One-Pot Three Component Synthesis, Biological Evaluation and Molecular Docking Studies of Novel Indolyl-3,4-dihydroacridinones

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Abstract: Simple and efficient one-pot method has been developed for the synthesis of indolyl-3,4dihydroacridinone analogues (**4a-4p**) in good yields using 2,5-Disubstituted indole-3-carboxaldehyde, 5,5dimethylcyclohexane-1,3-dione and 4-substitued anilines. The structures of all newly synthesized compounds have been characterized based on IR, ¹H, ¹³C NMR, Mass spectral data and elemental analysis. All the compounds have shown very good in vitro antimicrobial and antioxidant activities. Further, in silico molecular docking study indicates that the synthesized compounds are novel inhibitors of the enzyme Glucosamine-6-phosphate synthase.

Keywords: Indole-3-carboxaldehyde, Dihydroacridinone, Antimicrobial Activity, Antioxidant Activity, Molecular Docking.

INTRODUCTION

Over a period of past few decades microbial infections have been evolved on terrifying levels across the world as a result of antimicrobial resistance¹⁻². However, literature survey reveals that the slow rate of novel antimicrobials discovery coupled with rapid multidrug resistance signifies an urgent need to develop new and efficient antimicrobial agents with less side effects³. This framework involves the design of novel motifs with different mode of action than that of contemporary medicines. In search of a new potential target for antimicrobial therapy, the enzyme Glucosamine-6-phosphate synthase has been found to involve in constituting cell wall⁴⁻⁵. The product of this enzyme D-glucosamine-6-phosphate (GlcN-6-P), which is a precursor for the synthesis of essential macromolecules containing chitin in fungi and peptidoglycan in bacteria, as well as glycoproteins in mammals ⁶. Inhibition of GlcN-6-P synthase leads to disruption of microbial cell wall and subsequently, cell lysis without damaging mammalian cell.

Indole and its derivatives have emerged as important skeletons of various biologically active compounds such as antimicrobials⁷, antioxidants⁸, anti-carcinogens⁹ and antivirals¹⁰. Moreover, indole nucleus has fascinated increased attention due to its various potential bioactivities, like anti-rheumatoidal¹¹, anti-inflammatory¹², and anti-HIV¹³ activities and also play a vital role in tubulin polymerization inhibitory action¹⁴. On the other hand, acridine and acridinone derived scaffolds have acquired therapeutic success due to various interesting biological properties such as antioxidant¹⁵, antimalarial¹⁶, antihelmintic¹⁷, and antitumor¹⁸. In addition many acridines have reported as DNA intercalators, topoisomerase inhibitors¹⁹ and drugs in the treatment of Alzheimer's disease²⁰.

Incited by these studies and our research interest to launch the programs of design and synthesis of bioactive privileged structures containing indole ring²¹⁻²⁴, we have devised an efficient and economically viable protocol for the synthesis of indole integrated acridinones with a view to generate promising biologically active analogues. All compounds were evaluated for their *in vitro* antimicrobial and antioxidant activities. Molecular docking studies were performed on aforesaid GlcN-6-P synthase in order to investigate the better understanding of protein-drug binding affinity.

MATERIALS AND METHODS

All chemicals and reagents were bought commercially from Himedia, Spectrochem and SD Fine chemical companies and used without further purification. Open capillaries were used to determine the melting points of synthesized compounds and are uncorrected. The completion of reaction was examined by thin-layer chromatography (TLC) using pre-coated aluminium sheets with silica gel 60 F_{254} (MERCK). Mixture of ethyl acetate and hexane (1:1) was used as mobile phase and visualised under UV light and iodine. Infrared spectra (in KBr-pellet) were recorded using Perkin Elmer - Spectrum RX-IFTIR (ν_{max} in cm⁻¹). ¹H and ¹³C NMR spectra recorded on BRUKER Avance II NMR Spectrometer. The LC-MSD-Trap-SL instrument was used to record Mass spectra. The elemental analysis was performed using FLASH EA 1112 SERIES instrument. All the compounds furnished C, H and N values within ±0.5% of the theoretical values.

General synthesis of 2,5-substituted lindole-3 carboxaldehydes (1a-1d)

The 2,5-disubstituted indoles were fomylated under Vilsmeier-Haack conditions to afford precursors 2,5disubstituted indole-3-carboxaldehydes (**1a-1d**) which are used to obtain the target compounds²⁵.

General synthesis of 3,3-dimethyl-9-(2,5-disubstituted-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4a-4p)

A mixture of 2,5-disubstituted-indole-3-carbaxaldehyde (**1a-1d**) (0.01mol) 5,5-dimethylcyclohexane-1,3-dione (**2**) (0.01mol) and 4-substitued anilines (**3a-3d**) (0.01mol) in 12ml of ethanol was refluxed for 5-7hrs with catalytic amount of acetic acid. The progress of the reaction is monitored by TLC. After completion of the reaction, the excess of solvent was removed under reduced pressure. Crude product obtained after cooling to room temperature was filtered and then recrystallized from suitable solvent to get the target compounds (**4a-4p**).



