Antimicrobial activity of some pyrimidine derivatives in DMF and DMSO

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ABSTRACT. A series of pyrimidine derivatives have been synthesized and their structures were confirmed by IR, 1H NMR and mass spectral data. All these synthesized compounds were tested in vitro for their antimicrobial potential against Gram positive, Gram negative strains of bacteria as well as fungal strains in N, N-dimethyl formamide and dimethyl sulfoxide.

1. INTRODUCTION

Literature survey shows that a large number of pyrimidine derivatives are reported to exhibit antimycobacterial [1], antitumor [2], antiviral [3], anticancer [4], anti-inflammatory, analgesic, antifolate [5], antimicrobial [6], anti-fungal [7], antiproliferative [8] etc.

In the present work, some novel pyrimidine derivatives were synthesized. The characterization of synthesized compounds was done by IR, NMR and mass spectral analysis.

The antimicrobial activity of the synthesized compounds was done against some pathogenic Gram positive and Gram negative bacteria and fungi in N, N-dimethyl formamide and dimethyl sulfoxide.

2. EXPERIMENTAL

Synthesis:

Synthesis of Int-I (2-(benzo[d]thiazol-2-yl) acetonitrile):

To the ethanolic solution of 2-amino thiophenol (0.1 mol), malononitrile (0.1 mol) was added at 0-5 °C temperature, and glacial acetic acid was added drop wise. By the removal of ammonia gas, cyclized product was formed (Int-I). The completion of reaction was confirmed by analytical thin layer chromatography (TLC) using (0.75 : 0.25 Hexane : Ethyl acetate) as mobile phase. The resulting product was filtered, washed with hexane and dried under vacuum. This product was used further in next step.

\[
\begin{align*}
\text{NH}_2 \\
\text{SH}
\end{align*}
\] + \[\text{NC-CN}\] \[\xrightarrow{\text{gla. CH}_3\text{COOH dropwise}}\] \[\text{gla. CH}_3\text{COOH dropwise} \]

\[\begin{align*}
\text{Ethanol 0-5°c} \\
\text{30-35 min}
\end{align*}\]

\[\text{Int-I}\]

\[\begin{align*}
2\text{-amino thiophenol} & \\
\text{Malononitrile}
\end{align*}\]

Synthesis of pyrimidines:

In DMF solution of above synthesized product (Int-I), different substituted aldehydes and guanidine hydrochloride were added. The reaction mixture was refluxed for 8-10 hrs at 145-150 °C using piperidine as catalysis. The complecation of reaction was confirmed by analytical thin layer chromatography (TLC) using (ethyl acetate) as mobile phase. After completion of reaction, the reaction mixture was washed with diethyl ether in order to remove non polar impurities. The resulting solid was filtered.
The physical constants of synthesized compounds are shown in Table 1. The structure confirmation of these crystallized compounds was done by FTIR, $^1$H NMR and mss spectral data. IR spectra were recorded on IR affinity 1S (furier transport infra-red spectroscopy), $^1$H-NMR spectra were taken on a Bruker AVANCE II 400. In all the cases, $^1$H NMR spectra were obtained in DMSO-d$_6$ using TMS as an internal standard. The NMR signals are reported in δppm. Mass spectra were determined using direct inlet probe on a GCMS-QP-2010 mass spectrometer.

3. ANTIMICROBIAL ACTIVITY

Microorganisms tested:

The studied microorganisms were obtained from National Chemical Laboratory (NCL), Pune, India. The microorganisms were maintained at 4°C. The Gram positive bacteria studied were Bacillus cereus ATCC11778 (BC); Staphylococcus aureus ATCC29737 (SA), Corynebacterium rubrum ATCC14898 (CR), Listeria monocytogenes ATCC19112 (LM), Gram negative bacteria were Escherichia coli NCIM2931 (EC), Pseudomonas aeruginosa ATCC27853(PA), Salmonella typhimurium ATCC23564 (ST), Klebsiella pneumoniae NCIM2719 (KP) and fungal strains were Candida glabrata NCIM3448 (CG), Candida epicola NCIM3367 (CE), Candida albicans ATCC2091 (CA) and Cryptococcus neoformans NCIM3542 (CN). The microorganisms studied are clinically important ones causing several infections and food spoilage. In vitro antimicrobial activity of the pyrimidine was studied against pathogenic microbial strains by the agar well diffusion method [9].

