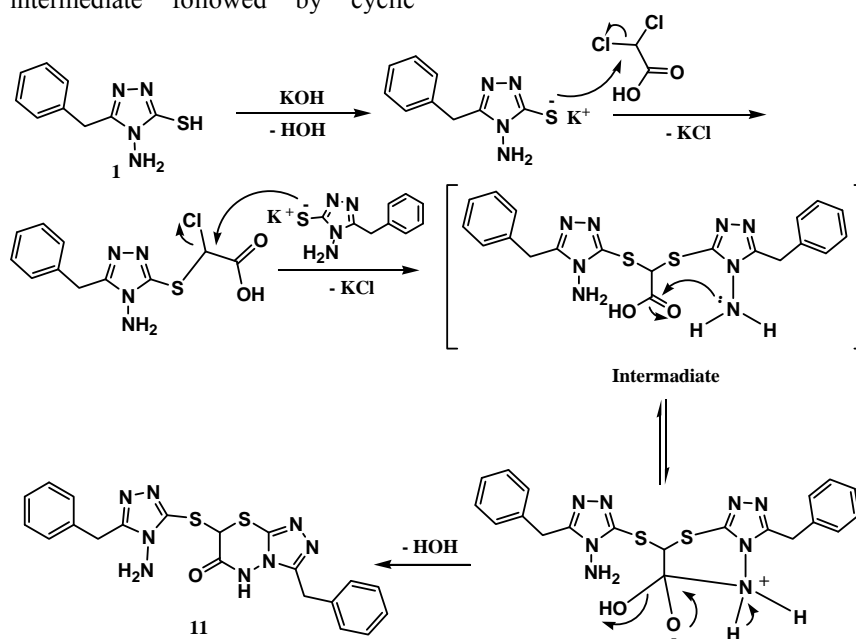


Scheme 5

Scheme 5: Mass fragmentation pattern of compound 11.

Formation of compound 11 may proceed via reaction of two molecules of 1 with one molecule of dichloroacetic acid to give an unstable intermediate followed by cyclic

condensation through losing of water according to the proposed mechanism (Scheme 6).



Scheme 6

Scheme 6: The suggested reaction mechanism of 1 to give compound 11

3.1 Antimicrobial Activity

Bacterial source and culture condition:

The used Bacterial strains were Gram negative bacteria including *E. Coli* (ATCC 25922) and Gram positive bacteria *Enterococcus faecalis* (ATCC 29212). Mueller-Hinton Agar was used as culture media (gl^{-1}) [29], Beef extract, 3.0; Peptone, 17.5; Starch, 1.5; Agar, 17, pH= 7.3 \pm 0.1. The plates were incubated at 37°C for 24 – 48 hrs.

Paper disc technique: Antibacterial activity was determined against the above strains using the paper disc assay method [30]. Whatman number 1 filter paper disc of 6.0 mm diameter was sterilized by autoclaving for 20 min at 121 °C. The sterile discs were impregnated with the spaced apart and plates were incubated at 37°C for 24- 48 hrs [31]. Chloramphenicol 30 μg /disc was used as a positive control. Diameter of the growth inhibition halos caused by the tested compounds were measured and expressed in millimeter. All the assays were carried out in triplicate.

Table 1: Effect of the synthesized compounds (1- 11) on bacterial growth (mm).

Sample No.	Bacterial growth inhibition zone diameter (mm)	
	Gram (-ve) Bacteria	Gram (+ve) Bacteria
	<i>E. Coli</i>	<i>Enterococcus faecalis</i>
1	7	5
2a	6	-----
2b	6	-----
3a	8	7
3b	6	-----
4a	9	7
4b	-----	-----
5	9	-----
6	9	7
7	9	-----
8	7	7
9	7	7
10	7	6
11	-----	-----
Choramphenicol 30 μg (Control)	18	18

E. coli (Escherichia coli) is the name of a germ, or bacterium that lives in the digestive tracts of humans and animals. Many types of *E. coli* can cause bloody diarrhea and urinary tract infections. Some strains of *E. coli* bacteria may also cause severe anemia or kidney failure [32]. Also, *Enterococci* are Gram-positive cocci that often occur in pairs (diplococci) or short chains. The important clinical infections caused by *Enterococcus* include urinary tract infections, bacteremia, bacterial endocarditis, diverticulitis, and meningitis [33].

The antibacterial activity of the synthesized compounds **1-11** were carried out on the growth of two pathogenic bacteria (*E. Coli* and *Enterococcus faecalis*). The data obtained in Table (1) indicate that 12/14 of these compounds have effects on *E. Coli* bacteria where the great inhibition (9 mm) was observed by the Mannich base **4a**, benzenesulfonamide **5**, acetamide **6** and the triazolothiadiazine **7**, but the low

inhibition (6 mm) was appeared by Schiff bases **2a, b** and the thiazolidinone **3b**. The triazole **1**, thiazolidinone **3a**, Phenyl-thioacetic acid ester **8**, thiadiazinone **9** and thiadiazinedione **10** showed moderate inhibition (7- 8 mm), while the Mannich base **4b** and the triazolothiadiazinone **11** showed no activity.

The inhibition effect was decreased on *Enterococcus faecalis* and also 7/14 only of the tested compounds showed moderate and low inhibition effect (5- 7 mm).

4. Conclusion

In summary, 4-amino-5-benzyl-4H-[1,2,4]triazole-3-thiol (**1**) has been utilized as a key starting material in the synthesis of many novel heterocyclic compounds **2- 11**. The constitution of these compounds assigned on the basis of IR, ^1H ^{13}C NMR, mass spectra and elemental analyses were found to be in correlation with the desired structure. The antimicrobial activity screening revealed that the compounds **1- 10** have significant antimicrobial activity. Compounds **4a, 5, 6** and **7** have high biological activity against gram (-ve) bacteria, and compounds **1, 3a, 8, 9** and **10** showed moderate inhibition effect.

On the other hand, compounds **1, 3a, 4a, 6, 8, 9** and **10** showed moderate inhibitions effect against gram (+ve) bacteria, while compounds **2a, 2b, 3b, 4b, 5, 7** and **11** showed no activity. The results are promising and show that the fine tuning of the structures **4a, 5, 6** and **7** can lead to some new antimicrobial agents in treating microbial infections.

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