[Reaction Scheme]

Antimicrobial Activities

Antimicrobial activities of the synthesized compounds were evaluated according to broth dilution assay method²⁶. The medium containing various concentrations of synthesized compounds *viz.*, 1000 μ g – 62.5 μ g per ml in DMSO prepared by serial dilution. After inoculation of culture, the tubes were incubated for 72 hours at 28°C. The minimum inhibitory concentration (MIC) of each sample was determined by measuring the optical density in the spectrophotometer (Electronics India) at 520 nm and compared the result with those of the non-inoculated broth used as blank. Control was prepared using media and inoculum without synthetic compound.

DPPH Radical Scavenging activity

The synthesized compounds were screened for free radical scavenging activity by DPPH method²⁷. 1 mL of different concentrations (25 μ g/mL, 50 μ g/mL 75 μ g/mL and 100 μ g/mL) of each compound as well as reference

standards Ascorbic Acid (AA) and Butylated Hydroxy Anisole (BHA) in methanol were taken in different test tubes. 1 mL methanolic solution of DPPH (0.1 mM) was added to these test tubes. The volume was adjusted to 5 mL by adding 3 mL of methanol and shaken vigorously. After incubation for 30 minutes at 27 °C, the absorbance was measured at 517 nm. The control was prepared as above without any compound. The percentage of radical scavenging activity was calculated using the following formula:

% radical scavenging activity = [(Control OD – Sample OD) / (Control OD)] \times 100

Ferric reducing antioxidant power activity

Compounds were evaluated for ferric reducing antioxidant power activity by using Ascorbic Acid (AA) and Butylated Hydroxy Anisole (BHA) as standards. Different concentrations of compounds along with standards (25 μ g/mL, 50 μ g/mL 75 μ g/mL and 100 μ g/mL) were mixed with 2.5 mL of 1% potassium ferricyanide in 2.5 mL of sodium phosphate buffer (0.2 M, pH 6.6). The mixture was incubated at 50 °C for 20 minutes and then 2.5 mL of trichloroacetic acid (10%, w/v) was added. Next, the contents were centrifuged at 3000 rpm for 10 minutes and supernatant layer (5 mL) mixed with 5 mL of distilled water followed by 1 mL of 0.1% ferric chloride solution. Absorbance was measured at 700 nm against the blank.

Molecular Docking Study

The ligands 2D structures were drawn using ChemDraw Ultra 12.0 and converted to energy minimized 3D structures with proper orientations using Marvin view (Chem Axon) program. The crystal structure of receptor enzyme D-glucosamine-6-phosphate synthase (PDB ID: 2VF5) was obtained from the RCSB Protein Data Bank, http://www.rcsb.org/pdb in pdb file format. To make the receptor enzyme free from any ligand, all heteroatoms and water molecules were removed while the Kollman charges, polar hydrogen atoms and solvation parameters added. All ligand's single bonds were assigned as free to revolve. AutoDock 4.2 and AutoDock Tools 1.5.6 docking programs were used to perform the docking simulation²⁸. The docking grid box was set at 70, 64, and 56 Å for x, y and z respectively with a spacing of 0.375 Å which cantered at active site and surrounded by the residues Ala 602, Val 399, Ala 400, Gly 301, Thr 302, Ser 303, Cys 300, Gln 348, Ser 349, Thr 352, Ser 347 and Lys 603. AutoGrid 4 program was used in order to set the grid maps and other parameters were fixed to default standard values. The AutoDock 4 and Lamarckian genetic algorithm (LGA) was used for energy calculation with a maximum of ten runs for each tested

ligand. The outputs were visualized by means of the Accelrys Discovery Studio client package and inspected for the binding modes with possible hydrogen bonding, polar and hydrophobic interactions of ligands with receptor active site. All these AutoDock simulation were performed in Intel (R) Core(TM) *i*3 CPU M380 @ 2.53 GHz of Dell system origin, with 2 GB DDR3 RAM. AutoDock 4.2 was compiled and run under Windows 7 Ultimate operating system.

RESULTS AND DISCUSSION

Synthesis of novel dihydroacridinone analogues of 2,5-disubstituted indoles (**4a-4p**) via one-pot three component methodology as outlined in Reaction Scheme. The progress of the reaction was monitored by TLC. The excess of solvent was removed under reduced pressure and reaction mixture was cooled to room temperature. The resulting crude product obtained was filtered, washed and recrystallized from suitable solvent. The structures of the synthesized compounds were confirmed by their melting points, IR, ¹H, ¹³C NMR, Mass and analytical data. This environmentally benign protocol presents mild reaction condition, simple procedure, with good yields and purity.