Preparation of solutions of compounds:

For all the compounds, DMF and DMSO were used for screening of antimicrobial activity. The solution of 20 mg/ml concentration was prepared for all the compounds.

4. RESULTS AND DISCUSSION:

In total, 10 compounds (AMG-1 to AMG-10) were synthesized. Table 1 shows the physical parameters of these compounds. The IR, $^1$H NMR and mass spectra of compound AMG-2 are shown in Figures 1, 2 and 3 respectively.

Spectral data:

$AMG\text{-}1$: IR (cm$^{-1}$, KBr): 3547.39 (-NH (pri.) str.), 3099.57 (Ar-H str.), 2938.74 (-CH$_2$ sym. str.), 1623.14 (-NH bending vib. Secondary amine), 1358.62 (-CH bending.), 1347.56 (C-$N$ (sec) bending), 1242-1010 (C-H in plane bending, phenyl ring), 752.34 (C-H str. 5-adjecent c atoms), 730.52 (C-Cl str.), $^1$H NMR (DMSO-$d_6$) δ(ppm) : 5.136 (2H, singlet, -NH$_2$), 7.287-7.314 (2H, -CH, multiplet), 7.024 (1H, singlet, -NH), 7.264-7.428 (4H, multiplet, -CH), 7.535-7.610 (1H, multiplet, -NH), 7.630-7.650 (1H, doublet, -CH), 7.904-7.925 (1H, doublet, -CH), MS: (m/z) = 353
AMG-2: IR (cm⁻¹, KBr): 3517.26 (-NH (sec.) str.), 3026.39 (Ar-H str.), 2969.63 (-CH₂ sym. str.), 2896.35 (C-H Str. (alkane), 1627.28 (-NH bending vib. Secondary amine), 1347.69 (-CH bending), 1369.24 (C-N (sec) bending), 1242-1010 (C-H in plane bending, phenyl ring), 751.38 (C-H str. 5-adjacent c atoms), ¹H NMR (DMSO-d₆) δ(ppm): 3.827 (3H, singlet, -OCH₃), 3.939 (3H, singlet-OCH₃), 5.072 (2H, singlet, -NH₂), 6.890-7.925 (2H, -CH, multiplet), 7.024 (1H, singlet, -NH), 7.264-7.428 (4H, multiplet, -CH), 7.630-7.650 (1H, doublet, -CH), 7.904-7.925 (1H, doublet, -CH), MS: (m/z) = 379

AMG-3: IR (cm⁻¹, KBr): 3471.87 (-NH (pri.) str.), 3161.33 (Ar-H str.), 3007.14 (-CH₂ sym. str.), 2227.86 (-CN str.), 1664.62(C=C str. α, β unsaturated 6-member ring), 1604.83(-NH bending vib. Secondary amine), 1381.08 (-CH bending.), 1315.50(C-N (sec) bending), 1242-1010(C-H in plane bending, phenyl ring), 767.69 (C-H str. 5-adjacent c atoms), 767.69(C-Cl str.), ¹H NMR (DMSO-d₆) δ(ppm): 6.789 (2H, singlet, -NH₂), 7.287-7.314 (2H, -CH, multiplet), 7.264-7.428 (4H, multiplet, -CH), 7.440-7.493 (1H, multiplet, -NH), 7.862-7.921 (1H, doublet, -CH), 7.970-8.021 (1H, doublet, -CH) 8.499 (1H, broad, -NH), MS: (m/z) = 353

