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In Silico Molecular Docking and Synthetic Elaboration of New Phthalazine Scaffolds Functionalized with a Phthalyl Moiety for Putative Antimicrobial Potency



Aya S. Makhlouf^{1*}, Maher A. El-Hashash², Ashraf A. F. Wasfy¹, Mohamed S. Behalo¹

¹Chemistry Department, Faculty of Science, Benha University, Benha, Egypt ²Chemistry Department, Faculty of Science, Ain Shams University, Cairo, Egypt

Abstract

This study investigated the synthesis and evaluation of novel phthalazine derivatives bearing phthalyl amino acid moieties as potential antibacterial and antifungal agents . A series of phthalazine derivatives (3 a-g) were synthesized by condensing phthalazin -1(2H) – one with various phthalyl amino acids (glutamine , methionine , tyrosine , histidine , alanine , phenylalanine and glycine) using N,N'-dicyclohexylcarbodiimide (DCC) as a dehydrating agent . Subsequent removal of the phthalyl protecting group yielded the corresponding free amino acid derivatives (4 a-g) The bioactivity of the synthesized compounds was scrutinized for antimicrobial efficacy, with certain candidates exhibiting auspicious potential. Computational molecular docking analyses were executed to decipher the intricate binding affinities and interaction dynamics between the synthesized compounds and their respective targets and their putative targets, revealing a correlation between binding affinity and antibacterial activity. The molecular architectures of the synthesized compounds were unequivocally authenticated through advanced spectroscopic modalities, including 'H-NMR, mass spectrometry, and infrared (IR) spectroscopy, alongside comprehensive elemental composition analysis.

Keywords: phthalazinone, phthalyl amino acid, Anti-bacterial, ati-fungal, molecular docking

1. Introduction

Heterocyclic compounds constitute a broad class of cyclic organic molecules that incorporate at least one heteroatom typically nitrogen, oxygen, or sulfurwithin their ring structure. These frameworks play a fundamental role in medicinal chemistry due to their widespread occurrence in biologically significant molecules such as vitamins, enzymes, and natural products. They exhibit a diverse range of biological activities, including antifungal, anti-inflammatory, antibacterial, antioxidant, anticonvulsant, antiallergic, enzyme inhibitory, herbicidal, anti-HIV, antidiabetic, anticancer, and insecticidal effects [1,2]. Among these, nitrogen-containing fused heterocyclesparticularly diaza-heterocycleshave garnered considerable attention as core pharmacophores in drug development. Notable examples include benzodiazine derivatives such as cinnoline (1,2-), quinazoline (1,3-), quinoxaline (1,4-), and phthalazine (2,3-), which feature two nitrogen atoms embedded within a fused aromatic ring system. These structures are commonly found in natural bioactive compounds and synthetic therapeutic agents, and they display a wide spectrum of pharmacological properties [3-6]. Phthalazine (benzopyridazine), with the molecular formula C₈H₆N₂, is considered a privileged scaffold in drug design. Its derivatives have shown a broad range of biological activities, including cardiotonic [7-9], anticonvulsant [10-15], vasorelaxant [16,17], anti-inflammatory [18,19], anticancer [20–23], antibacterial [24–29], antifungal [30–34], antihypertensive [35–38], and carbonic anhydrase inhibitory effects [39– 41], as well as applications in agrochemistry [42-45]. Owing to its versatile biological potential and synthetic flexibility, the phthalazine scaffold continues to be a focal point in medicinal chemistry. In this study, a series of 4-(2,4dimethylphenyl)phthalazine-1(2H)-one derivatives were synthesized and evaluated for their antibacterial and antifungal activities. The findings were further supported by molecular docking studies to explore their potential mechanisms of action.

2. Material and method Materials

All melting points were ascertained without correction utilizing an electrothermal melting point apparatus. Elemental composition assessments were executed via a Heraeus CHN Rapid Analyzer within the Microanalytical Unit at Cairo University. Thin-layer chromatography (TLC) analyses were conducted on Merck TLC aluminum plates coated with silica gel 60F₂₅₄, with visualization facilitated through UV quenching at 254 nm. Infrared (IR) spectral data were acquired using a Unicam SP-1200 spectrophotometer, employing the KBr pelletization technique. ¹H-NMR spectra were recorded in DMSO-d₆ using a Varian Plus spectrometer (300 MHz). Mass spectrometric profiles were generated on a Shimadzu GC-MS QP 1000 EX system, operating at 70 eV under electron ionization (EI) conditions.