The IR spectrum of 7-chloro-9-(5-chloro-2-phenyl-1H-indol-3-yl)-3,3-dimethyl-3,4-dihydroacridin-1(2H)-one (**4a**) showed a strong absorption at 3443 cm⁻¹ corresponding to indole NH, absorptions at 1663 cm⁻¹ and 1593 cm⁻¹ correspond to carbonyl and aromatic C=N stretching respectively. The ¹H NMR spectrum of (**4a**) has exhibited a singlet at δ 12.39 due to indole NH which is also D₂O exchangeable. A singlet at δ 8.21, doublet at δ 8.10, and multiplet between 7.96-7.11 for eleven aromatic protons in the molecule. Signals between 2.43-2.36 and at δ 1.01 were assigned for two methylene and methyl protons respectively. The ¹³C NMR of the compound (**4a**) has shown a signal at 193 is due to carbonyl carbon and at δ 164 corresponds to C=N. The two methylene carbons appeared at δ 55 and 51 whereas two equivalent methyl carbon displayed a peak at δ 31. The Mass spectrum of (**4a**) has shown molecular ion peak at m/z = 484 corresponds to its molecular weight, with isotopic peaks at m/z = 486 and 488. These spectral data support the structure of compound (**4a**).

7-Chloro-9-(5-chloro-2-phenyl-1H-indol-3-yl)-3,3-dimethyl-3,4-dihydroacridin-1(2H)-one (4a)

Yield 69%, mp 194-195 °C. IR spectrum, v, cm⁻¹: 3443(NH), 1663(C=O), 1593(C=N). ¹H NMR spectrum, δ , ppm: 12.39 (s, 1H, indole NH), 8.21 (s, 1H, Ar-H), 8.10 (d, J = 9.0 Hz, 1H, Ar-H), 7.96 (dd, J = 9.0 and 2.0 Hz, 1H, Ar-H), 7.75-7.72 (m, 2H, Ar-H), 7.54-7.47 (m, 3H, Ar-H), 7.39 (s, 1H, Ar-H), 7.34 (d, J = 8.4 Hz, 1H, Ar-H), 7.11 (dd, J = 8.4 and 1.6 Hz, 1H, Ar-H), 2.43 (s, 2H, CH₂), 2.36 (s, 2H, CH₂), 1.01 (s, 6H, 2CH₃). ¹³C NMR spectrum, δ , ppm:

193, 164, 146, 144, 138, 136, 137, 134, 132, 130, 125, 121, 55, 51, 33, 31. Found, %: C 71.69; H 4.52; N 5.80. Calculated, %: C 71.76; H 4.57; N 5.77. MS, *m/z*: 484 [M]⁺, 486 [M+2]⁺, 488 [M+4]⁺ (10:6.3:1).

7-Chloro-3,3-dimethyl-9-(5-methyl-2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4b)

Yield 60%, mp 196-198 °C. IR spectrum, *v*, cm⁻¹: 3432(NH), 1652(C=O), 1601(C=N). ¹H NMR spectrum, *δ*, ppm: 12.42 (s, 1H, indole NH), 8.13 (d, *J* = 8.8 Hz, 1H, Ar-H), 7.99 (s, 1H, Ar-H), 7.92 (dd, *J* = 8.7 and 2.0 Hz, 1H, Ar-H), 7.71-7.68 (m, 2H, Ar-H), 7.58-7.49 (m, 3H, Ar-H), 7.41 (s, 1H, Ar-H), 7.31 (d, *J* = 8.3 Hz, 1H, Ar-H), 7.10 (dd, *J* = 8.3 and 1.6 Hz, 1H, Ar-H), 2.40 (s, 2H, CH₂), 2.33 (s, 2H, CH₂), 2.29 (s, 3H, CH₃), 1.00 (s, 6H, 2CH₃). Found, %: C 77.51, H 5.38, N 6.05. Calculated, %: C 77.49, H 5.42, N 6.02. MS, *m/z*: 465 [M+1]⁺, 467 [M+3]⁺ (3:1).

7-Chloro-3,3-dimethyl-9-(2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4c)

Yield 64%, mp 190-192 °C. IR spectrum, *ν*, cm⁻¹: 3448(NH), 1658(C=O), 1599(C=N). ¹H NMR spectrum, *δ*, ppm: 12.39 (s, 1H, indole NH), 8.10 (d, *J* = 9.0 Hz, 1H, Ar-H), 7.98 (dd, *J* = 9.0 and 2.2 Hz, 1H, Ar-H), 7.85 (d, *J* = 8.6 Hz, 1H, Ar-H) 7.78-7.62 (m, 5H, Ar-H), 7. 51 (d, *J* = 8.6 Hz, 1H, Ar-H), 7.42 (s, 1H, Ar-H) 7.28-7.19 (m, 2H, Ar-H), 2.43 (s, 2H, CH₂), 2.31 (s, 2H, CH₂). 0.98 (s, 6H, 2CH₃). Found, %: C 77.21, H 5.10, N 6.18. Calculated, %: C 77.24, H 5.14, N 6.21. MS, *m/z*: 451 [M+1]⁺, 453 [M+3]⁺ (3:1).