AMG-4: IR (cm⁻¹, KBr): 3474.27 (-NH (pri.) str.), 3199.68 (Ar-H str.), 2970.38 (-CH₂ sym. str.), 1602.85 (-NH bending vib. primary amine), 1365.60 (-CH bending.), 1242-1010 (C-H in plane bending, phenyl ring), 721.38 (C-H str. 5-adjacent c atoms), 651.94 (C-Br str.), ¹H NMR (DMSO-d₆) δ(ppm): 6.028 (2H, singlet, -NH₂), 7.189-7.204 (2H, -CH, multiplet), 7.324 (1H, singlet, -NH), 7.458-7.624 (4H, multiplet, -CH), 7.815-7.898 (1H, doublet, -CH), 8.014-8.147 (1H, doublet, -CH), 8.106-8.247 (1H, broad, -NH), MS: (m/z) = 398

AMG-5: IR (cm⁻¹, KBr): 3649.32 (-NH (pri.) str.), 3012.57 (Ar-H str.), 2954.95 (-CH₂ sym. str.), 2921.57 (C-H Str. (alkane), 1606.70 (-NH bending vib. primary amine), 1431.18 (-CH bending (alkane)), 1365.60 (-CH bending.), 1242-1010 (C-H in plane bending, phenyl ring), 729.09 (C-H str. 5-adjacent c atoms), ¹H NMR (DMSO-d₆) δ(ppm): 6.369 (3H, singlet, -OCH₃), 6.675 (2H, singlet, -NH₂), 6.962-6.996 (2H, multiplet -CH), 7.024 (1H, multiplet, -NH), 7.180-7.602 (4H, multiplet, -CH), 7.360-7.452 (1H, doublet, -CH), 7.631-7.784 (1H, doublet, -CH), 7.968 (1H, broad, -NH), MS: (m/z) = 349

AMG-6: IR (cm⁻¹, KBr): 3612.57 (-NH (pri.) str.), 3147.28 (Ar-H str.), 2970.41 (-CH₂ sym. str.), 1645.28 (-NH bending vib. primary amine), 1365.60 (-CH bending.), 1242-1010 (C-H in plane bending, phenyl ring), 721.38 (C-H str. 5-adjacent c atoms), 1091.71 (C-F str.), ¹H NMR (DMSO-d₆) δ(ppm): 6.743 (2H, singlet, -NH₂), 6.904-6.948 (2H, -CH, multiplet), 7.000 (1H, multiplet, -NH), 7.258-7.501 (4H, multiplet, -CH), 7.835-7.943 (1H, doublet, -CH), 7.963-8.062 (1H, doublet, -CH), 8.259-8.612 (1H, multiplet, -NH), MS: (m/z) = 337

AMG-7: IR (cm⁻¹, KBr): 3597.46 (-NH (pri.) str.), 3030.28 (Ar-H str.), 2970.38 (-CH₂ sym. str.), 2914.75 (C-H Str. (alkane), 1641.73 (-NH bending vib. primary amine), 1435.04 (-CH bending (alkane)), 1365.60 (-CH bending.), 1242-1010 (C-H in plane bending, phenyl ring), 727.16 (C-H str. 5-adjacent c atoms), ¹H NMR (DMSO-d₆) δ(ppm): 3.512 (3H, singlet, -CH₃), 6.145 (2H, singlet, -NH₂), 6.358-6.476 (2H, multiplet -CH), 6.742 (1H, multiplet, -NH), 6.823-7.015 (4H, multiplet, -CH), 7.263-7.324 (1H, doublet, -CH), 7.534-7.627 (1H, doublet, -CH), 8.017 (1H, broad, -NH), MS: (m/z) = 333

AMG-8: IR (cm⁻¹, KBr): 3481.51 (-NH (pri.) str.), 3151.69 (Ar-H str.), 2970.38 (-CH₂ sym. str.), 2890.24 (C-H Str. (alkane), 1587.42 (-NH bending vib. primary amine), 1435.04 (-CH bending (alkane)), 1365.60 (-CH bending.), 1242-1010 (C-H in plane bending, phenyl ring), 729.09 (C-H str. 5-adjacent c atoms), ¹H NMR (DMSO-d₆) δ(ppm): 3.721 (3H, singlet, -OCH₃), 6.748 (2H, singlet, -NH₂), 6.962-6.996 (2H, multiplet -CH₃), 7.024 (1H, multiplet, -NH), 7.180-7.602 (4H,
multiplet, -CH), 7.813-7.832 (1H, doublet, -CH), 7.943-7.962 (1H, doublet, -CH), 8.439 (1H, broad, -NH), *MS: (m/z) = 349*  

