

## DESIGN AND SYNTHESIS OF BENZHYDROL DERIVATIVES AS ANTITUBERCULOSIS AGENTS



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Title	Design and Synthesis of Benzhydrol Derivatives as Antituberculosis
	Agents
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MISS PITIKAN KANJANAPRUK : DESIGN AND SYNTHESIS OF BENZHYDROL DERIVATIVES AS ANTITUBERCULOSIS AGENTS THESIS ADVISOR : ASSISTANT PROFESSOR SATHIT NIRATISAI, Ph.D.

Benzhydrol derivatives were designed, synthesized, and evaluated for antituberculosis activities. Benzhydrol derivatives (compounds 4a-f, 9, 10, 13a-c) and its diacetylated derivatives (compounds 5a, 5c, 5f), which are analogs of the natural product 1'-acetoxychavicol acetate (ACA) from Alpinia galangal rhizome, were designed as new antituberculosis agents. In this synthetic approach, compounds 4a-f, 9 were synthesized in three steps, i.e. (1) coupling reactions between benzoic acid or substituted benzoic acids and phenol to form ester linkages, (2) Fries rearrangement of phenylbenzoates using aluminium chloride as a catalyst to get benzophenones, and (3) reduction of ketones with sodium borohydride to obtain benzhydrol compounds. Compounds 4a-f and 9 had overall yields of 20.26%, 26.98%, 56.59%, 49.71%, 31.41%, 23.97%, and 40.54%, respectively. Benzhydrol 10 was prepared by amide hydrolysis of compound 4f with a 6.57% overall yield. By acetylating 4a, 4c, and 4f with acetic anhydride, the diacetylated derivatives 5a, 5c, and 5f were obtained with overall yields of 18.82%, 39.22%, and 21.96%, respectively. The amide derivatives 13a-c were prepared: (1) amide formation between 4-aminobenzophenone and alkyl, aryl, or long-chain aryl acids, and (2) ketone reduction. Compounds 13a-c had overall yields of 57.53%, 29.74%, and 24.26%, respectively. The structures of desired compounds were elucidated by spectral data (<sup>1</sup>H NMR, <sup>13</sup>C NMR and IR). These synthesized benzhydrol compounds were evaluated for antituberculosis activities by agar-dilution method. It was found that they had antituberculosis efficacy on 20 clinical isolates and MTB H37Rv reference strains (ATCC 27294). Compound 4d, 4hydroxy- $\alpha$ -(4'-butylphenyl)benzyl alcohol possessed the highest inhibitory activity in this series with an MIC value of 20-40 µg/ml. The structure-activity of the synthesized benzhydrol derivatives were also summarized.

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# **TABLE OF CONTENTS**

ABSTRACT	D
ACKNOWLEDGEMENTS	E
TABLE OF CONTENTS	F
LIST OF TABLES	I
LIST OF FIGURES	J
CHAPTER 1 INTRODUCTION	1
1.1 Statement and significance of the research problem	1
1.2 Objectives of the study	3
1.3 Hypotheses of the study	3
1.4 Scope of the study	5
CHAPTER 2 LITERATURE REVIEW	8
2.1 Tuberculosis	8
2.1.1 Tuberculosis Overview	8
2.1.2 Causes of Tuberculosis	9
2.2 Anti-tuberculosis drugs	11
2.2.1 Targets and mode of action of current anti-TB drugs	12
2.3 1'-S-Acetoxychavicol acetate (ACA)	15
2.3.1 ACA modification for anti-TB activity	15
2.3.2 Stability of ACA	17
2.4 Chemical synthesis	20
2.4.1 Benzhydrol synthesis	20
2.4.2 Benzophenone synthesis	21
2.4.3 Fries rearrangement reaction	22
2.4.4 Reduction	24
2.5 Chemical structure of anti-TB agents	25

