

### Synthesis, Characterization, Anti-mycobacterial evaluation, In-Vitro and Computational analysis of Pyrazole derivative

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#### **ABSTRACT**

Pyrazole derivatives have gained considerable interest in medicinal chemistry due to their diverse pharmacological properties, including antituberculosis activity. This study focuses on the synthesis, characterization, ADMET-SAR predictions, and biological evaluation of substituted pyrazole derivatives of 3-(4-methoxyphenyl)-3-oxopropanoate.) The synthesized compounds were characterized using spectral techniques, including NMR, IR, and mass spectroscopy. ADMET-SAR analysis predicted the pharmacokinetics and drug-likeness of these compounds, and their antituberculosis activity was evaluated against *Mycobacterium tuberculosis* strains. The results indicated promising biological activity and favourable drug-like properties, suggesting their potential as lead molecules for the development of antituberculosis drugs.

#### **KEYWORDS**

Pyrazole, Hydrazides, antituberculosis, ADMET-SAR, drug design, pharmacokinetics.

### **INTRODUCTION**

Tuberculosis (TB), caused by *Mycobacterium tuberculosis*, remains a global health challenge.<sup>1-3</sup> The emergence of multidrug-resistant (MDR) and extensively drug-resistant (XDR) strains necessitates the discovery of novel therapeutic agents<sup>4</sup>. Pyrazole derivatives have emerged as promising scaffolds due to their diverse biological activities, including antituberculosis properties<sup>5-8</sup>. This study explores the synthesis and biological evaluation of pyrazole derivatives of 3-(4-methoxyphenyl)-3-oxopropanoate, a chemical scaffold with potential antituberculosis activity. Various pyrazole derivatives containing quinoline, furan, indole, pyridine, pyrimidine, coumarin, purine, pyrrole, benzofuran, benzoxazoles, etc, are reported in the literature showing anti-tuberculosis activity<sup>9</sup>.

Tuberculosis (TB) remains one of the deadliest infectious diseases worldwide, with an estimated 10 million cases and 1.5 million deaths annually, according to the World Health Organization (WHO)<sup>10-11</sup>. Despite significant advancements in medical science, TB continues to pose a global health threat, particularly in developing and underdeveloped nations. One of



the critical challenges in combating TB is the emergence of multidrug-resistant (MDR) and extensively drug-resistant (XDR) strains of *Mycobacterium tuberculosis*. These strains have rendered many first-line and second-line TB drugs ineffective, necessitating the urgent development of novel therapeutic agents <sup>12-15</sup>.

Pyrazole is a five-membered heterocyclic compound containing two adjacent nitrogen atoms in its ring structure. The pyrazole scaffold has attracted substantial attention in medicinal chemistry due to its versatility and broad spectrum of biological activities. As a five-membered heterocyclic compound containing two nitrogen atoms, pyrazole derivatives have demonstrated diverse pharmacological properties, including anticancer, anti-inflammatory, antimalarial, and antituberculosis activities <sup>16-20</sup>. In particular, the structural adaptability of pyrazole derivatives enables extensive modifications, thereby enhancing their interactions with specific biological targets. It is very important to note that studies published by Desale et al (2019), Haroon ur Rashid et al, Diego G. Ghiano et al, and Rafal M. Mohareb, et al. have shown very peculiar anti-tubercular activities [1.5- 12.5 μg/ml]<sup>21-24</sup>.

### Natural Products and Drugs Containing Pyrazole Moieties<sup>25</sup>:

Among the pyrazole derivatives, 3-(4-methoxyphenyl)-3-oxopropanoate is a promising precursor, offering unique structural features suitable for antituberculosis drug design. The incorporation of functional groups into this scaffold can improve its pharmacokinetic properties, including absorption, distribution, metabolism, excretion, and toxicity (ADMET). Advances in computational techniques, such as ADMET-SAR (Structure-Activity Relationship) modelling, have enabled researchers to predict the pharmacokinetic and toxicological profiles of drug candidates, accelerating the drug discovery process <sup>26-30</sup>.



This study aims to synthesize and characterize substituted pyrazole derivatives of 3-(4-methoxyphenyl)-3-oxopropanoate and evaluate their antituberculosis potential. The synthesis of novel compounds involves optimizing reaction conditions to ensure high yields and purity. Characterization techniques such as nuclear magnetic resonance (NMR), infrared (IR) spectroscopy, and mass spectrometry are employed to confirm the molecular structures of the synthesized compounds.

