activities against *S. aureus* as indicated by their prominent inhibition zones. At 400 µg/ml, a slightly higher zone of inhibition was seen for Trimethoprim than that for BVSSS22 (18.33±0.57 mm vs. 17.33±1.15 mm). This unexpected observed difference could be due to the inadequate number of replicates (n=3). Additionally, BVSSS22 and Trimethoprim could have also been tested at an additional concentration which is higher than 400 µg/ml. The structure of BVSSS22 consists of a 4-methylumbelliferone moiety; A similar study was conducted by Emam S *et al.*, who designed and synthesized some coumarin Schiff-base derivatives and evaluated them for activity against *S. aureus;* they found from the agar well diffusion assay that the compounds '3a, 3b, 3f and 4a' displayed activity with inhibition zones of 25±4, 24±2, 35±2, and 24±3 mm, respectively, which however, were all less than that of the standard, ofloxacin i. e. 40±2 mm [10]. Another similar study is that conducted by Tiwari *et al.*, who designed and synthesized 7 benzamidocoumarins and evaluated them for activity against *S. aureus;* in the two-fold serial microdilution method of evaluation compound '21' possessed activity against *S. aureus,* exhibiting an MIC of 6.25 µg/ml, which was however, more than that of the standard, erythromycin i. e. 2.75 µg/ml [11]. The structures of all the compounds from these studies and BVSSS22 are given in fig. 7.

CONCLUSION

From the prior *in silico* investigation, BVSSS22 was found to possess a good binding score, drug-likeness, and ADMET characteristics. Overall, these *in silico* results were better than those of the standard Trimethoprim. BVSSS22 was successfully synthesized and characterized. In the agar well diffusion assay against *S. aureus*, it was found that Trimethoprim was found to have a slightly greater zone of inhibition than that of BVSSS22. Hence, we can safely conclude that BVSSS22 is a safe and potent drug for antibacterial activity against *S. aureus*. It could be tested at a higher concentration or structurally optimized to enhance its potency further, such as including at least one methoxy group to the structure [12-14].

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AUTHORS CONTRIBUTIONS

Ms. Shreya Shet-For carrying out the synthesis, characterization, *in vitro* activity, and writing Suma B V-. Writing – review and editing, Supervision, Investigation, Conceptualization.

CONFLICT OF INTERESTS

Declared none

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