Synthesis

4-(2,4 di methyl phenyl)phthalazin-(1H)- one (1)

Hydrazine hydrate (0.01 mol) was added to a stirred solution of 2-(2,4-dimethylbenzoyl)benzoic acid (A) (0.01 mol) in absolute ethanol (50 mL). The reaction mixture was then heated under reflux for 3 hours. After completion of the reaction, the mixture was allowed to cool to room temperature, resulting in the formation of a solid precipitate. The solid was collected by filtration, washed with cold ethanol, and recrystallized from ethanol to afford pure phthalazinone (1).

The obtained phthalazinone (1) was isolated as colorless crystalline solids in 60% yield; m.p. 230–232 °C.IR (cm⁻¹) v_{max} : 3320 (NH) ,3161and 2908(CH) ,1671(CO). H NMR (300 MHz, DMSO-d₆) δ : 2.31–2.40 (s, 6H, 2 × CH₃, m-xylene), 7.10–8.10 (m, 7H, Ar–H), 12.40 (s, 2H, NH₂, exchangeable with D₂O). ¹³C NMR(DMSO-d₆, 300MHz) δ ppm: 159.4(C=O), 138.8, 136.9, 134.7, 132.2, 131.1,131, 130.4, 130.8, 129, 127.5, 126.2, 123.7 (aromatic carbons), 21.6, 19.5 (aliphatic carbons). Anal. Calcd. for C₁₆H₁₄N₂O: C, 76.78; H, 5.64.; N, 11.19. Found C,76.2; H,5.5; N,11.02%.

General procedure for the synthesis of N-phthaloyl amino acid(2a-g)

A mixture of 0.01mol of amino acid (a-g) and 0.01mol of finally ground phthalic anhydride was placed in beaker, then the beaker was then immersed in an oil bath preheated to 180-185 °C for 15 minutes. The mixture was stirred occasionally during the first 10 minutes. After the reaction, the beaker was removed and allowed to cool until the liquid mass solidified. The solid material was recrystallized from methanol-water.

General procedure for synthesis of N-phthalyl amino acid-phthalazinone (3a-g)

A mixture of 4-(2,4-di methyl phenyl) phthalazin-1H-one 1 (0.01mole) and phthalyl amino acid 2a-g (0.01mole) was dissolved in 30mlTHF. The conglomerate of reactants was allowed to cool down to a cryogenic state of 0°C and N,N'-dicyclohexylcarbodiimideDCC(0.01mole) dissolved in THF and added to reaction mixture. The cohort of reactants was subject to stirring for a full 24-hour period at 0°C, transitioned to a further 24 hours of agitation at room temperature, and ultimately maintained at 0°C throughout the overnight interval. The precipitated N,N'-dicyclohexylurea was separated via filtration, and the resulting filtrate underwent successive washing with 1N hydrochloric acid, followed by 1N sodium bicarbonate, and subsequently with a solution that has reached the threshold concentration of sodium chloride as a solute, beyond which additional solute will no longer dissolveof 1N sodium chloride. This mixture was then dried using anhydrous sodium sulfate and allowed to stand overnight. Following this, it was subjected to filtration to eliminate any remaining solids, after which excess solvent was evaporated. The resultant residue was dissolved in benzene and set aside for a duration of two hours, leading to the re-precipitation of dicyclohexylurea, which was then removed by filtration. The solvent was subsequently eliminated under reduced pressure, and the resultant residue was crystallized from an appropriate solvent to yield compound (3 a-g) [47].

4-(2,4-dimethylphenyl) phthalazin-1-yl-2-acetate derived from 1,3-dioxoisoindolin-2-yl (3a).

Yield ,61%, m.p $181-183^{\circ}$ C, $IR(cm^{-1})v_{max}$: 2910(CH),1735,1719 (imidic CO), 1672 (C=O),1630(C=N).. ¹H NMR (300 MHz, DMSO-d₆) δ : 2.30–2.40 (s, 6H, 2 × CH₃, m-xylene), 4.80 (s, 2H, CH₂, COCH₂N), 7.10–8.40 (m, 11H, Ar–H). ¹³C NMR(DMSO-d₆,300MHz) δ ppm:171(C-O-C),168.2(2C=O), 168(C=O), 150, 138.3, 136.8, 133. 5,132.6,132.2, 131.4, 127.5,127.3,126.7,126.6,125.5,123.7,119.7(aromaticcarbon),43,1,21.6,19.5(aliphatic carbons) Anal.Calcd for C₂₆H₁₉N₃O₄:C,71.39;H,4.38;N,9.61%. Found C,71.19; H,4.18; N,9.41%.