Furthermore, the study leverages ADMET-SAR analysis to predict the pharmacological profiles of the compounds, ensuring their compatibility with drug-like properties. The synthesized compounds are then tested against *Mycobacterium tuberculosis* strains to assess their biological activity. This dual approach of computational and experimental evaluation offers a comprehensive understanding of the compounds' potential as antituberculosis agents. By minimizing the use of hazardous reagents and optimizing reaction conditions, the study not only contributes to the discovery of new TB drugs but also promotes environmentally friendly research methodologies.

#### **MATERIAL AND METHODS:**

All chemicals and solvents were procured from Merck chemicals and further purified them whenever necessary. The Thin layer chromatographic technique (TLC 0.25 mm E-Merck silica gel 60 F254 pre-coated plates) was implemented to conduct all necessary reactions. TLC plates after spotting were further visualised with UV light. All compounds were tested for their respective melting points using apparatus from Sunder Industrial Products, Mumbai, and they were all uncorrected. We have recorded <sup>1</sup>H-NMR spectra on a 300 MHz Agilent instrument. The FT–IR spectral analysis was carried out by using the Perkin Elmer Tensor-II model <sup>31-32</sup>.

### **SYNTHESIS OF PYRAZOLES:**

**Synthesis of different pyrazoles derivatives(3a-3j):** Pyrazoles derivatives were synthesized from substituted aryl hydrazide(1a-j) and ethyl 3-(4-methoxyphenyl)-3-Oxoproanoate (2)



 $(5-hydroxy-3-(4-methoxyphenyl)-1 \\ H-pyrazol-1-yl) (phenyl) methan one \\$ 

3 a-j

Add an equimolar quantity of 1,3-diketones/ketoesters and acyl hydrazide into absolute ethanol containing a catalytic amount of acetic acid, and reflux the reaction mixture till completion of the reaction in a water bath. The reaction takes 1-4 hrs to complete. Monitor the progress of the reaction by performing TLC time to time in ethyl acetate -pet ether (1:9) as the mobile phase. After completion of the reaction, distil out the solvent and cool it to room temperature stir for a few minutes. The solid product formed was filtered, washed with water and cold ethanol, dried, and then crystallized from ethanol.

TABLE 1: Experimental parameters and melting point of different synthesised pyrazole Derivatives

Hydrazides: 1a-1j	Pyrazole derivatives: 3a-3j	Yield (%)	Colour	m.p. (°C)
N NH <sub>2</sub>	O O O O O O O O O O O O O O O O O O O	68	Buff white	190
NO <sub>2</sub> NH <sub>2</sub>	O O <sub>2</sub> N OH	70	white	180
NO <sub>2</sub> NH <sub>2</sub>	O N N NO <sub>2</sub>	63	Buff white	204
O <sub>2</sub> N NH <sub>2</sub>	O O N NO <sub>2</sub>	60	Buff white	152



O <sub>2</sub> N NH <sub>2</sub> NH <sub>2</sub> NO <sub>2</sub> 1e	O N N OH NO <sub>2</sub>	65	Light Yellow	240
N NH <sub>2</sub>	O H O C C C C C C C C C C C C C C C C C	61	Buff white	180
O NH <sub>2</sub>	O O O CI	66	Buff white	272
CI NH2	O H O C C	72	Buff white	160
HO NH <sub>2</sub> HO OH	OH OH OH	62	Buff white	220
H <sub>3</sub> C N NH <sub>2</sub>	O O O O O O O O O O O O O O O O O O O	68	Buff white	320

TABLE 2: Spectral characterization of synthesised hydrazones (3a-3j)

Pyrazole	FT- IR da	ıta (cm <sup>-1)</sup>			<sup>1</sup> HNMR	chemical	shifts of	
derivatives				proton in δ				
	Enolic-	>N-	Ar-O -	C-O-C	Enolic-	> N-	O-CH <sub>3</sub>	
	ОН	C=O		Stretching	ОН	С=СН		
3a	3200	1666	1250	1029	10.34	5.75	3.82	
3b	3162	1635	1250	1029	10.95	5.71	3.81	
3c	3601	1667	1251	1029	11.05	5.70	3.82	
3d	3698	1670	1277	1030	10.98	5.72	3.81	



3e	3303	1608	1270	1031	11.69	5.73	3.81
3f	3698	1678	1251	1029	10.81	5.73	3.81
3g	3697	1681	1249	1029	10.61	5.52	3.81
3h	3123	1697	1250	1028	11.65	5.72	3.80
3i	3292	1670	1249	1028	11.65	5.72	3.79
3j	3698	1593	1250	1028	11.65	5.71	3.80

#### **EXPERIMENTAL DATA:**

### 1. (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl) (phenyl) methanone (3a) Yield (%): 68; m.p. (<sup>0</sup>C): 190.