CHAPTER 3 MATERIALS AND METHODS	30
3.1 Materials	30
3.1.1 Instruments	30
3.1.2 Chemical reagents and Instruments	30
3.1.3 Chemical preparations	32
3.2 Methods	33
3.2.1 General methods for chemistry	33
3.2.2 Synthesis of designed compounds	35
3.2.2.1 Synthesis of phenyl benzoate derivatives (2a-f)	35
3.2.2.2 Synthesis of 4-hydroxy benzophenone derivatives (3a-f)	37
3.2.2.3 Synthesis of (4-Hydroxyphenyl)phenylmethanol derivatives (4-Hydroxyphenylmethanol derivatives (4-H	4a-f)
	39
3.2.2.4 4-acetoxy-α-(phenyl)benzyl acetate derivatives (5a, 5d, 5g)	42
3.2.2.5 2-Naphthyl benzoate (7)	43
3.2.2.6 4-hydroxyphenyl(2-naphthyl)methanone (8)	44
3.2.2.7 4-hydroxyphenyl(2-naphthyl)methanol (9)	44
3.2.2.8 4-((4-aminophenyl)(hydroxy)methyl)phenol (10)	45
3.2.2.9 4-amido benzophenone derivatives (12a-c)	45
3.2.2.10 4-amido benzhydrol derivatives (13a-c)	47
3.3 Evaluation of antitubercular activities	48
CHAPTER 4 RESULTS AND DISCUSSION	50
4.1 Chemical synthesis	52
4.1.1 Phenyl benzoate compounds (2a-f)	52
4.1.2 Benzophenone compounds (3a-m)	54
4.1.2.1 Fries rearrangement reaction investigation by DSC analysis	54
4.1.3 Benzhydrol compounds (4a-f) and 9	59
4.1.4 Diacetyl benzhydrol compounds (5a, 5c, 5f)	60
4.1.5 4-hydroxy-α-(4'-amino phenyl)benzyl alcohol (10)	60
4.1.6 Amidobenzhydrol compounds (13a-c)	61

4.2 Structural characterizations $\epsilon$	54
4.2.1 IR spectroscopic characterization	54
4.2.2 NMR spectroscopic characterization	59
4.3 Structure-activity relationship of synthesized compounds and anti-tuberculosis activity	; 74
CHAPTER 5 CONCLUSION	78
REFERENCES	31
APPENDIX	38
VITA	17



# LIST OF TABLES

# Page

Table 1 Synthesized benzhydrol derivatives	6
Table 2 Anti-TB drugs and their targets	13
Table 3 % Compositions of Fries rearrangement products by LC-MS analysis at         variable temperatures	56
Table 4 The reaction conditions and percent yields of compounds 3a-f and 8	59
Table 5 The physical properties and percent yield of benzhydrol derivatives	63
Table 6 Characteristic IR bands of important functional groups of synthesized         compounds	68
Table 7 In vitro antitubercular activities of synthesized compounds against MTB <sup>a</sup> .	77



# LIST OF FIGURES

# Page

Figure 1 Chemical structure of 1'S-1'-Acetoxychavicol acetate (ACA)1
Figure 2 Design of benzhydrol derivatives
Figure 3 Proposed benzhydrol derivatives
Figure 4 Design strategy used for benzhydrol derivatives
Figure 5 The Mycobacterial cell wall structure10
Figure 6 Mycolic acids in MTB11
Figure 7 ACA and its analogs possessing anti-TB activities16
Figure 8 Hydrolysis of 1'-Actetoxychavicol acetate17
Figure 9 Proposed sigmatropic rearrangement of 1'-Acetoxychavicol acetate18
Figure 10 Chemical structure of 1'-Acetoxychavicol acetate (ACA), benzhydrol and benzhydrol diacetate
Figure 11 Diarylmethane and diarylmethanone derivatives possessing anti-TB activities
Figure 12 Synthesis of benzhydrol analogs
Figure 13 Synthesis of benzhydrol and acetoxybenzhydrol analogs
Figure 14 Synthesis of benzophenone derivatives
Figure 15 Mechanism of Fries rearrangement reaction
Figure 16 Mechanism of ketone reduction by NaBH <sub>4</sub> 25
Figure 17 Structure of amino containing anti-TB compounds25
Figure 18 Structure of amide containing anti-TB compounds26
Figure 19 Structure of halogen containing anti-TB compounds27
Figure 20 Structure of cinnamyl containing anti-TB compounds
Figure 21 Structure of long chain containing anti-TB compounds29
Figure 22 The scope of research project
Figure 23 The reaction mechanism of the esterification
Figure 24 DSC curves of the mixtures of compounds 2b, 2e, 2f and AlCl <sub>3</sub> 55

Figure 25 The reaction mechanism of Fries rearrangement.	.58
Figure 26 The reaction mechanism of sodium borohydride reduction	.60
Figure 27 The reaction mechanism of the base-catalyzed hydrolysis	.61
Figure 28 The reaction mechanism of the amide formation	.62
Figure 29 The structure of active benzhydrol derivatives of this series	.80



## CHAPTER 1 INTRODUCTION

#### 1.1 Statement and significance of the research problem

In recent years, tuberculosis (TB) remains as one of the ten leading causes of death worldwide in 2021.[1] There are an estimated 10.4 million new incident of TB annually, and the number of deaths is around 1.8 million. The high susceptibility of human immunodeficiency virus-infected persons to the disease, the emergence of multi-drug-resistant (MDR-TB) strains and extensively drug-resistant (XDR-TB) ones have brought this infectious disease into the focus of urgent scientific interest. However, the development of new anti-TB drugs has been slow, therefore, importantly efforts to discover new classes of chemical compounds acting with different mechanisms from currently used.