7-Chloro-9-(1H-indol-3-yl)-3,3-dimethyl-3,4-dihydroacridin-1(2H)-one (4d)

Yield 59%, mp 210-213 °C. IR spectrum, *ν*, cm⁻¹: 3258(NH), 1654(C=O), 1592(C=N). ¹H NMR spectrum, *δ*, ppm: 12.20 (s, 1H, indole NH), 8.29 (s, 1H), 8.06 (d, *J* = 8.7 Hz, 1H, Ar-H), 7.94 (dd, *J* = 8.7 and 1.8 Hz, 1H, Ar-H), 7.87-7.84 (m, 1H, Ar-H), 7.59-7.56 (m, 1H, Ar-H), 7.46 (s, 1H), 7.30-7.27 (m, 2H, Ar-H), 2.43 (s, 2H, CH₂), 2.36 (s, 2H, CH₂), 1.01 (s, 6H, 2CH₃). Found, %: C 73.65, H 5.16, N 7.50. Calculated, %: C 73.69, H 5.11, N 7.47. MS, *m/z*: 374 [M]⁺ 376 [M+2]⁺ (3:1).

7-Bromo-9-(5-chloro-2-phenyl-1H-indol-3-yl)-3,3-dimethyl-3,4-dihydroacridin-1(2H)-one (4e)

Yield 70%, mp 198-199 °C. IR spectrum, *v*, cm⁻¹: 3413(NH), 1649(C=O), 1610(C=N). ¹H NMR spectrum, δ , ppm: 12.41 (s, 1H, indole NH), 8.21 (s, 1H, Ar-H), 8.07 (d, *J* = 9.2 Hz, 1H, Ar-H), 7.92 (dd, *J* = 9.2 and 2.0 Hz, 1H, Ar-H), 7.78-7.74 (m, 2H, Ar-H), 7.60-7.55 (m, 3H, Ar-H), 7.42 (s, 1H, Ar-H), 7.36 (d, *J* = 8.4 Hz, 1H, Ar-H), 7.02 (dd, *J* = 8.4 and 1.6 Hz, 1H, Ar-H), 2.42 (s, 2H, CH₂), 2.35 (s, 2H, CH₂), 1.00 (s, 6H, 2CH₃). ¹³C NMR spectrum, δ , ppm:

196, 167, 143, 141, 140, 139, 138, 136, 131, 130, 126, 125, 54, 51, 32, 30. Found, %: C 65.75, H 4.17, N 5.30. Calculated, %: C 65.74, H 4.19, N 5.29. MS, *m/z*: 528 [M]⁺, 530 [M+2]⁺, 532 [M+4] (10:13:3).

7-Bromo-3,3-dimethyl-9-(5-methyl-2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4f)

Yield 65%, mp 202-203 °C, IR spectrum, *ν*, cm⁻¹: 3376(NH), 1652,(C=O), 1586(C=N). ¹H NMR spectrum, *δ*, ppm: 12.38 (s, 1H, indole NH), 8.15 (d, *J* = 8.8 Hz, 1H, Ar-H), 8.00 (s, 1H, Ar-H), 7.91 (dd, *J* = 8.7 and 2.0 Hz, 1H, Ar-H), 7.69-7.65 (m, 2H, Ar-H), 7.50-7.41 (m, 4H, Ar-H), 7.28 (d, *J* = 8.3 Hz, 1H, Ar-H), 7.09 (dd, *J* = 8.3 and 1.6 Hz, 1H, Ar-H), 2.41 (s, 2H, CH₂), 2.34 (s, 2H, CH₂), 2.30 (s, 3H, CH₃), 1.00 (s, 6H, 2CH₃). Found, %: C 70.70, H 4.97, N 5.55. Calculated, %: C 70.73, H 4.95, N 5.50. MS, *m/z*: 508 [M]⁺, 510 [M+2]⁺ (1:1).

7-Bromo-3,3-dimethyl-9-(2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4g)

Yield 67%, mp 204-206 °C, IR spectrum, ν, cm⁻¹: 3365(NH), 1670(C=O), 1594(C=N). ¹H NMR Spectrum, *δ*, ppm: 12.37 (s, 1H, indole NH), 8.02 (d, *J* = 8.8 Hz, 1H, Ar-H), 7.93 (dd, *J* = 8.8 and 2.0 Hz, 1H, Ar-H), 7.86 (d, *J* = 8.2 Hz, 1H, Ar-H) 7.76-7.61 (m, 5H, Ar-H), 7.49 (d, *J* = 8.2 Hz, 1H, Ar-H), 7.40 (s, 1H, Ar-H) 7.29-7.23 (m, 2H, Ar-H), 2.43 (s, 2H, CH₂), 2.31 (s, 2H, CH₂). 0.98 (s, 6H, 2CH₃). Found, %: C 70.36, H 4.65, N 5.68. Calculated, %: C 70.31, H 4.68, N 5.65. MS, *m/z*: 494 [M]⁺, 496 [M+2]⁺ (1:1).