FT-IR (cm<sup>-1</sup>): 3200, 3000.67, 2836,31, 1666, 1628, 1618, 1532, 1577, 1487, 1445, 1286.15, 1249.94, 1068, 1029.8, 869, 836, 795.

<sup>1</sup>HNMR (300 MHz, D<sub>6</sub> DMSO) δ ppm): 3.82 (s) 3H, 5.75 d (s) 1H, 6.91 (d, J:8.7Hz) 2H, 7.99 (d, J:7.5Hz) 2H, 7.47 (t, J:7.5Hz) 2H, 7.55 (t, J:8.7Hz) 3H, 10.34 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{14}N_2O_3$  found C, 69.38; H, 4.79; N, 9.52; O, 16.37. m/Z: 294.10

# 2. (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl) (2-nitrophenyl) methanone (3b)

Yield (%):70; m.p. (°C): 180.

FT-IR (cm-1): 3162, 3012, 2837, 1598, 1580, 1573, 1531, 1498, 1485, 1442, 1250, 1029, 836, 761, 660.

<sup>1</sup>H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.81 (s) 3H,5.71 (s) 1H, 6.90 (d, J:8.7Hz) 2H, 7.56 (d, J:8.7Hz) 2H, 7.69 (m) 2H, 8.07 (d, J:8.1Hz) 2H, 10.95 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{13}N_3O_5$  found C, 60.18; H, 3.86; N, 12.38; O, 23.58. m/Z: 339.09

## 3. (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl) (3-nitrophenyl) methanone (3c)

Yield (%): 63; m.p. (<sup>0</sup>C): 204.

FT-IR (cm-1): 3601, 3073, 2849, 1667, 1608, 1573, 1531, 1351, 1251, 1029, 835, 772, 726, 658.

<sup>1</sup>H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.82 (s) 3H, 5.7 (s) 1H, 6.91 (d, J:9.0 Hz) 2H, 7.71 (d, J:7.5 Hz) 2H, 7.75 (m, J:7.5 Hz) 2H, 8.41 (s) 1H, 8.96 (s) 1H, 11.05 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{13}N_3O_5$  found C, 60.18; H, 3.86; N, 12.38; O, 23.58. m/Z: 339.09

# 4. (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl) (4-nitrophenyl) methanone (3d)

Yield (%): 60; m.p. (<sup>0</sup>C): 152.

FT-IR (cm-1): 3698, 3169, 2840,1580, 1514,1445, 1470, 1298, 1277, 1177, 838, 790, 710.



<sup>1</sup>H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.81 (s) 3H, 5.72 (s) 1H, 6.90 (d, J:6.9 Hz) 2H, 7.56 (d, J:6.9 Hz) 2H, 7.87 (d, 7.2Hz) 2H, 8.24 (d, J:7.2Hz) 2H, 10.98 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{13}N_3O_5$  found C, 60.18; H, 3.86; N, 12.38; O, 23.58. m/Z: 339.09

## 5. (3,5-dinitrophenyl) (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl) methanone (3e)

Yield (%): 65; m.p. (°C): 240.

FT-IR (cm-1): 3303, 3214, 3106, 2979, 1608, 1583, 1542.47, 1520, 1495, 1485,1440, 1323,1270, 1257, 1178, 1031, 837, 816, 730, 653.

<sup>1</sup>H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.81 (s) 3H, 5.73 (s) 1H, 6.9 d (d) 2H, 7.56 (d) 2H, 9.16 (s) 1H, 9.35 (s) 2H, 11.69 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{12}N_4O_7$  found C, 58.13; H, 3.15; N, 14.58; O, 29.14; m/Z: 384.07

### **6.** (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl) (pyridin-3-yl) methanone (3f) Yield (%): 61; m.p. ( $^{0}$ C): 180.

FT-IR (cm-1): 3698, 3383.64, 3196.90, 2980, 1678, 1651, 1594.71, 1542, 1476, 1251, 1029, 837, 773, 697.