1'S-1'-Acetoxychavicol acetate (ACA) (Figure 1), the major constituent isolated from *Alpinia galangal* rhizome, and its derivatives were reported to be efficacious antituberculosis agents. ACA was studied against *M. tuberculosis* H37Ra and *M. tuberculosis* H37Rv strains with minimum inhibitory concentrations (MIC) 0.1-0.5  $\mu$ g/mL and 0.6-1.6  $\mu$ g/mL, respectively.[2] ACA is reasonably a lead compound for the development of new potent anti-TB drugs. However, ACA still has a few disadvantages i.e. poor water solubility and degradation in aqueous solution. Because of unstable structure, it is hard to develop ACA as edible drugs and modification of chemical structure is needed to improve stability. Thus, while the possibility to develop ACA itself as an oral anti-TB drug may be remote, synthetic compounds that contain the ACA pharmacophore were considered to be viable lead.



Figure 1 Chemical structure of 1'S-1'-Acetoxychavicol acetate (ACA)

To avoid the instability of ACA, our attempt began to investigate the replacement effect of the vinyl on ACA with phenyl, as shown in Figure 2. In a review study, benzhydrol diacetate analog was reported as a stable analog of natural ACA.[3] Therefore, in this research, benzhydrol derivatives were designed to obtain promising stable compounds, synthesized, and tested for anti-TB activity. Benzhydrol was selected as a scaffold for designing of proposed compounds. To further understanding the SAR of the benzhydrol derivatives, substitution at aromatic ring A and B was explored. The benzydrol derivatives were designed and synthesized into three groups. 1) 4-Hydroxybenzhydrols containing the para- position of ring A was substituted with hydroxyl group. The acetamido substitution at ring B was substituted at meta- or para- to compare the effects of substitution positions on anti-TB activities. At the para- position of ring B, various functional groups, including halide, amino, amide, aromatic, and aliphatic groups, were attached. 2) Benzhydrol diacetates were designed to mimic the ACA structure to evaluate the acetoxy group's necessity. The di-hydroxy compounds that show effective activity were converted to di-acetoxy compounds to investigate their anti-TB activities further. 3) 4-Amidobenzhydrols were generated from the coupling between benzhydrol and the alkyl, aryl, or cinnamyl to explore the molecular target. (Figure 3) In addition, interesting intermediate compounds, benzophenone derivatives were tested for anti-TB activity.



Figure 2 Design of benzhydrol derivatives



Figure 3 Proposed benzhydrol derivatives

### 1.2 Objectives of the study

The goal of the study is to design and synthesize benzhydrol derivatives as antituberculosis agents. Objectives of research in order to achieve this goal are 1. To design and synthesize benzhydrol derivatives 2. To evaluate antitubercular activities of the synthesized benzhydrol derivatives

3. To determine the structure-activity relationship (SAR) of the synthesized

benzhydrol derivatives after evaluation of MIC values against M.tuberculosis strains

#### 1.3 Hypotheses of the study

A literature survey described that both ACA and diphenyl compounds possess anti-TB activity with good MIC values on *M. tuberculosis* strains. [2, 4-6] The two structural fragments, i.e., 1'-hydroxychavicol obtained from ACA and diphenyl rings, were structurally combined to generate benzhydrol derivatives, which can be employed to explore target sites. (Figure 4). The benzhydrol derivatives were modified as follows:

1. Substitution of halogen atoms, fluorine and bromine, at *meta-* or *para*positions of the benzene ring (R3) may enhance anti-TB activity due to attractive interaction between the high electronegativity nature of halogen atoms with electrophilic regions of target sites. The different sizes of halogen atoms may describe the size of the binding sites of biological targets. 2. Substitution at R3 with various chemical groups, e.g., alkyl long chain or replacing benzene ring with naphthalene ring (compound 9), may also give information on the size and shape of binding sites as in hydrophobic pockets of the targets.

3. Amide substitution (R3) may be employed, assuming that its ionization giving positive ions at physiological pH may attract negative ions of target binding sites. In addition, the amino group can be coupled with various side-chain acids, e.g., aliphatic acid, aromatic acid, or cinnamic acid, by forming an amide linkage to explore the interaction of amide side chains with biological targets.

4. Acetoxy substitution at  $\alpha$ -carbon (R1) and benzene ring (R2) positions may increase biological activity with promising acetyl groups as essential substituents similar to that of the lead compound (ACA).