7-Bromo-9-(1H-indol-3-yl)-3,3-dimethyl-3,4-dihydroacridin-1(2H)-one (4h)

Yield 71%, mp 218-220 °C. IR spectrum, *v*, cm⁻¹: 3298(NH), 1638(C=O), 1616(C=N). ¹H NMR spectrum, *δ*, ppm: 12.20 (s, 1H, indole NH), 8.25 (s, 1H, Ar-H), 8.01 (d, *J* = 8.8 Hz, 1H, Ar-H), 7.91 (dd, *J* = 8.8 and 1.8 Hz, 1H, Ar-H), 7.85-7.82 (m, 1H, Ar-H), 7.60-7.57 (m, 1H, Ar-H), 7.42 (s, 1H, Ar-H), 7.33-7.30 (m, 2H, Ar-H), 2.43 (s, 2H, CH₂), 2.34 (s, 2H, CH₂), 0.99 (s, 6H, 2CH₃). Found, %: C 65.90, H 4.55, N 6.71. Calculated, %: C 65.88, H 4.57, N 6.68. MS, *m/z*: 418 [M]⁺, 420 [M+2]⁺ (1:1).

9-(5-Chloro-2-phenyl-1H-indol-3-yl)-3,3,7-trimethyl-3,4-dihydroacridin-1(2H)-one (4i)

Yield 63%, mp 280-282 °C. IR spectrum, ν , cm⁻¹: 3350(NH), 1655(C=O), 1603(C=N). ¹H NMR spectrum, δ , ppm: 12.42 (s, 1H, indole NH), 8.17 (s, 1H, Ar-H), 8.00 (d, J = 8.6 Hz, 1H, Ar-H), 7.86 (dd, J = 8.6 and 1.8 Hz, 1H, Ar-H), 7.69-7.63 (m, 2H, Ar-H), 7.55-7.49 (m, 3H, Ar-H), 7.34 (s, 1H, Ar-H), 7.21 (d, J = 8.0 Hz, 1H, Ar-H), 7.01 (dd, J = 8.0 and 1.6 Hz, 1H, Ar-H), 2.47 (s, 2H, CH₂), 2.35 (s, 2H, CH₂), 2.31 (s, 3H, CH₃), 1.02 (s, 6H, 2CH₃). ¹³C NMR

spectrum, *δ*, ppm: 195, 167, 143, 141, 140, 139, 138, 136, 131, 130, 126, 125, 55, 50, 32, 30, 22. Found, %: C 77.45, H 5.46, N 5.95. Calculated, %: C 77.49, H 5.42, N 6.02. MS, *m/z*: 463 [M-1]⁺, 465 [M+1]⁺ (3:1).

3,3,7-Trimethyl-9-(5-methyl-2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4j)

Yield 58%, mp 202-204 °C. IR spectrum, *ν*, cm⁻¹: 3387(NH), 1673(C=O), 1599(C=N). ¹H NMR spectrum, *δ*, ppm: 12.34 (s, 1H, indole NH), 8.08 (d, *J* = 8.7 Hz, 1H, Ar-H), 7.95 (s, 1H, Ar-H), 7.86 (dd, *J* = 8.7 and 2.0 Hz, 1H, Ar-H), 7.72-7.69 (m, 2H, Ar-H), 7.60-7.52 (m, 3H, Ar-H), 7.32 (s, 1H, Ar-H), 7.23 (d, *J* = 8.2 Hz, 1H, Ar-H), 7.03 (dd, *J* = 8.2 and 1.4 Hz, 1H, Ar-H), 2.41 (s, 2H, CH₂), 2.34 (s, 2H, CH₂), 2.30 (s, 3H, CH₃), 2.29 (s, 3H, CH₃) 1.00 (s, 6H, 2CH₃). Found, %: C 83.70, H 6.38, N 6.34. Calculated, %: C 83.75, H 6.35, N 6.30. MS, *m/z*: 444 [M]⁺.

3,3,7-Trimethyl-9-(2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4k)

Yield 68%, mp 148-150 °C. IR spectrum, *ν*, cm⁻¹: 3340(NH), 1658(C=O), 1591(C=N). ¹H NMR spectrum, *δ*, ppm: 12.36 (s, 1H, indole NH), 8.00 (d, *J* = 8.6 Hz, 1H, Ar-H), 7.90 (dd, *J* = 8.6 and 2.0 Hz, 1H, Ar-H), 7.86 (d, *J* = 8.2 Hz, 1H, Ar-H) 7.76-7.60 (m, 5H, Ar-H), 7. 52 (d, *J* = 8.2 Hz, 1H, Ar-H), 7.40 (s, 1H, Ar-H) 7.26-7.15 (m, 2H, Ar-H), 2.42(s, 2H, CH₂), 2.33(s, 2H, CH₂) 2.30 (s, 3H, CH₃), 1.00 (s, 6H, 2CH₃). Found, %: C 83.63, H 6.18, N 6.46. Calculated. %: C 83.69, H 6.09, N 6.51. MS, *m/z*: 429 [M-1]⁺.