1H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.81 (s) 3H, 5.73 (s) 1H, 6.9 (d) 2H, 7.55 (d) 2H, 7.93 (s, J: 6.6Hz), 1H, 8.30 (s), 1H, 8.75 (s) 1H, 9.17 (s) 1H, 10.81 (s) 1H.

Elemental analysis Calculated for:  $C_{16}H_{13}N_3O_3$  found C, 65.08; H, 4.44; N, 14.23; O, 16.25; m/Z: 295.10.

# 7. (4-chlorophenyl)(5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl)methanone (3g)

Yield (%): 66; m.p. (<sup>0</sup>C): 272.

FT-IR (cm-1): 3697, 3168, 2980, 2865, 1681, 1639, 1589, 1561, 1437, 1296, 1249, 1175, 1063, 1029, 1010, 079, 835, 680.

1H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.81 (s) 3H, 5.52 (s) 1H, 6.92 (s, J:7.5Hz) 2H, 7.56 (d, J:7.5Hz) 2H, 7.97 (d, J:7.2Hz) 2H, 8.15(d) 2H, 10.61 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{13}ClN_2O_3$  found C, 62.11; H, 3.99; N, 13.58; O, 14.60.

m/Z: 328.06

### 8. (2,4-dichlorophenyl)(5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl) methanone (3h)

Yield (%): 72; m.p. (<sup>0</sup>C): 160.

FT-IR (cm-1): 3123, 2959, 2836, 1697, 1616, 1582, 1492, 1474, 1297, 1250, 1028, 817, 743, 678.

1H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.80 (s) 3H, 5.72 (s) 1H, 6.91 (d, J:8.1Hz) 2H, 7.44 (d) 2H, 7.56 (m, J:8.1Hz) 2H, 8.10 (s) 1H, 11.65 (s)1H.

Elemental analysis Calculated for:  $C_{17}H_{12}Cl_2N_2O_3$  found C, 56.22; H, 3.33; N, 8.52; O, 13.22. m/Z: 362.02



# 9. (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl)(3,4,5-trihydroxyphenyl) methanone (3i)

Yield (%): 62; m.p. (<sup>0</sup>C): 220.

FT-IR (cm-1): 3292, 2980, 1670, 1599, 1249, 1028, 817, 743, 678;

1H-NMR (δ in ppm, 300 MHz, D<sub>6</sub> DMSO): 3.79 (s) 3H, 5.72 (s) 1H, 6.91 (d) 2H, 7.56 (d) 2H, 7.81 (s) 2H, 9.91 (s) 3H, 11.65 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{14}N_2O_6$  found C, 59.65; H, 4.12; N, 8.18; O, 28.05. m/Z: 342.09

## 10. (5-hydroxy-3-(4-methoxyphenyl)-1H-pyrazol-1-yl)(6-methylpyridin-3-yl) methanone(3j)

Yield (%): 68; m.p. (<sup>0</sup>C): 320.

FT-IR (cm-1): 3698, 2923, 2853, 1583, 1473, 1428, 1410, 1250, 1175, 1028, 794, 675;

1H-NMR ( $\delta$  in ppm, 300 MHz, D<sub>6</sub> DMSO): 1.25 (s) 3H, 3.80 (s) 3H, 5.71 (s) 1H, 6.91 (d) 2H, 7.56 (d) 2H, 8.06 (m) 3H, 11.65 (s) 1H.

Elemental analysis Calculated for:  $C_{17}H_{15}N_3O_3$  found C, 67.01; H, 4.89; N, 13.58; O, 15.52. m/Z: 309.11

#### **RESULT AND DISCUSSION:**

<sup>1</sup>H-NMR, FT-IR Spectral characterization: The pyrazoles synthesized showed medium absorption between 3600-3200 cm<sup>-1</sup>, which is absent in the parent molecule, depicting an enol -OH group. In a molecule. Further absorption of the amide carbonyl group (pyrazoles) of the amide group. in the region 1660-1600 cm<sup>-1</sup> (medium to strong) and at 1420 cm due to C=N stretching confirmed the formation of the pyrazoles ring. Further, all newly synthesized pyrazole derivative molecules have a medium absorption pattern at 1340-1100 cm<sup>-1</sup> due to bending vibration and at 1307-1246 cm<sup>-1</sup> due to stretching vibration of the C-N bond. All the compounds showed an absorption band in the region of 1250-1150 cm<sup>-1</sup> due to asymmetrical stretching and at 1020-1075 due to symmetrical stretching, confirming the ethereal linkage(C-O-C) in them.