Figure 4 Design strategy used for benzhydrol derivatives

#### **1.4 Scope of the study**

A set of new analogs were designed and synthesized to evaluate the structural requirements for the antituberculosis activity. Benzhydrol derivatives prepared from organic synthesis are shown in Table 1. All synthesized compounds were purified by chromatography. Percent yields of products were calculated for each purified compound. Chemical structures were elucidated by using spectroscopic methods i.e. Fouirer Transform Infrared Spectroscopy (FTIR), Proton Nuclear Magnetic Spectroscopy (<sup>1</sup>H-NMR), Carbon-13 Nuclear Magnetic Spectroscopy (<sup>13</sup>C-NMR), and Mass Spectroscopy (MS). Then, all compounds were subjected for anti-TB testing. Agar dilution method were used in order to evaluate MICs of synthesized compounds. For the SAR study, the presence of each functional group of benzhydrol derivatives that induce antitubercular activity may potentially be explained by the interaction between ligands and possible target sites.



Entry	Compound	Chemical structures Molecul		Molecular
	code		formula	weight
1	<b>4</b> a	OH	$C_{13}H_{12}O_2$	200.23
		но		
2	4b	OH	$C_{13}H_{11}FO_2$	218.22
		HOFF		
3	4c	OH HO Br	C <sub>13</sub> H <sub>11</sub> BrO <sub>2</sub>	279.13
4	4d	ОН	C <sub>17</sub> H <sub>20</sub> O <sub>2</sub>	256.34
	à	HO 3		
5	4e	OH H N CH <sub>3</sub>	C <sub>15</sub> H <sub>15</sub> NO <sub>3</sub>	257.28
6	4f	HO OH HO HO HO CH <sub>3</sub>	C <sub>15</sub> H <sub>15</sub> NO <sub>3</sub>	257.28
7	9	НО	C <sub>17</sub> H <sub>14</sub> O <sub>2</sub>	250.29
8	10	HO NH <sub>2</sub>	C <sub>13</sub> H <sub>13</sub> NO <sub>2</sub>	215.25

 Table 1 Synthesized benzhydrol derivatives

Entry	Compound	Chemical structures	Molecular	Molecular	
	code		formula	weight	
9	5a	Aco QAc	C <sub>17</sub> H <sub>16</sub> O <sub>4</sub>	284.31	
10	5d	AcO Br	C <sub>17</sub> H <sub>15</sub> BrO <sub>4</sub>	363.20	
11	5f	Aco CH <sub>3</sub>	C19H19NO5	341.36	
12	<b>13</b> a	OH OH NH CH <sub>3</sub>	C <sub>15</sub> H <sub>14</sub> NO <sub>2</sub>	240.28	
13	13b	OH N H H	C <sub>20</sub> H <sub>17</sub> NO <sub>2</sub>	303.35	
14	13c	OH O N H	C <sub>22</sub> H <sub>19</sub> NO <sub>2</sub>	329.39	

### CHAPTER 2 LITERATURE REVIEW

#### 2.1 Tuberculosis

#### 2.1.1 Tuberculosis Overview

Tuberculosis (TB) is a threat to worldwide public health, mainly caused by Mycobacterium tuberculosis (MTB) bacteria species. In 2021, an estimated 10.4 million people fell ill with TB worldwide and the largest number of new TB cases occurred in the South-East Asian region with 44% of new cases. TB has led to an estimated death of around 1.3 million including human immune deficiency virus (HIV) positive people in 2021 and is the second leading cause of deaths among the infectious disease [1]. With its high death rate, tuberculosis remains a difficult disease to treat. There are other forms of tuberculosis that affect the kidney, brain, lungs, etc., but mycobacteria mostly damage the lungs because MTB, an aerobic mycobacterium, receives a great deal of oxygen from the lungs. Due to the emergence of MDR-TB, XDR-TB, and TDR-TB, the directly observed treatment short therapy (DOTS) with the first line anti-TB medications isoniazid (INH), rifampin (RIF), ethambutol (EMB), and pyrazinamide (PZA) is no longer sufficient to combat this lethal disease. [7-9]. Bedaquiline recommended by United States Food and Drug Administration (USFDA) [10] and delamanid approved by European Medical Agency (EMEA) [11] for treatment of MDR-TB have shown cases of resistance. In addition to QT prolongation, bedaquiline and delamanid cause hepatotoxicity [12] and CNS toxicity [13, 14], respectively. There have been reports that long-term treatment with bedaquiline is related with cardiac arrhythmia as a fatal side effect [15]. Recently, USFDA has approved pretomanid, a nitroimidazopyran derivative, in conjunction with bedaquiline and linezolid for XDR-TB or MDR-TB to treat adult TB patients [16]. Due to the deteriorating tuberculosis condition, there has been an urgent need for the discovery of novel antitubercular agents. Poor chemotherapeutics, crossresistance, and the dearth of effective anti-TB medications highlight the need for the discovery of innovative anti-tuberculous agents that may have a novel mode of action. Novel anti-TB drugs with benzhydrol as its pharmacophoric characteristic are described here.