5-Amino-2-(1,3-dioxoisoindolin-2-yl)-5-oxo pentanoate of 4-(2,4-dimethylphenyl)phthalazin-1-yl. (3b).

Yield ,63%, m.p 163-165°C, IR(cm⁻¹) v_{max} :2916(CH),1730,1719 (imidic CO), 1667 (C=O),1660(C=N). ¹H NMR (300 MHz, DMSO-d₆) δ : 2.05–2.10 (m, 4H, CH₂CH₂, COCHCH₂CH₂), 2.30–2.40 (s, 6H,2 × CH₃, m-xylene), 4.30 (t, J = 6.6 Hz, 1H, CH,COCHN),7.10–8.30(m,11H,Ar–H),7.10(brs,2H,NH₂,exchangeablewithD₂O). Anal.Calcd.for C₂₉H₂₄N₄O₅:C,68.5;H,4.7;N,11.02%.Found C,68.3; H,4.5; N,10.82%.

3-(4-Hydroxyphenyl)-2-(1,3-dioxoisoindolin-2-yl) propanoate of 4-(2,4-dimethylphenyl)phthalazin-1-yl (3c).

Yield ,81%, m.p 180-182°C, IR(cm⁻¹) v_{max} : 3325(OH),1771,1730(imidic CO), 1714(C=O),1668(C=N) .¹H NMR (300 MHz, DMSO-d₀) δ: 2.30–2.40 (s, 6H,2 × CH₃, m-xylene), 3.10 (d, J = 6.8 Hz, 2H, CH₂–Ph), 4.80 (t, J = 6.8 Hz, 1H, CH, COCHN), 6.90–8.40(m,15H,Ar–H),9.08(s,1H,OH, exchangeable with D₂O).Anal. Calcd. for C₃₃H₂₅N₃O₅:C,72.9;H,4.64;N,7.7%.Found C,72.6;H,4.31;N,7.4%.

2-(1,3-Dioxoisoindolin-2-yl) propanoate of 4-(2,4-dimethylphenyl) phthalazin-1-yl (3d).

Yield ,52%, m.p 180-182°C, IR(cm⁻¹) v_{max} :1725,1718(imidic CO), 1673 (C=O ester),1610(C=N). H NMR (300 MHz, DMSO-d₆) δ :1.50 (d,J = 6.6 Hz, 3H, CH₃,HCCH₃), 2.30–2.40 (s, 6H,2×CH₃, m-xylene), 5.30 (q,J = 6.6 Hz, 1H, CH, COCHN),7.10–8.40 (m,11H,Ar–H). NMR (DMSO-d₆, 300MHz) δ ppm:171(C-O-C), 168.3(C=O), 167.9(2C=O), 150, 138.3, 136.8, 133. 5,132.6,132.2, 131.4, 127.5,127.3, 126.7,126.6, 125.5,123.7, 119.7 (aromatic carbons),55.6, 43,1, 21.6, 19.5,13.2 (aliphatic carbons). Anal. Calcd. for C₂₇H₂₁N₃O₄:C,71.8; H,4.65; N,9.3%. Found C,71.6; H,4.45; N,9.1%.

3-Phenyl-2-(1,3-dioxoisoindolin-2-yl) propanoate of 4-(2,4-dimethylphenyl) phthalazin-1-yl (3e).

Yield ,75%, m.p 182-184°C, IR(cm⁻¹) v_{max} :1720,1714(imidic CO), 1667 (C=O ester),1610(C=N). ¹H NMR (300 MHz, DMSO-d₆) δ: 2.30–2.40 (s, 6H, 2 × CH₃, m-xylene), 3.50 (d, J = 6.8 Hz, 2H, CH₂–Ph), 4.80 (t, J = 6.8 Hz, 1H, CH, COCHN), 7.10–7.90 (m, 16H, Ar–H). Anal.Calcd.for C₃₃H₂₅N₃O₄:C,75.14; H,4.74; N,7.96%. Found C,74.94; H,4.54; N,7.76%.

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3-(1H-Imidazol-4-yl)-2-(1,3-dioxoisoindolin-2-yl) propanoate of 4-(2,4-dimethylphenyl) phthalazin-1-yl(3f).

Yield: 72%; m.p. 190-192°C. IR (cm⁻¹) v_{max} : 1735, 1709 (imidic C=O), 1672 (C=O ester), 1627 (C=N). ¹H-NMR (DMSO-d₆, 300 MHz): δ 2.3–2.4 (2s, 6H, 2 × CH₃, m-xylene), 3.3 (d, J = 7.2 Hz, 2H, CHCH₂), 4.8 (t, J = 7.8 Hz, 1H, COCHN), 7.1–8.3 (m, 11H, aromatic protons), 7.8 and 8.3 (2s, 2H, two CH groups from the imidazole ring), 12.7 (s, 1H, NH, exchangeable with D₂O). Analytical data calculated for C₃₀H₂₃N₅O₄: C, 69.62; H, 4.48; N, 13.53%. Found: C, 69.4; H, 4.32; N, 13.2%.