The peak in the <sup>1</sup>HNMR due to the -CH<sub>2</sub>-CO- proton in the region of 3.7 to 4.2 ppm is absent, indicating the formation of an enolic form, which is confirmed by a peak showing a weak singlet in the deshielded area at 10-11 ppm. The data of chemical shift of >N-C=CH of newly synthesized pyrazole molecules 3a-j are summarized in Table 5. The prominent singlet at 3.79-3.81 ppm occurred due to the presence of the etherical proton of CH<sub>3</sub>-O in all newly synthesized pyrazole derivatives in the deshielded zone.

We synthesized all ten pyrazoles for the sake of an efficient and cost-effective anti-tuberculosis agent and the biological importance of pyrazoles.



### STUDY OF IN VITRO ANTI-MYCOBACTERIUM TUBERCULOSIS ACTIVITY:

Mycobacteria strain used for the in vitro anti-mycobacterial tuberculosis (vaccine strain, H37RV strain): ATCC no. 27294. The anti-Tuberculosis study was performed by using the popular Microplate Alamar blue assay (MABA), as it is one of the best methods for analysis <sup>33</sup>. We added 200 μL of sterile deionised water into 96 perimeter wells of a sterile plate in order to minimise the evaporation of test medium in the wells during incubation. All the wells received 100 μL of the middle broth 7H9 and were allowed to be serially diluted with the test compound directly on plates. The test compounds' concentration varied from 100 to 0.2 μg/mL. All the plates were covered and sealed with parafilm and incubated at 37 °C for 5 days. After that, 25 μL of freshly prepared 1:1 mixture of Alamar blue reagent and 10% Tween 80 was added to each plate and incubated further for 24 hours. Blue colour in the well was interpreted as no bacterial growth, and pink colour was treated as bacterial growth. We recorded MIC (lowest drug concentration that prevented colour changes from blue to pink) values with reference to three anti-TB drugs, namely Pyrazinamide, ciprofloxacin, and streptomycin. All the results of newly synthesised compounds are in Table 4.

All the newly synthesized hydrazones (3a-3j) showed good to moderate in vitro antituberculosis activity as tested with the Microplate Alamar Blue assay (MABA) against Mycobacterium tuberculosis (vaccine strain, H37RV strain) (34). Compounds 3b MIC value:  $12.5 \,\mu\text{g/mL}$ ) and 3a is  $25 \,\mu\text{g/mL}$ ) were found to be the active compounds (Pyrazinamide and ciprofloxacin, MIC value:  $3.125 \,\mu\text{g/mL}$ ). 3c-j showed moderate MIC values of  $50 \,\mu\text{g/mL}$ . It is worth noting that, after some structural modifications, these candidates may become promising anti-TB drug candidates in the future.

Table 3: Minimum Inhibitory Concentration Values for Compounds (3a-3j) against H37rv Strain

Test samples	3a	3b	3c	3d	3e	3f	3g	3h	3i	3j	Pyr.	cipro	Strepto
MIC value (μg/mL)	25	12.5	50	50	50	50	50	50	50	50	3.125	3.125	6.250

Table 4: Probable Values of In-Silico ADMET Properties of 3a-3j molecules using AdmetSAR 3.0

Description/		Probable value of Pyrazole derivatives									
Properties	3a	3b	3c	3d	3e	3f	3g	3h	3i	3j	
Molecular weight	294.31	339.30	339.30	339.30	384.30	295.29	328.75	363.2	342.30	309.32	
A log P	3.368	3.17	3.134	3.149	3.104	1.819	3.939	4.893	1.62	2.571	

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Synthesis, Characterization, Anti-mycobacterial evaluation, In-Vitro and Computational analysis of Pyrazole derivative