#### 2.1.2 Causes of Tuberculosis

Tuberculosis is caused by the bacillus Mycobacterium tuberculosis (MTB). MTB is a rather large, non-motile, rod-shaped bacterium, measuring 2 to 4 µm in length and 0.2 to 0.5  $\mu$ m in width, and dividing every 16 to 20 hours, which is a very slow rate compared to other bacteria. MTB is neither Gram-positive nor Gramnegative because it lacks the chemical characteristics of either category. Its cell wall possesses both Gram-positive and Gram-negative features.[17] They are categorized as acid-fast Gram-positive bacteria (AFB) because they lack an outer cell membrane and their fatty cell walls prevent acid solutions from decolorizing the cells during diagnostic staining. The cell wall structure of MTB deserves special attention because it appears to allow MTB to survive in its preferred environment. It is a major determinant of virulence for the bacterium. The cell walls of all Mycobacterium species are thicker than those of most other bacteria, being hydrophobic, waxy, and abundant in mycolic acids/mycolates. The cell wall consists of the hydrophobic mycolate layer and a peptidoglycan layer held together by a polysaccharide, arabinogalactan.[18, 19] The arabinogalactan and mycolic acid layer is covered by a polypeptide layer. Other glycolipids include phosphatidyinositol mannosides and lipoarabinomannan. Porins are involved in the movement of tiny hydrophilic molecules through the cell wall's outer membrane. (Figure 5)

*นั้นว่าม*ยาลัยศิลปาก



Source: From http://student.ccbcmd.edu/courses/bio141/lecguide/unit1/prostruct/u1fig11.html

## Figure 5 The Mycobacterial cell wall structure

Lipids make up more than sixty percent of the mycobacterial cell wall. Mycolic acids, cord factor, and wax-D are the three primary components that make up the lipid fraction of the MTB cell wall [20, 21]. Mycolic acids impart *M. tuberculosis* with unique properties that defy medical treatment. They increase an organism's resistance to chemical damage and dehydration and reduce the efficacy of hydrophilic antibiotics and biocides [22].

Mycolic acids are high-molecular weight fatty acid. They are composed of a longer beta-hydroxy chain with a shorter alpha-alkyl side chain. Each molecule contains between 60 and 90 carbon atoms. Three main types of mycolic acids are alpha-, methoxy-, and keto-mycolic aicd. Alpha-mycolic acids constitute at least 70% of the organism's mycolic acids and include several cyclopropane rings. Between 10 and 15 percent of the mycolic acids in an organism are methoxy-mycolic acids, which contain several methoxy groups. 10% to 15% of the remaining mycolic acids are keto-mycolic acids, which contain several ketone groups (Figure 6).



Source: From https://pl.csl.sri.com/mycolate-overview.html

Figure 6 Mycolic acids in MTB

### 2.2 Anti-tuberculosis drugs

In 1944, chemotherapy for tuberculosis was initiated using the natural substance streptomycin. Then, in the early 1950s, the alternative synthetic chemicals isoniazid and pyrazinamide were discovered, followed by ethambutol and rifampicin in 1961 and 1963, respectively.[23, 24] After the approval of rifampicin in 1967, no new anti-tubercular drugs were developed until the introduction of bedaquiline as an anti-tubercular medicine in 2012. There was a forty-year gap in identifying innovative drugs to treat this disease. TB is a priority disease for the discovery and development of novel, safe compounds because MDR and XDR strains are emerging at a rapid rate. It is synergistic of the HIV pandemic, pharmacokinetic interaction between anti-TB and anti-retroviral drugs, complex regimens for MDR-TB, longer treatment duration, relapse, and toxic side effects.[25] The first-line drugs, including isoniazid, rifampin,

ethambutol and pyrazinamide, are the most effective and have the lowest toxicity. The second-line drugs are less effective and have more toxic effects than first-line drugs. These drugs may not be available in many developing countries. There are streptomycin, *p*-amino salicylic acid, ciprofloxacin, and moxifloxacin. The third-line drugs are Amikacin, linezolid, clarithromycin, kanamycin, cycloserine, and capreomycin. These drugs were categorized as "third-line treatments" because they are not highly effective or their efficacy has not been demonstrated.