4-(2,4- di methylphenyl) phthalazin-1-yl -2-(1,3-dioxoisoindolin-2-yl) -4-(methyl thio)butanoate(3g)

Yield: 77%; Melting Point: 170-172°C. IR (cm⁻¹) v_{max} : 1730, 1716 (imidic C=O), 1669 (C=O ester), 1580 (C=N). H-NMR (DMSO-d₆, 400 MHz): δ 2.1 (m, 3H, overlapping signals from SCH₃ and CHCH₂CH₂), 2.3–2.4 (3s, 6H, 2 × CH₃, m-xylene), 2.6 (t, J = 7.6 Hz, 2H, CH₂CH₂S), 4.3 (t, J = 8.1 Hz, 1H, COCHN), 7.1–7.9 (m, 11H, aromatic protons). Anal. Calcd . for C₂₉H₂₅N₃O₄S: C, 68.08; H, 4.93; S, 6.26; N, 8.21%. Found values: C, 67.88; H, 4.73; S, 6.16; N, 8.01%

General procedure for the synthesis of amino acid-phthalazinone (4a-g).

To a solution of compound 3a-g (0.01 moles) in an appropriate alcoholic solvent, hydrazine hydrate (0.01 moles) was meticulously introduced. The resultant reaction mixture was subjected to reflux conditions for a span of three hours. Following the completion of the reaction, the mixture was allowed to cool, leading to the precipitation of a solid product. This solid was then collected by filtration and underwent recrystallization from ethanol, resulting in the formation of the desired compounds (4a-g).

4-(2,4- di methyl phenyl) phthalazin-1-yl-glycinate(4a)

Yield ,80%, m.p 238-240°C, IR(cm⁻¹) ν_{max} : 3436,3300,3161(NH₂ group),1671(C=O ester),1609(C=N). ¹H-NMR (DMSO-d₆, 400 MHz): δ 2.3–2.4 (2s, 6H, 2 × CH₃ of m-xylene), 4.1 (s, 2H, COCH₂NH₂), 7.1–7.8 (m, 7H, Ar-H), 8.3 (s, 1H, NH, exchangeable with D₂O). ¹³C NMR(DMSO-d₆, 300MHz) δ ppm: 171(C-O-C), 168(C=O), 150, 138.3, 136.8, 133.5, 131.4, 127.5,127.3, 126.7,126.6 119.7 (aromatic carbons),45.5, 21.6, 19.5 (aliphatic carbons). Anal. Calcd .for C₁₈H₁₇N₃O₂:C,70.34;H,5.58;N,13.67%. Found C,67.68;H,5.38;N,13.47%.

4-(2,4- di methyl phenyl) phthalazin-1-yl-glutaminate(4b).

Yield,83% , m.p. 218-220 °C, IR (cm⁻¹) v_{max} : 3437, 3303 (NH₂), 1672 (C=O, ester), 1610 (C=N). ¹H-NMR (DMSO-d₆, 300 MHz): δ 2.05–2.1 (m, 4H, COCHCH₂CH₂, J = 7.8 Hz, with interference), 2.3–2.4 (2s, 6H, 2 × CH₃ of m-xylene), 3.4 (t, J = 7.5 Hz, 1H, COCHNH₂), 7.1 (s, 2H, CONH₂, exchangeable with D₂O), 8.4 (s, 2H, NH₂, exchangeable with D₂O), 7.1–7.9 (m, 7H, aromatic protons, Ar-H). ¹³C NMR(DMSO-d₆, 300MHz) δ ppm: 173.6(C=O), 171(C-O-C), 168.3(C=O), 150, 138.3, 136.8, 133. 5, 131.4, 127.5,127.3, 126.7,126.5,119.7 (aromatic carbons),52.2, 33, 28, 21, 19.1 (aliphatic carbons). Anal. Calcd. For C₂₁H₂₂N₄O₃: C,66.65; H.5.86; N.14.81%. Found C,66.45; H.5.66; N.14.61%.

4-(2,4-di methyl phenyl) phthalazin-1-vl-tyrosinate(4c).