H-bond Acceptor	5	7	7	7	9	6	5	5	8	6
H-bond Donor	1	1	1	1	1	1	1	1	4	1
Rotatable Bonds	3	4	4	4	5	3	3	3	3	3
Human intestinal	0.947	0.958	0.961	0.951	0.938	0.988	0.972	0.972	0.652	0.984
absorption (HIA)	0.747	0.736	0.701	0.731	0.756	0.766	0.772	0.772	0.032	0.704
Caco-2	0.607	0.374	0.464	0.348	0.361	0.957	0.767	0.741	0.052	0.887
Blood brain	0.617	0.711	0.574	0.475	0.446	0.886	0.707	0.741	0.032	0.87
barrier (BBB)	0.017	0.711	0.574	0.473	0.440	0.880	0.712	0.764	0.113	0.67
OATP2B1	0.362	0.446	0.459	0.458	0.526	0.103	0.393	0.486	0.517	0.205
inhibitor	0.302	0.770	0.737	0.436	0.520	0.103	0.575	0.400	0.517	0.203
OATP1B1	0.893	0.832	0.793	0.736	0.7	0.968	0.85	0.814	0.798	0.939
inhibitor	0.093	0.032	0.793	0.730	0.7	0.908	0.65	0.014	0.796	0.939
OATP1B3	0.893	0.788	0.783	0.742	0.682	0.982	0.862	0.799	0.782	0.955
inhibitor	0.893	0.700	0.763	0.742	0.082	0.982	0.802	0.799	0.782	0.933
MATE1 inhibitor	0.222	0.252	0.246	0.246	0.275	0.12	0.207	0.239	0.253	0.159
OCT2 inhibitor	0.16	0.175	0.157	0.158	0.191	0.308	0.159	0.192	0.244	0.293
BSEP inhibitor	0.759	0.933	0.909	0.922	0.904	0.644	0.904	0.949	0.345	0.838
P-glycoprotein	0.439	0.821	0.816	0.814	0.886	0.307	0.640	0.760	0.217	0.667
inhibitor	0.00	0.125	0.120	0.215	0.157	0.214	0.075	0.004	0.200	0.521
P-glycoprotein	0.09	0.135	0.139	0.215	0.157	0.314	0.075	0.084	0.298	0.531
substrate	0.202	0.60	0.606	0.704	0.672	0.467	0.574	0.554	0.056	0.070
CYP3A4 substrate	0.283	0.68	0.606	0.724	0.673	0.467	0.574	0.774	0.056	0.878
CYP2C9 substrate	0.342	0.687	0.672	0.721	0.703	0.463	0.638	0.766	0.036	0.755
CYP2D6 substrate	0.183	0.253	0.244	0.317	0.245	0.263	0.249	0.328	0.022	0.577
CYP3A4	0.333	0.813	0.705	0.607	0.703	0.462	0.447	0.494	0.170	0.238
inhibition										
CYP2C9	0.831	0.943	0.897	0.843	0.88	0.745	0.897	0.919	0.407	0.576
inhibition										
CYP2D6	0.095	0.126	0.104	0.082	0.156	0.24	0.126	0.174	0.226	0.134
inhibition										
CYP1A2	0.795	0.699	0.694	0.314	0.663	0.806	0.799	0.833	0.803	0.626
inhibition										
Rodent	0.335	0.408	0.502	0.468	0.547	0.432	0.368	0.332	0.474	0.396
Carcinogenicity										
Eye corrosion	0.245	0.027	0.035	0.013	0.037	0.002	0.186	0.165	0.092	0.001
Eye irritation	0.96	0.486	0.57	0.19	0.467	0.03	0.903	0.858	0.902	0.024
Ames	0.326	0.716	0.852	0.792	0.893	0.276	0.492	0.47	0.264	0.471
mutagenesis										
Human either-a-	0.374	0.772	0.807	0.753	0.866	0.239	0.757	0.909	0.549	0.796
go-go inhibition										
(10-30 uM)										
Micronuclear	0.509	0.849	0.822	0.857	0.832	0.827	0.614	0.6	0.409	0.882
Nephrotoxicity	0.502	0.534	0.644	0.534	0.641	0.393	0.598	0.57	0.424	0.381
Respiratory	0.078	0.29	0.213	0.449	0.365	0.917	0.097	0.0093	0.409	0.82
toxicity										
Acute Oral	0.591	0.466	0.499	0.666	0.443	0.824	0.673	0.52	0.094	0.765
Toxicity (c)										
Estrogen receptor binding	0.346	0.358	0.448	0.453	0.425	0.059	0.541	0.677	0.401	0.22
Androgen	0.297	0.523	0.501	0.479	0.509	0.114	0.418	0.5	0.315	0.349
receptor binding	0.271	0.525	0.501	0.7/3	0.509	0.117	0.710	0.5	0.515	0.577
Thyroid receptor	0.193	0.287	0.373	0.334	0.383	0.117	0.412	0.587	0.167	0.244
binding	0.173	0.207	0.575	0.557	0.505	0.11/	0.712	0.507	0.107	0.277
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Glucocorticoid receptor binding	0.227	0.45	0.468	0.498	0.529	0.152	0.423	0.574	0.237	0.308
Aromatase binding	0.474	0.452	0.441	0.347	0.395	0.419	0.532	0.526	0.392	0.46
Honey bee toxicity	0.073	0.059	0.035	0.052	0.046	0.045	0.065	0.109	0.032	0.033
Biodegradation	0.098	0.037	0.025	0.017	0.41	0.56	0.025	0.023	0.266	0.036
Crustacea aquatic toxicity	0.527	0.652	0.537	0.518	0.563	0.173	0.728	0.871	0.165	0.209
Fish aquatic toxicity	0.736	0.754	0.709	0.658	0.711	0.327	0.638	0.903	0.38	0.349
Water solubility (log S)	-3.854	-5.106	-4.963	-5.026	-4.977	-3.354	-4.565	-5.603	-3.252	-4.455
Plasma protein binding (100%)	0.978	0.988	0.939	0.937	0.9	0.821	0.996	0.971	0.11	0.953