Novel drugs required to combat tuberculosis must contain bactericidal activity against both drug-susceptible and drug-resistant MTB strains and be effective in real-world settings. In general, novel and safe anti-TB drugs aim to decrease treatment duration, develop new mechanisms of action against drug-resistant strains, be effective against MDR-TB and XDR-TB, prevent drug-drug interactions, and reduce toxic side effects. In addition, new anti-tuberculosis medications must be oral-only regimens in adults, children, and HIV patients. Their posology must be once daily or less frequently for treatment course tolerability. The next challenge is to be cost competitive with current drugs.

### 2.2.1 Targets and mode of action of current anti-TB drugs

Current standard treatment for tuberculosis (TB) involves drugs that interfere with bacterial metabolism, particularly those that prevent cell wall production. Cell wall inhibitors include isoniazid, ethambutol, ethionamide, and cycloserine; nucleic acid synthesis inhibitors include rifampicin and quinolones; protein synthesis inhibitors include streptomycin and kanamycin; and second-line TB medications include those that block the metabolism of membrane energy (pyrazinamide) [26]. New drugs have emerged recently as potential candidates for the treatment of TB. The new anti-tuberculosis agents are targeting mycolic acid biosynthesis, protein biosynthesis, cofactor biosynthesis, menaquinone biosynthesis, ATP biosynthesis, regulatory proteins, and the stringent response enzyme [27]. Targets and mechanisms of action of current TB drugs are summarized in Table 2.

Table 2 Anti-TB	drugs	and	their	targets
-----------------	-------	-----	-------	---------

No.	Drug	Chemical class	Mechanism of action	MIC (µg/ml)	Ref.
1	Isoniazid	Isonicotinic acid derivative	Inhibits cell wall mycolic acid biosynthesis	0.02-020	[26]
2	Rifampicin	Amide derivatives	Inhibits of RNA synthesis	0.05-0.50	[26]
3	Pyrazinamide	Niacinamide derivative	Depletion of membrane energy	5.5-6.0	[26]
4	Ethambutol	Ethylenediamine	Inhibits cell-wall arabinogalactan synthesis	1.0-5.0	[26]
5	Streptomycin	Amino- glycoside	Inhibits protein synthesis	1.0-8.0	[26]
6	Kanamycin	Amino- glycoside	Inhibits protein synthesis	1.0-4.0	[26]
7	Amicacin	Semi-synthetic aminoglycoside	Inhibits protein synthesis	1.0-4.0	[26]
8	Capreomycin	Cyclic peptide	Inhibits protein synthesis	2.0-4.0	[26]
9	Quinolones	Fluroquinolone	Inhibits DNA replication and transcription	0.2-4.0	[28]
10	Ethionamide	Isonicotinic acid derivative	Inhibits mycolic acid synthesis	2.5-10.0	[26]
11	Prothionamide	Thioamide derivative	Disrupts cell wall biosynthesis	0.5	[29]
12	Para-amino salicylic acid	Salicylic acid	Inhibits folate biosynthesis	0.50-8.0	[26]
13	D-cycloserine	D-analine analogue	Inhibits peptidoglycan biosynthesis in the cell wall	1.5-40.0	[26]
14	Rifabutin	A rifamycin class	Inhibits RNA synthesis	0.015 - 0.125	[30]
15	Rifapentine	Cyclopentyl rifampicin	Inhibits RNA synthesis	0.06-0.5	[31]

No.	Drug	Chemical class	Mechanism of action	MIC (µg/ml)	Ref.
16	Clofazimine	A lipophilic riminophena-zine antibiotic	Probably acts on membrane transport	0.01- 0.25	[26]
17	Bedaquiline	Diarylquinoline	Inhibits ATP synthesis	0.03- 0.12	[32]
18	Delamanid	Dihydronitro imidazo oxazole derivative	Probably inhibiting cell wall	0.006- 0.024	[33]
19	Linezolid	Oxazolidinone	Inhibits protein synthesis	0.06-1.0	[34]
20	Thioaceta- zone	Thiosemi- carbazone	Inhibits cell wall synthesis	0.1	[35]
21	Pretomanid	Nitroimidazole	Probably inhibition of cell wall synthesis and causing respiratory	0.015- 0.25	[33]
22	Tedizolid	Oxazolidinone	Inhibits protein synthesis	0.015– 16	[36]
23	Sutezolid	Oxazolidinone	Inhibits protein synthesis	≤ 0.062	[37]
24	SQ 109	1,2- ethylenediamine	Inhibits cell wall synthesis	0.7-1.56 (µM)	[38]
25	TBA-354	Pyridine biaryl nitroimidazole	Inhibits cell wall synthesis and causing respiratory poisoning	0.011	[39]
26	CPZEN-45	Caprazamycin derivative	Inhibits cell wall biosynthesis	1.56	[40]
27	Q203	Imidazopyridine amide	Oxidative phosphorylation	10 nM	[41]
28	SQ641	Carboxamide	Inhibits cell wall biosynthesis	1.0	[42]