Yield ,76%, m.p 238-240°C, IR(cm⁻¹) $_{max}$.3431broad band for(OH group), 3163(NH₂ group),1669(C=O ester). ¹H-NMR (DMSO-d₆, 300 MHz): δ 2.3–2.4 (2s, 6H, 2 × CH₃ of m-xylene), 3.4 (d, J = 7.2 Hz, 2H, CH₂-Ph), 4.2 (t, J = 7.6 Hz, 1H, COCHNH₂), 6.6–7.9 (m, 11H, aromatic protons, Ar–H), 8.9 (s, 2H, NH₂, exchangeable with D₂O), 9.1 (s, 1H, OH, exchangeable with D₂O). Anal. Calcd. for C₂₅H₂₃N₃O₃:C,72.62;H,5.61;N,10.16%. Found C,72.42.45;H,5.41;N,9.96%

.4-(2,4-di methyl phenyl)phthalazin-1-yl alaninate(4d)

Yield ,89%, m.p 2 39- 2 41 $^\circ$ C, IR(cm $^-$ 1)v_{max}: 3326,3160(NH₂ group),1670(C=O ester). 1 H-NMR (DMSO-d₆, 300 MHz): δ 1.2 (d, J = 6.8 Hz, 3H, CHCH₃), 2.3–2.4 (s, 6H, 2 × CH₃ from m-xylene), 3.5 (q, J = 6.8 Hz, 1H, COCHNH₂), 7.1–7.9 (m, 7H, aromatic protons, Ar–H), 8.7 (s, 2H, NH₂, exchangeable with D₂O). Anal. Calcd .for C₁₉H₁₉N₃O₂:C,71.01;H,5.96;N,13.08%. Found C,70.81;H,5.61;N,12.8%.

4-(2,4-di methyl phenyl) phthalazin-1-yl phenylalaninate(4e).

Yield ,71%, m.p 200°C, IR(cm⁻¹) v_{max} ;3327,3162(NH₂ group),1670(C=O ester),1628(CN).¹H-NMR (DMSO-d₆, 300 MHz): δ 2.3–2.4 (2s, 6H, 2 × CH₃ of m-xylene), 3.2 (d, J = 7.2 Hz, 2H, CH₂-Ph), 4.1 (t, J = 7.6 Hz, 1H, COCHNH₂), 7.1–7.9 (m, 12H, Ar–H), 8.7 (s, 2H, NH₂, exchangeable with D₂O).Anal. Calcd. for C₂₅H₂₃N₃O₂:C,75.55; H 5. 83;N,10.57%. Found C,75.32;H,5.61;N,10.2%.

4-(2,4-di methyl phenyl) phthalazin-1-yl histidinate(4f).

Yield ,78%, m.p 220°C, IR(cm⁻¹) \mathbf{v}_{max} : 3327,3162(NH₂ group),1671(C=O ester),1626(CN). 'H-NMR (DMSO-d₆, 300 MHz): δ 2.3–2.4 (2s, 6H, 2 × CH₃ from m-xylene), 3.2 (d, J = 7.2 Hz, 2H, CHCH₂), 4.1 (t, J = 7.6 Hz, 1H, COCHNH₂), 7.1–7.9 (m, 7H, Ar–H), 7.6 and 8.6 (2s, 2H, imidazole CH), 8.7 (s, 2H, NH₂, exchangeable with D₂O), 12.7 (s, 1H, imidazole NH, exchangeable with D₂O). Anal. Calcd. For C₂₂H₂₁N₅O₂: C,68.20; H,5.46;N,18.08%. Found C,68.1;H,5.11;N,17.9%.

4-(2,4-di methyl phenyl) phthalazin-1-yl methioninate(4g).

Yield ,50%, m.p 210°C, IR(cm⁻¹) v_{max} :3437,3326(NH₂ group),1671(C=O ester),1630(CN). ¹H-NMR (DMSO-d₆, 300 MHz): δ 2.1 (m, 3H, overlapping signals from SCH₃ and CHCH₂CH₂), 2.3–2.4 (2s, 6H, 2 × CH₃ of m-xylene), 2.6 (t, J = 7.4 Hz, 2H, CH₂SCH₃), 3.4 (t, J = 7.2 Hz, 1H, COCHNH₂), 7.1–7.9 (m, 7H, Ar–H), 8.7 (s, 2H, NH₂, exchangeable with D₂O). Anal. Calcd. For C₂₁H₂₃N₃O₂S:C,66.12; H,6.08;S,8.39;N,11.01%. Found C,66.01.32;H,6.01;S,8.25;N,10.9%.