TABLE 5: Chemical toxicity prediction of Pyrazole derivatives using STopTox 1.0 software

Mole	Acute in Toxicity	halation	Acute or toxicity	al	Acute de Toxicity	rmal	Eye irri Corrosi	tation & on	Skin sensitiz	zation	Skin irritati Corrosion	on &
- cule	Predi ction	Confi dence	Predi ction	Confi dence	Predi ction	Confi dence	Predi- ction	Confi- dence	Predi ction	Confi dence	Predi ction	Confi dence
3a	Non Toxic	65	Toxic	58	Non Toxic	74	Toxic	78	Non- Sensitizer	70	Negative	90
3b	Non Toxic	66	Non- Toxic	51	Non Toxic	63	Toxic	79	Non sensitizer	60	Negative	90
3c	Non Toxic	66	Toxic	51	Non toxic	63	Toxic	79	Non Sensitizer	60	Negative	90
3d	Non toxic	66	Toxic	51	Non toxic	63	Toxic	79	Non Sensitizer	60	Negative	90
3e	Non Toxic	66	Toxic	51	Non Toxic	63	Toxic	79	Non Sensitizer	60	Negative	90
3f	Non Toxic	65	Toxic	59	Non toxic	75	Toxic	78	Non Sensitizer	70	Negative	90
3g	Non Toxic	66	Toxic	58	Non toxic	75	Toxic	76	Non sensitizer	70	Negative	80
3h	Non Toxic	66	Toxic	58	Non toxic	75	Toxic	76	Non sensitizer	60	Negative	70
3i	Non Toxic	73	Non Toxic	66	Non Toxic	78	Toxic	76	Non sensitizer	60	Negative	80
3j	Non Toxic	78	Toxic	58	Non Toxic	77	Toxic	83	Non Sensitizer	70	Negative	90

TABLE 6: Drug likeness, Drug Score, Toxicity Risk, and TSPA of 3a-3j using the Osiris program.

	c log P	solubility	Mol wt.	Drug likeness	Drug score	TSPA
3a	3.16	-4.9	294.31	4.07	0.89	64.35
3b	2.24	-5.36	339.31	3.16	0.90	110.17
3c	2.24	-5.36	339.31	-2.33	0.88	110.17
3d	2.24	-5.36	339.31	-8.14	0.86	110.17
3e	1.32	-5.82	384.30	-9.99	0.08	155.99
3f	2.16	-4.10	295.30	-5.21	0.93	77.24
3g	3.77	-5.64	328.75	-5.46	0.83	64.35



3h	4.37	6.37	363.20	4.92	0.76	64.35
3i	2.12	-4.01	342.31	5.28	0.91	125.04
3j	2.56	-4.47	309.32	3.92	0.91	77.24

### **IN-VITRO SILICO ADMET PREDICTIONS:**

All synthesised pyrazoles showed good drug score and drug likeness, along with good bioactivity scores. Their Admet SAR analysis suggested that there might be strong interaction with different receptors, ligands, and enzymes<sup>34</sup>. The result of most of the newly synthesised compounds physiochemical parameters were found to be within the range set by Lipinski's five rules. We have taken Osiris program to make a prediction of the toxicity profile (against mutagenic, tumorigenic, irritant, and reproductive effects). The outcome of all parameters indicated a low toxicity risk profile of all compounds except 3d and 3i (found to be irritant. Compounds 3f and 3j displayed high Caco-2 cell permeability with a value of 0.957 and 0.887, respectively. Similarly, it was also noted that compounds 3f and 3j might have a high tendency to cross the BBB with a probability of 0.886 and 0.870, respectively, as comes from in-silico treatment.