**Table 2** Anti-TB drugs and their targets (continued)

#### 2.3 1'-S-Acetoxychavicol acetate (ACA)

1'-S-Acetoxychavicol acetate (ACA) is the major constituent isolated from *Alpinia galangal* (Kha) rhizome. Greater galangal is also known as *Alpinia galangal* (L.) Sw. (Zingiberaceae), and it is a perennial herb that has rhizomatous root stocks and tall stems that are covered with leaves. The plant has been discovered in the Western Ghats, Mysore, Goa, Malabar, and Gujarat, as well as Thailand, Indonesia, China, and Malaysia. ACA was reported to possess activities as antioxidant, anticancer, antifungal, antimicrobial, anti-inflammatory, gastroprotective agent, anti-HIV, and anti-TB.[2, 3, 43-58]

For anti-TB activity, ACA was studied against *M. tuberculosis* H37Ra and *M. tuberculosis* H37Rv strains with minimum inhibitory concentrations (MIC) 0.1-0.5  $\mu$ g/mL and 0.6-1.6  $\mu$ g/mL, respectively.[2] In addition, ACA was reported that it could inhibit efflux pump in *M. smegmatis* mc2 cells.[44] This is highly advantageous because inhibition of efflux pump can decrease the resistance of bacteria to antibiotic and reduce the frequency of the emergence of resistant mutant strains, while the toxic level of ACA is low in various mammalian cells with higher MICs than the MIC against M.tb H37Ra.[2]

# 2.3.1 ACA modification for anti-TB activity

Due to the various advantages of ACA, it is a likely candidate for creating new effectively anti-tuberculosis drugs. Bunthitsakda W. reported the study of anti-TB of ACA analogs to clarify the structure–activity relationship for drug design. There were four compounds (I, II, III, and IV) showed potent anti-TB activity against *M. tuberculosis* H37Ra (Figure 7). This report showed that the presence of two acetyl groups replacing hydroxyl groups did not significantly affect anti-TB activity.[43] Electron-withdrawing halogen substitutions at the *para-* and *ortho-* position of benzene ring of compounds (I, II) showed high inhibitory activity against *M. tuberculosis* H37Ra. The presence of the aliphatic side chain at C1' of chavicol of compounds (III, IV) was important for improving activity.[59] This research informed that the essential anti-tuberculosis structure of ACA analogs should contain the following structural requirements: 1) the *para-*substitution of halide atom at the benzene ring was required; 2) the presence of aliphatic side chain at C1' of chavicol should have fewer than 8 carbons; 3) The presence of an acetyl group rather than a

hydroxyl group at the C1' position of an active molecule (with an eight-carbon aliphatic side chain) was crucial for enhancing its biological activity; 4) A phenyl group connected to a carbon atom with a highly electronegative group at the *para*-position of the benzene ring, such as a chloride or bromide atom, possesses the physical properties necessary to enter the target cell more effectively than other aromatics. [43] The MICs of *S*-ACA against *M. tuberculosis* H37Ra and H37Rv were lower than those of the racemic ACA. Both *S*- and *rac*-ACA had potent bactericidal activity and were able to kill the *M. tuberculosis* H37Rv strain.[4] The structural requirements of ACA for displaying activity have described that phenyl acetate moiety was important, substitution of vinyl group by phenyl still possessed the activity and the chirality was not influential factor for eliciting the response. [46]



S-ACA MIC H37Ra = 0.2 μg/ml rac-ACA MIC H37Ra = 0.5 μg/ml Rifampicin MIC H37Ra = 0.005 μg/ml

S-ACA MIC H37Rv = 0.7 μg/ml rac-ACA MIC H37Rv = 2.7 μg/ml Rifampicin MIC H37Rv = 0.1 μg/ml



INH MIC = 0.023-0.046 µg/ml

Figure 7 ACA and its analogs possessing anti-TB activities

#### 2.3.2 Stability of ACA

In an aqueous solution, ACA lacked stability. After a few hours at room temperature, ACA undergoes hydrolysis and structural rearrangement. Three degraded products were identified as 1'-hydroxychavicol acetate, *p*-coumaryl diacetate, and *p*-acetoxycinnamic alcohol (Figure 8) [60]. It can also be postulated that ACA is metabolically unstable compound. As hydrolysis products, *p*-acetoxycinnamic alcohol is the result of ACA's SN1 reaction mechanism, and *p*-coumaryl diacetate is the consequence of [3,3]-sigmatropic isomerization (Figure 9) [62]. Kubota K. and co-workers reported the antioxidant activity of ACA and its related compounds from *A. galanga* fresh rhizomes imported from Thailand. They heated rhizomes in aqueous medium or lard and antioxidant assay was investigated by ferric thiocyanate and thiobarbituric acid (TBA) method. It was found degraded compounds of *A. galanga* after heating in aqueous medium or lard [61]. As a result, the clinical applications of ACA as an antituberculosis drug are limited by some restrictions.



p-Acetoxy cinnamic alcohol

Figure 8 Hydrolysis of 1'-Actetoxychavicol acetate.