9-(1H-Indol-3-yl)-3,3,7-trimethyl-3,4-dihydroacridin-1(2H)-one (4l)

Yield 62%, mp 212-214 °C. IR spectrum, ν, cm⁻¹: 3359(NH), 1664(C=O), 1597(C=N). ¹H NMR spectrum, *δ*, ppm: 12.29 (s, 1H, indole NH), 8.24 (s, 1H, Ar-H), 8.00 (d, *J* = 8.6 Hz, 1H, Ar-H), 7.94 (dd, *J* = 8.6 and 1.8 Hz, 1H, Ar-H), 7.87-7.84 (m, 1H, Ar-H), 7.60-7.57 (m, 1H, Ar-H), 7.46 (s, 1H, Ar-H), 7.29-7.26 (m, 2H, Ar-H), 2.40 (s, 2H, CH₂), 2.32 (s, 2H, CH₂), 2.29 (s, 3H, CH₃), 1.01 (s, 6H, 2CH₃). Found, %: C 81.30, H 6.30, N 7.83. Calculated, %: C 81.33, H 6.26, N 7.90. MS, *m/z*: 354 [M]⁺.

9-(5-Chloro-2-phenyl-1H-indol-3-yl)-7-methoxy-3,3-dimethyl-3,4-dihydroacridin-1(2H)-one (4m)

Yield 63%, mp 272-274 °C. IR spectrum, *ν*, cm⁻¹: 3424(NH), 1643(C=O), 1589(C=N). ¹H NMR spectrum, *δ*, ppm: 12.26 (s, 1H, indole NH), 8.15 (s, 1H, Ar-H), 8.00 (d, *J* = 9.2 Hz, 1H, Ar-H), 7.93 (dd, *J* = 9.0 and 2.0 Hz, 1H, Ar-H), 7.80-7.76 (m, 2H, Ar-H), 7.58-7.54 (m, 3H, Ar-H), 7.40 (s, 1H, Ar-H), 7.35 (d, *J* = 8.4 Hz, 1H, Ar-H), 7.00 (dd, *J* = 8.4 and 1.6 Hz, 1H, Ar-H), 3.49 (s, 3H, OCH₃), 2.43 (s, 2H, CH₂), 2.36 (s, 2H, CH₂), 1.01 (s, 6H, 2CH₃). ¹³C NMR

spectrum, *δ*, ppm: 195, 164, 145, 143, 142, 141, 139, 138, 133, 132, 126, 125, 56, 49, 34, 32, 29. Found, %: C 74.93, H 5.21, N 5.80. Calculated, %: C 74.91, H 5.24, N 5.82. MS, *m/z*: 480 [M]⁺, 482 [M+2]⁺ (3:1).

7-Methoxy-3,3-dimethyl-9-(5-methyl-2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (4n)

Yield 65%, mp 206-208 °C. IR spectrum, *ν*, cm⁻¹: 3369(NH), 1664(C=O), 1613(C=N). ¹H NMR spectrum, *δ*, ppm: 12.36 (s, 1H, indole NH), 8.10 (d, *J* = 8.6 Hz, 1H, Ar-H), 7.97 (s, 1H, Ar-H), 7.90 (dd, *J* = 8.6 and 1.8 Hz, 1H, Ar-H), 7.71-7.66 (m, 2H, Ar-H), 7.49-7.40 (m, 4H, Ar-H), 7.26 (d, *J* = 8.2 Hz, 1H, Ar-H), 7.02 (dd, *J* = 8.2 and 1.4 Hz, 1H, Ar-H), 3.48, (s, 3H, OCH₃), 2.40 (s, 2H, CH₂), 2.33 (s, 2H, CH₂), 2.29 (s, 3H, CH₃), 1.00 (s, 6H, 2CH₃). Found, %: C 80.85, H 6.10, N 6.06. Calculated, %: C 80.84, H 6.13, N 6.08. MS, *m/z*: 460 [M]⁺.

7-Methoxy-3,3-dimethyl-9-(2-phenyl-1H-indol-3-yl)-3,4-dihydroacridin-1(2H)-one (40)

Yield 59%, mp 194-195 °C. IR spectrum, *v*, cm⁻¹: 3290(NH), 1656(C=O), 1602(C=N). ¹H NMR spectrum, *δ*, ppm: 12.36 (s, 1H, indole NH), 8.02 (d, *J* = 8.8 Hz, 1H, Ar-H), 7.93 (dd, *J* = 8.8 and 2.0 Hz, 1H, Ar-H), 7.86 (d, *J* = 8.4 Hz, 1H, Ar-H) 7.77-7.62 (m, 5H, Ar-H), 7. 48 (d, *J* = 8.4 Hz, 1H, Ar-H), 7.40 (s, 1H, Ar-H) 7.28-7.20 (m, 2H, Ar-H), 3.51 (s, 3H, OCH₃), 2.42(s, 2H, CH₂), 2.33(s, 2H, CH₂), 1.00 (s, 6H, 2CH₃). Found, %: C 80.71, H 5.90, N 6.29. Calculated, %: C 80.69, H 5.87, N 6.26. MS, *m/z*: 446 [M]⁺.