The potency of the molecule to inhibit the hERG-K<sup>+</sup> channel may lead to heart arrhythmia and potentially death in the primary stage of the easy drug development process. We have studied external predicative binary and multiclass models of the molecule using "Pred-hERG"<sup>35-37</sup>. We saw positive (blocker) response for 3e and 3h molecules, whereas all other molecules showed a negative (non-blocker) response

All new synthesised molecule shows no any activeness towards MATE, OCT2 inhibition, CYP2D6 substrate, and UGT catalysed reaction.

All newly synthesized compounds were predicted to show non-skin sensitization. Compound 3i was found to be a non-toxic molecule, and all others showed oral toxicity from the in-silico model. The human colon adenocarcinoma cell line (Caco-2) is used to design good intestinal absorption and defensive properties of the mucosa of the intestine. Each molecule showed good in silico intestinal absorption (HIA% % was> 30%). Ames mutagenesis test is used to identify by revert mutants and mutagenicity of environmental samples <sup>38-39</sup>.

All our synthesised compounds did not exhibit acute inhalation toxicity and dermal toxicity against OECD TG 403,436 and OECD TG 402, respectively, as checked in silico rat models. All compounds presented inactiveness towards MATE 1, OCT2 inhibition, CYP2D6 substrate, and UGT catalysed reactions. Compounds 3a to 3j were predicted to have skin non-



sensitization. Compound 3i was found to be Non-toxic against OECD TG 401, 420, 423, and 425 of rats. All the compounds showed eye irritation and corrosion toxicity against OECD TG 405 of the rabbit. Molecules 3a and 3i showed human oral toxicity as predicted from in-silico models. The human colon adenocarcinoma cell line (Caco-2) is used to design good intestinal absorption. Ames mutagenesis test is used to identify revertant mutations and mutagenicity of environmental samples. It is also used to detect suitable mutants. Compounds (mutant) 3b,3c, 3d, and 3e were also observed to be positive in the Ames mutagenesis test. All the compounds displayed Ames's test probability within the range 0.50-0.6. All the molecules were predicted to be non-nephrotoxic. Only 3f shows respiratory toxicity. All the molecules were observed for moderate acute oral toxicity and class III toxicity as calculated from in-silico results. All the compounds were found to be in silico non-carcinogenic. Moreover, we also noticed the fact that our compounds might have a mitochondrial subcellular localization profile. Compounds 3a TO 3j did not display any honey bee toxicity and also showed fish aquatic non-toxicity, except compound 3i.

#### **CONCLUSION:**

The synthesized pyrazole derivatives exhibited promising structural and pharmacokinetic properties. ADMET-SAR analysis indicated good absorption and low toxicity profiles, supporting their potential as drug candidates. Biological assays demonstrated moderate to good antituberculosis activity, correlating with their predicted pharmacokinetics. Compound 3b (MIC value 12.5 µg/ml) did not show any oral toxicity and was found to be most active against Mycobacterium tuberculosis (H37 RV)<sup>40-42</sup>. We get to know from the in-silico test, all were non-carcinogenic. However, further in vivo studies are essential to confirm efficacy and safety. Further studies, including in vivo evaluations and mechanism-of-action investigations, are recommended to advance these molecules as therapeutic candidates.

### **CONFLICTS OF INTEREST**

None

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EVALUATION OF ANTITUBERCULAR AND ANTI FUNGAL ACTIVITY OF SOME NOVEL 6-(4-SUBSTITUTED ARYL)-2-(3,5-DIMETHYL-1H-PYRAZOL-1-YL) IMIDAZO[2,1-B] [1,3,4] THIADIAZOLE DERIVATIVES [Internet]. Citeseer. 2013 [cited 2020 Sep 20]. Available from: http://citeseerx.ist.psu.edu/viewdoc/download?doi=10.1.1.837.4315&rep=rep1&type=pdf

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Synthesis, Characterization, Anti-mycobacterial evaluation, In-Vitro and Computational analysis of Pyrazole derivative

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