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Antimicrobial assay

The antimicrobial activity of the synthesized compounds was evaluated using the agar diffusion method. All compounds were screened *in vitro* for their antibacterial activity against **Staphylococcus aureus** (Gram-positive bacteria), and **Klebsiella pneumoniae** (Gram-negative bacteria) using nutrient agar as the culture medium. The antifungal activity was assessed against **Candida albicans** and **Aspergillus niger** using Sabouraud dextrose agar medium. **Ampicillin** and **gentamicin** served as standard reference drugs for Gram-positive and Gram-negative bacteria, respectively, while **nystatin** was used as the standard antifungal agent. **Dimethyl sulfoxide** (**DMSO**) was employed as a negative control. All compounds were tested at a final concentration of **15 mg/mL** against both bacterial and fungal strains.

Method of Testing[48]

Sterilized nutrient agar medium (20–25 mL) was aseptically poured into sterile Petri dishes and allowed to solidify at room temperature. A microbial suspension was prepared in sterile normal saline to match the turbidity of a 0.5 McFarland standard, equivalent to approximately 1.5×10^8 CFU/mL. The turbidity was adjusted to an optical density (OD) of 0.13 using a spectrophotometer at 625 nm. Within 15 minutes of turbidity adjustment, a sterile cotton swab was immersed in the microbial suspension and used to evenly inoculate the surface of the solidified agar. The plates were left to dry for 15 minutes with the lids in place to ensure proper adherence of the inoculum. Subsequently, wells of 6 mm diameter were aseptically punched into the agar using a sterile cork borer. Each well was filled with 100 μ L of the test compound solution using a micropipette. The plates were then incubated at 37 °C for 24 hours to assess antibacterial activity. The experiment was performed in triplicate, and the resulting zones of inhibition were measured in millimeters (mm) using a standard ruler.

Molecular Docking Investigations

Molecular docking is a sophisticated computational method utilized to anticipate the interaction between a protein (receptor) and a small molecule (ligand) within the protein's binding site. This information is crucial in drug discovery and design.

3. Results and Discussion

The synthesis of new phthalazinone derivatives was successfully achieved through a Friedel-Crafts acylation reaction. The reaction between phthalic anhydride and m-xylene in the presence of AlCl₃ as a catalyst led to the formation of 2-(2,4-dimethylbenzoyl) benzoic acid (compound A), as depicted in Scheme 1. The structure of compound A was confirmed through infrared (IR) spectroscopy, which displayed a characteristic carbonyl stretch (ν C=O) at 1673 cm⁻¹, confirming the presence of the acylated moiety. Subsequently, compound A was subjected to a nucleophilic cyclization reaction using hydrazinium hydroxide in ethanol, resulting in the formation of 4-(2,4-dimethylphenyl)phthalazine-1(2H)-one (1)[24,49,50,51] , as illustrated in Scheme (1).

Infrared (IR) spectroscopic analysis of compound (1) demonstrated the complete disappearance of the characteristic hydroxyl group absorption band, accompanied by the emergence of a strong carbonyl stretching band at 1671 cm⁻¹, consistent with the presence of a cyclic amide functionality. These findings confirm the successful formation of the phthalazinone core structure. Compound (1) was subsequently employed as a key synthetic intermediate for the generation of a structurally diverse array of amino acid-conjugated derivatives. These derivatives were synthesized through the coupling of (1) with a series of *N*-protected amino acids. The rationale behind this structural diversification lies in the well-documented enhancement of biological activity imparted by the introduction of amino acid residues to heterocyclic cores such as phthalazinones.

To investigate this effect further, compound (1) was reacted with *phthalyl*-protected derivatives of amino acids including glutamine, alanine, phenylalanine, tyrosine, methionine, glycine, and histidine. The coupling reactions were carried out in the presence of N,N'-dicyclohexylcarbodiimide (DCC), a widely used dehydrating agent, to facilitate amide bond formation. This procedure afforded a series of new phthalazinone derivatives, designated as compounds (3a–g), as illustrated in Scheme 2.

phthalic anhydride
$$Amino \ acids$$
 $-H_2O$
oil bath (fusion)

Ar $Ar = m$ -xylene scheme(2)

The N-deprotection of compounds 3a-g was accomplished via thermal reflux in anhydrous ethanol. Each compound was subjected to continuous reflux under boiling conditions for a duration of 3 hours, ensuring complete removal of the phthalyl protecting group from the amino moiety. This deprotection strategy efficiently afforded the corresponding free aminoacyl derivatives (4a-g) in good yields. The transformation is depicted in **Scheme 3**, highlighting the successful conversion of protected intermediates to their respective unprotected analogues.