9-(1H-Indol-3-yl)-7-methoxy-3,3-dimethyl-3,4-dihydroacridin-1(2H)-one (4p)

Yield 66%, mp 185-186 °C. IR spectrum, *ν*, cm⁻¹): 3427(NH), 1669(C=O), 1600(C=N). ¹H NMR spectrum, *δ*, ppm: 12.39 (s, 1H, indole NH), 8.26 (s, 1H, Ar-H), 8.03 (d, *J* = 8.8 Hz, 1H, Ar-H), 7.94 (dd, *J* = 8.8 and 1.8 Hz, 1H, Ar-H), 7.82-7.79 (m, 1H, Ar-H), 7.59-7.55 (m, 1H, Ar-H), 7.39 (s, 1H, Ar-H), 7.31-7.28 (m, 2H, Ar-H), 3.49 (s, 3H, OCH₃), 2.40 (s, 2H, CH₂), 2.32 (s, 2H, CH₂), 1.01 (s, 6H, 2CH₃). Found, %: C 77.79, H 6.02, N 7.53. Calculated, %: C 77.81, H 5.99, N 7.56. MS, *m/z*: 370 [M]⁺.

Antimicrobial activity

Newly synthesized compounds (**4a-4p**) were assessed for *in vitro* antimicrobial activity against a group of two Gram positive (*Bacillus subtilis* and *Staphylococcus aureus*) and two Gram negative (*Escherichia coli* and *Pseudomonas aeruginosa*) bacteria using Streptomycin sulphate as standard. Antifungal activity was performed against three fungal species (*Aspergillus niger*, *Aspergillus flavus* and *Fusarium oxysporum*) by using Ketoconazole as standard. Minimum Inhibitory Concentration (MICs) were determined by broth dilution assay method²⁶. The results of antimicrobial activity are presented in **Table 1**. Among the screened compounds **4a**, **4e**, **4f**, **4g** and **4m** (MIC = 125 μ g/mL) were found to be active against the various bacterial strains under study compared to Streptomycin sulphate (MIC = 62.5 μ g/mL). However, the compounds **4a**, **4b**, **4f**, **4g**, **4i**, **4j**, **4m** and **4n** showed inhibition effect at MIC = 250 μ g/mL against fungal strains *A*. *niger* and *A*. *flavus* while the compound **4e** exhibited same potency as the standard Ketoconazole at MIC = 125 μ g/mL against *A*. *niger*. Analysis of these results shows that indole ring with chloro substitution at five position accounts for increased antibacterial activity. Chloro and methyl groups are responsible for enhanced antifungal activity.

DPPH radical scavenging activity

The interaction of synthesized compounds (4a-4p) with the stable DPPH free radical was investigated *in vitro*²⁷. These interactions will generate α, α -diphenyl- β -picryl hydrazine which leads to discoloration. The degree of discoloration is considered as an indicator of free-radical-scavenging potential of the antioxidant. Butylated hydroxy anisole (BHA) and Ascorbic Acid (AA) were taken as standard antioxidants. Among the compounds 4a and 4i with chloro group at five position of indole ring have shown excellent radical scavenging potential, whereas the compounds 4b, 4c, 4i and 4n with methyl and hydrogen at five position of indole ring exhibited relatively low scavenging ability. In contrast, rest of the compounds have shown least scavenging property. These findings are illustrated in Figure 1.

Ferric reducing antioxidant power activity

All the synthesized compounds were screened *in vitro* to explore their ability to reduce preformed ferric to ferrous ion by ferric reducing antioxidant power assay method²⁹. Butylated hydroxy anisole (BHA) and Ascorbic Acid (AA) were used as reference standard. The compounds showed reducing capabilities in concentration dependent mode. Similar to DPPH assay results, compounds **4a**, **4e** and **4i** with chloro substituted indole ring revealed excellent activity and moderate activity is observed for methyl indolylacridinone analogues **4b**, **4f** and **4j**. Alongside, other indolylacridinone analogues showed least reducing power. The results are compared with standard in **Figure 2**.

Molecular Docking Study

To recognize the structural insight into the potency of synthesized novel indolylacridinone analogues (4a-4p), their binding interactions with the GlcN-6-P synthase crystal structure (PDB ID: 2VF5) were investigated. In the present study, the AutoDock 4.2 software instilled with Lamarckian genetic algorithm was used to envisage best fit ligand's conformation inside the active packet of target enzyme. The identified active packet of GlcN-6-P synthase comprises

12 amino acid residues Ala 602, Val 399, Ala 400, Gly 301, Thr 302, Ser 303, Cys 300, Gln 348, Ser 349, Thr 352, Ser 347 and Lys 603³⁰.

Table 2 summarizes database generated from a total set of 16 compounds docked with the known GlcN-6-P synthase receptor active site. The docking results obtained from ten different ligand-receptor complex conformations of each tested compound were scrutinized and conformation with lowest binding energy was further analyzed for the binding interactions, including hydrogen bonding, hydrophobic and electrostatic interactions. It has been observed that all ligands as well as the reference drug *i.e.* Streptomycin interact inside the receptor active site with binding energies ranging from -8.89 to -4.29 kcal/mol. Best docked conformations of compounds (4a-4p) with GlcN-6-P synthase displayed multiple hydrophobic interactions and hydrogen bonding with active site residues such as Ala 602, Thr352 and Lys 603 by means of one hydrogen bond whereas Streptomycin interacts with residues Ala 602, Cys 300, Thr 352 and Val 399 by four hydrogen bonds (Table 3). The nitrogen of actidinone and indole rings of compounds 4a and 4e were able to form hydrogen bond with residues Thr 352 and Ala 602 respectively. However, the oxygen of methoxy group present at actidinone ring of compound 4p forming hydrogen bond with Ala 602. Figure 3 illustrates the binding interactions of compounds 4a, 4e, 4p and Streptomycin.