**AMG-9: IR (cm⁻¹, KBr):** 3514.27 (-NH (pri.) str.), 3215.36 (Ar-H str.), 2970.41 (-CH₂ sym. str.), 1602.57 (-NH bending vib. primary amine), 1365.60 (-CH bending.), 1242-1010 (C-H in plane bending, phenyl ring), 704.02 (C-H str. 5-adjacent c atoms), ¹H NMR (DMSO-d₆) δ(ppm) : 6.028 (2H, singlet, -NH₂), 6.745-6.821 (2H, -CH, multiplet), 7.247 (1H, singlet, -NH), 7.432-7.536 (3H, multiplet, -CH), 7.639-7.782 (1H, doublet, -CH), 7.839-7.904 (1H, doublet, -CH), 8.439 (1H, broad, -NH), *MS: (m/z) = 325*  

**AMG-10: IR (cm⁻¹, KBr):** 3377.63 (-NH (sec.) str.), 2903.47 (Ar-H str.), 1596.21 (-NH bending vib. Secondary amine), 1346.35 (-CH bending.), 1302.39 (C-N (sec) bending.), 1242-1010 (C-H in plane bending, phenyl ring), 725.63 (C-H str. 5-adjacent c atoms), ¹H NMR (DMSO-d₆) δ(ppm) : 3.523 (3H, singlet, -OCH₃), 3.963 (3H, singlet-OCH₃), 6.254 (2H, singlet, -NH₂), 6.635-6.702 (2H, multiplet, -CH₂), 7.114 (1H, multiplet, -NH), 7.632-7.728 (3H, multiplet, -CH), 7.830-7.912 (1H, doublet, -CH), 7.934-7.996 (1H, doublet, -CH), 8.012-8.236 (1H, broad, -NH), *MS: (m/z) = 379*  

**Antibacterial activity:**

Figure 4 shows the zone of inhibition of synthesized compound against Gram positive bacteria in DMF and DMSO. In DMF (Figure 4[A]), against BC AMG-5 exhibited maximum inhibition and minimum is shown by AMG-4. The compounds AMG-1, AMG-6 and AMG-8 showed moderate inhibition against BC. Other compounds had no effect at all. In DMSO, against BC, AMG-1 had maximum inhibition whereas minimum inhibition is due to AMG-6. Some of the compounds had no effect against BC. The results suggest that inhibition depends not only on structure of compounds but also on the solvent. Against BC, in DMF, maximum compounds had inhibition as compared to DMSO. So, for this strain, DMF is more effective. The comparison of inhibition among different compounds shows that all the compounds have the same central moiety but different substitution groups. In DMF, AMG-5 containing 4-methoxy group showed maximum inhibition against BC. The compound AMG-8 also contains methoxy group but at 3rd position but its effect is less as compared to AMG-5. This suggests that position of group also affect inhibition. In DMSO, against BC AMG-1 containing 4-chloro group had maximum effect but when chloro group is at 3rd position as in AMG-3, it had no effect at all.

Against SA, AMG-6 exhibited maximum inhibition in DMF whereas AMG-3 had minimum inhibition. Other compounds showed moderate inhibition except AMG-5 and AMG-8. Thus, in DMF against this strain 4-fluoro group is very effective whereas methoxy group at 3rd and 4th position had no effect at all. However, in DMSO against SA, only AMG-1, AMG-5, AMG-6 and AMG-9 showed inhibition and maximum is exhibited by AMG-6 containing 4-fluoro group. The comparison of inhibition against this strain in the two solvents again suggests DMF to be good solvent.

AMG-1, AMG-5, AMG-6, AMG-8, and AMG-9 could inhibit CR in DMF and effect is maximum for AMG-1 containing 4-chloro group. In DMSO, only AMG-5 containing 4-methoxy and AMG-9 containing thiophene inhibit CR.  

For LM, not a single compound was effective in DMF whereas in DMSO, only AMG-5 exhibited inhibition.