The success of the deprotection process was unambiguously confirmed by infrared spectroscopic analysis. The spectra of compounds **4a–g** exhibited the appearance of a characteristic **N–H stretching vibration**, indicative of a free amino group, typically observed in the region of **3300–3500 cm**⁻¹. Concurrently, the **disappearance of the imide-related absorption band** confirmed the complete removal of the *N*-phthalyl protecting group. These spectral changes collectively provide definitive evidence for the successful generation of the deprotected aminoacyl-phthalazinone derivatives.

Statistical analysis :-

Statistical differences among samples of the same bacterial or fungal type were assessed using one-way analysis of variance (ANOVA), followed by Duncan's multiple range test for post hoc comparisons, utilizing the SPSS software package (version 22.0, IBM Corp., Armonk, NY, USA). Data are presented as mean \pm standard deviation ($\bar{X} \pm SD$). A p-value of less than 0.05 was considered statistically significant, while p-values less than 0.01 and 0.001 were regarded as highly significant and very highly significant, respectively. The observed inhibition zones are systematically presented in Tables 1 and 2

Table (1) delineates the antibacterial and antifungal bioactivity profiles of the newly synthesized compounds (3a-g), providing a comprehensive evaluation of their inhibitory efficacies.

Sample	3a	3b	3c	3d	3e	3f	3g	Standard antibiotic
Microorganism								antiblotic
Gram negative bacteria								Gentamicin
Klebsiella pneumonia (ATCC: 10031)	11±1.0 +Ve	NA	15.3±0.6 ++Ve	14.7±0.6 ++Ve	22.0±1.0 ++++Ve	9.7±0.6 +Ve	12.3±0.6 ++Ve	25.3±0.6ª
Gram positive bacteria								Ampicillin
Staphylococcus aureus (ATCC: 13565)	NA	NA	NA	NA	NA	NA	NA	21.3±0.6ª
Fungi								Nystatin
Candida albicans (ATCC: 10231)	NA	NA	NA	NA	NA	NA	NA	21.6±0.6
Asperagillus Nigar (ATCC: 16404)	NA	NA	NA	NA	NA	NA	NA	19.3±0.6

Table (2) delineates the antibacterial and antifungal efficacies of the newly synthesized compounds (4a-g), offering an intricate comparative analysis of their inhibitory potentials against targeted microbial strains.

Sample microorganism	4a	4b	4c	4d	4e	4f	4g	Standard antibiotic
Gram negative bacteria								Gentamicin
Klebsiella pneumonia (ATCC: 10031)	28.3±0.6 ++++Ve	14 ±1.0 ++Ve	20±1.0 + ++Ve	25±0.6 ++++Ve	19.0±0.6 +++Ve	11.7±0.6 +Ve	13±1.0 ++Ve	25.3±0.6 ^a
Gram positive bacteria								Ampicillin
Staphylococcus aureus (ATCC: 13565)	16.7±0.6	NA	NA	20.3±0.6	NA	NA	NA	21.3±0.6ª
Fungi								Nystatin
Candida albicans (ATCC: 10231)	NA	NA	NA	NA	NA	NA	NA	21.6±0.6
AsperagillusNigar (ATCC: 16404)	NA	NA	NA	NA	NA	NA	NA	19.3±0.6

- Zone of inhibition is expressed in the form of mean± Standard deviation (mm).
- NA: Noactivity.
- Well diameter (6mm).
- 100 μ was tested.

Molecular docking Studies

Binding with hydroxylethylthiazolekinase

To check the binding ability of compounds with hydroxylethylthiazolekinase, five compounds (4a,4d, 3e,TZE,ampicilin) were docked separately in the active pocket of this enzyme. Strong binding with such enzyme suggests the chance to be used as bacteriocidal and hence the mode of action for their activity as antibiotic agent.