	Gram positive bacteria		Gram negative bacteria		Fungi			в. subili Bacill
Compound	B. subtilis	S. aureus	E. coli	P. aeruginosa	A. niger	A. flavus	F. oxysporum	subtilis, aureu
4a	125	250	125	125	250	500	500	Staphylococci
4b	250	250	250	250	250	250	500	Escherich
4c	500	500	250	500	500	500	>1000	coli,
4d	500	250	500	500	1000	500	>1000	aeruginos
4e	125	125	250	- 250	125	500	500	Pseudomon
4f	250	250	125	125	250	250	500	nige
4g	250	500	250	125	500	250	>1000	Aspergilli
4h	1000	500	500	500	500	500	>1000	niger, A. flavu
4i	500	500	1000	500	250	250	500	Aspergilli
4j	1000	1000	500	500	>1000	250	500	oxysporun
4k	500	500	500	1000	500	500	500	Fusariu
41	500	500	250	500	500	500	>1000	oxysporum.
4 m	125	250	125	250	250	250	500	[A
4n	500	500	250	250	250	>1000	500	Streptomyc
4o	250	500	1000	500	500	500	500	sulphate, [B
4 p	1000	500	1000	1000	500	1000	>1000	Ketoconazol
[A]	n.t.	n.t.	n.t.	n.t.	125	125	250	n.t.: not tested
[B]	62.5	62.5	62.5	62.5	n.t.	n.t.	n.t.	

Table 1. The antimicrobial activity results (MIC, µg/mL) of compounds (4a-4p)



Figure 1. DPPH Radical Scavenging Activity of the synthesized compounds (4a-4p).

Figure 2. Ferric Reducing Antioxidant Power Activity of the synthesized compounds (4a-4p).



Table 2. Molecular Docking Interaction of synthesized novel compounds (**4a-4p**) against the GlcN-6-P synthase enzyme (2VF5) Active Site.

^{a,b} Calculated by AutoDock. ^c Number of hydrogen bonds formed between the GlcN-6-P synthase active site and the novel leads.^d

Compound	Binding Energy (Kcal/mol) ^a	Inhibition Constant (µM) ^b	No. of hydrogen bonds ^c	Bond distance ^d	Interacting residues of 2VF5 ^e
4a	-8.00	1.36	1	1.86	Ala 602
4b	-8.64	0.46	1	1.96	Thr 352
4c	-8.47	0.623	1	1.95	Thr 352
4d	-7.71	2.23	1	2.06	Lys 603
4 e	-8.89	0.306	1	2.05	Thr 352
4f	-8.81	0.346	1	2.00	Thr 352
4g	-8.75	0.387	1	2.08	Thr 352
4h	-7.79	1.93		2.07	Thr 352
4i	-8.56	0.530	1	1.97	Thr 352
4j	-8.56	0.527		1.96	Thr 352
4k	-7.74	2.13	1	1.87	Ala 602
41	-7.74	2.12	1	2.01	Lys 603
4m	-8.36	0.745	1	1.94	Thr 352
4 n	-8.54	0.547	1	1.96	Thr 352
40	-8.12	1.12		2.14	Thr 352
4p	-8.87	0.315	1	1.74	Ala 602
Streptomycin Sulphate	-4.29	0.718	4	1.94 2.05 2.19 2.21	Ala 602 Cys 300 Thr 352 Val 399

Hydrogen bond distance ^e The interacting active site residues of GlcN-6-P synthase enzyme with novel leads in the ligand-receptor complex.



Figure 3. The stable and best conformers with least binding energy of the novel analogues 4a, 4e, 4p including the standard streptomycin. (**A**) 4a forming 1H-bond with Ala 602. (**B**) 4e forming 1H bond with Thr 352. (**C**) 4p forming 1H bond with Ala 602. (**D**) Streptomycin forming 4H bond with Ala 602, Cys 300, Thr 352 and Val399. The green dotted lines represent Hydrogen bonds.

CONCLUSION

In conclusion, we have developed a direct, one-pot three component and environmentally benign protocol for the synthesis of novel indole integrated acridinone scaffolds under mild reaction conditions, simple experimental procedure, with good yields and purity. The synthesized compounds were then evaluated for in vitro antimicrobial and antioxidant activities. The compounds 4a, 4e, 4f and 4m exhibited very good antimicrobial activity against the tested microbial strains whereas compounds 4a and 4i emerged as potent radical scavengers with excellent ferric ion reducing ability. Therefore, acridinone ring is responsible for the enhanced activity of indole system. The docking study revealed that all synthesized compounds exhibited strong affinity towards the enzyme GlcN-6-P synthase. Hence, these findings will attract chemists and pharmacologists for further studies in the indole field in search of strong antioxidant and anticancer agents. . K'I'I K

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