All over, AMG-6 (containing 4-fluoro group) could inhibit 75% of Gram positive bacteria in DMF solvent and AMG-5 could inhibit 100% the zone of inhibition in DMSO solvent. Thus overall, AMG-5 having 4-OCH₃ substitution is more effective for studied Gram positive bacteria in both the solvents and DMF is good solvent for the studied Gram positive bacteria.

Figure 5 shows zone of inhibition against Gram negative bacteria in DMF and DMSO. Only AMG-2 and AMG-3 containing 3,4-dimethoxy and 3-chloro groups respectively inhibit EC in
DMF whereas AMG-6 and AMG-8 containing 4-flouro and 3-methoxy groups respectively could inhibit EC in DMSO. Thus, position of group and solvent affect inhibition.

Against PA, only one compound AMG-8 containing 3-methoxy group showed inhibition in DMF whereas in DMSO, AMG-2 and AMG-3 exhibited inhibition. Thus, in DMSO, 3,4-dimethoxy and 3-chloro groups are effective.

In DMF, against ST only AMG-1 and AMG-10 showed inhibition and maximum inhibition is by AMG-1 containing 4-chloro group. AMG-2, AMG-4, AMG-6 and AMG-8 could inhibit ST in DMSO and 3,4-dimethoxy group present in AMG-2 is most effective.

In DMF, AMG-4, AMG-8 and AMG-9 showed inhibition against KP and maximum inhibition is by AMG-4 containing 4-bromo group. None of the compound could inhibit KP in DMSO.

Thus, among studied Gram negative bacteria, for the studied compounds, DMSO is better solvent and KP is the most resistant bacteria. Further, the studied compounds are found to be not as effective against Gram negative bacteria as compared to Gram positive bacteria.

The zones of inhibition against four fungal strains are shown in Figure 6 for all the compounds in DMF and DMSO. Not a single compound could inhibit CG in both DMF and DMSO. Thus, CG is most resistant fungal strain among the selected fungal strains for the studied compounds.

All the studied compounds show moderate activity against CE in DMF whereas only AMG-5 containing 4-methoxy group inhibited CE in DMSO. So, -OCH₃ group at 4ᵗʰ position is more effective for CE in DMSO. Whereas all the groups are effective in DMF.

Except AMG-8 and AMG-9, all the studied compounds showed inhibition against CA in DMF. Whereas only AMG-4 and AMG-5 could inhibit CA in DMF. Against CN, all the compounds except AMG-4, AMG-5 and AMG-10, exhibited inhibition in DMF. However, in DMSO, only AMG-3 and AMG-5 could inhibit.

AMG-5 and AMG-6 having 4-OCH₃ and 4-F substitution respectively. AMG-5 could inhibit 75% zone of inhibition in both the solvents. AMG-6 could inhibit 75% of the studied fungal strains in DMF whereas in DMSO, this compound had no effect at all.

Again, for the studied fungal strains, DMF is better solvent.

5. CONCLUSION

It is concluded that inhibition depends on solvent used, molecular structure of compound and bacterial strain. Overall, DMF is better solvent for the studied compounds and compounds containing methoxy group is most effective. Thus, such compound can act as lead molecule for the synthesis of new therapeutic drug.

<table>
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<th>Compound Code</th>
<th>Substitution R</th>
<th>Molecular formula</th>
<th>Molecular Weight g/mol</th>
<th>Yield (%)</th>
<th>Rf value</th>
<th>Melting point °C</th>
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<td>AMG-1</td>
<td>-4-Cl</td>
<td>C₁₂H₁₂ClN₂S</td>
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Table 1: The physical constants of all the synthesized compounds
Figure 1: IR spectra of compound AMG-2
**Figure 2:** $^1$H NMR spectra of compound AMG-2.

**Figure 3:** Mass spectra of compound AMG-2.
Figure 4: Activity of compounds AMG-1 to AMG-10 against Gram positive bacteria in [A] DMF and [B] DMSO.
Figure 5: Antimicrobial activity of compounds AMG-1 to AMG-10 against Gram negative bacteria in [A] DMF and [B] DMSO.
**Figure 6**: Antifungal activity of compounds AMG-1 to AMG-10 in DMF

**References**