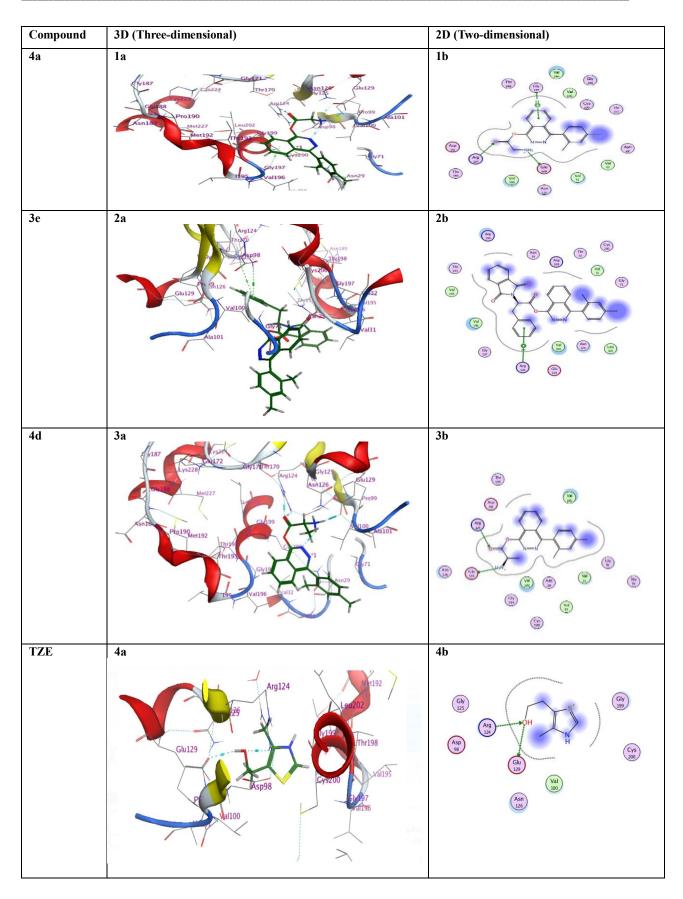
By screening the data depicted in the Table 3 it can be noticed that compound 4a showed binding affinity to hydroxylethylthiazole kinase without forming any Arene-cation, only with one Arene-H with Gly-197 and H-bond with Glu 129 and Arg 124 as show in (fig 1a,1b). We also noticed that compound 3eshowed the strongest binding profile (B.E=6.049Kcal/mole) to hydroxylethylthiazole kinase without forming any Hydrogen – bond and Arene-H, only one Arene –cation with Arg 129 as show in (fig 2a,2b). Compound 3e had marvelous results as antibiotic compound compared to Ampiciline which has binding energy (B.E=-5.1532Kcal/mole) as show in fig2a,2b, fig5a,5b.

On the other hand compounds 4d,Tzeligand and Ampicline showed binding affinity to hydroxyethylthiazole kinase with H-bond with Glu129, Arg 124, without forming any Arene-H and Arene –cation as show in (fig3a,3b,fig4a,4b, fig5a,5b).Tze ligand showed the worth binding mode (B.E= -3.621932Kcal/mole) although it could form interaction with H-bond with (Glu129 and Arg 124) .This may be attributed to it's small rigid structure and minor flexible functionality.Compund4d has a similar mood withTzeligand, but 4d has more binding energy value(-5.2 Kcal/mole) than binding energy of Tze ligand (-3.6219 Kcal/mole). So,we conclude that 3e has the strongest effect as Anti-biotic.

Table (3). binding data of phthalazine derivative hydroxylethylthiazole kinase B.E= Binding energy , RMSD= root mean square deviation

Ligand	B.E	RMSD	H-Bond	Arene-H	Arenecation
4a	-5.2343	1.7699	Glu 129	Gly 197	-
			Arg 124	·	
4d	-5.2025	1.4853	Glu129	-	-
			Arg124		
3e	-6.0497	1.8565	-	-	Arg124
Tzeligand	-3.6219	1.4534	Glu129	-	-
Ü			Arg124		
Ampicillin	-5.1533	1.8978	Gly199	-	-
•			Arg124		

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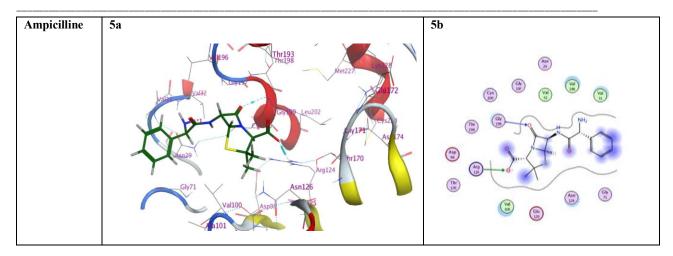


Figure (1a,2a,3a,4a,5a) 2D-(1b,2b,3b,4b,5b) 3D- binding profiles of hydroxylethylthiazole kinase with (3e,4a,4d,ampiciline,tze) respectively showing H-bonds, arene-H and arene- cation .

Conclusion

The present study describes the efficient synthesis of a series of innovative phthalazine-based amino acid conjugates. Their noteworthy antimicrobial activity, supported by molecular docking analyses, underscores their potential as promising lead compounds for the design and development of new antimicrobial therapeutics.

Conflict of interest

The authors unequivocally affirm that no conflicting or divergent interests exist that could be construed as compromising the integrity of this manuscript's publication.

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