1'-Acetoxychavicol acetate (ACA)

p-Coumaryl diacetate

Figure 9 Proposed sigmatropic rearrangement of 1'-Acetoxychavicol acetate.

Because ACA has a few disadvantages, i.e. poor water solubility and degradation in aqueous solution, it is hard to develop ACA as edible drugs and chemical structure modification is needed to improve stability. While the possibility of developing ACA as an oral anti-TB medication is remote, synthesized molecules containing the ACA pharmacophore may be a feasible lead. Benzhydrol derivatives were designed as stable ACA analogs. The vinyl group of ACA was substituted by the phenyl group, converting to benzhydrol diacetate (Figure 10). Benzhydrol derivatives were reported to possess antiallergic and anticancer activities. [3, 46] It has been proved that these derivatives are stable and become promising lead compounds for anti-TB activity for this study.



Figure 10 Chemical structure of 1'-Acetoxychavicol acetate (ACA), benzhydrol and benzhydrol diacetate

Due to the advantages of diaryl containing benzhydrol scaffold, design and synthesis of benzhydrol derivatives were investigated in several studies. Furthermore, diarylmethane and diarylmethanone derivatives with promising antitubercular activities were also well documented (Figure 11). [4-6]



Figure 11 Diarylmethane and diarylmethanone derivatives possessing anti-TB activities

From literature reviews [4-6], it was found that most of compounds containing diaryl rings with nitrogen atom and halogen atom showed effective anti-TB activity. The electron donating and electron withdrawing groups of nitrogen and halogen atoms show a significant effect on anti-TB activity. However, the mechanism of action of these compounds on *M. tuberculosis* is unknown and there are only few SAR studies of ACA and benzhydrol derivatives on anti-TB activity. Hence, it is of great interest to discover novel compounds with high potency for *M. tuberculosis* inhibition. Based on the structures of ACA and benzhydrol, we attempted to design and synthesis a series of benzhydrol derivatives and determine the SAR of novel benzhydrol derivatives against *M. tuberculosis* H37Rv activities.

#### **2.4 Chemical synthesis**

#### 2.4.1 Benzhydrol synthesis

Benzhydrol or diphenylmethanol can be synthesized by the reduction of benzophenone or by a Grignard reaction between phenyl magnesium bromide and benzaldehyde. The synthesis pathway of benzhydrol type 1'-acetoxychavicol acetate (ACA) analogs was reported by Misawa et al. [3] The hydroxyl group of 4-hydroxybenzaldehyde was protected with *tert*-butyldimethylchlorosilane (TBS-Cl). The intermediates were then made by treating 4-hydroxybenzaldehyde with the right Grignard or aryl lithium reagents. The TBS group was deprotected with *tetra-n*-butylammonium fluoride (TBAF), and acetylation with acetic anhydride yielded benzhydrols and their acetyl analogs. On the other hand, the hydroxyl group of 4-bromophenol was protected with TBS-Cl, followed by treatment with n-butyllithium, yielding aryllithium species, which were quenched with different benzaldehydes to yield intermediates. TBS group deprotection with TBAF and subsequent acetylation with acetic anhydride yielded benzhydrols and acetylated compounds (Figure 12).



Figure 12 Synthesis of benzhydrol analogs

Yasuhara et al. synthesized the acetoxybenzhydrols as a potent antiallergic. The benzhydrol analogues were created by protecting the hydroxyl of 4-hydroxy benzophenone with *tert*-butyldimethylsilyl chloride (TBSCl), which resulted in the corresponding silyl ether compound, which was then reduced with sodium borohydride to yield a racemic mixture of TBS-O-benzhydrol. 4-hydroxybenzhydrol was produced by treating protected benzhydrol with tetrabutylammonium fluoride in THF. Under usual conditions, 4-hydroxybenzhydrol was treated with acetic

anhydride, suitable acyl chlorides, or methyl chlorocarbonate to obtain corresponding acetoxybenzhydrols. (Figure 13).



Figure 13 Synthesis of benzhydrol and acetoxybenzhydrol analogs

## 2.4.2 Benzophenone synthesis

According to the benzhydrol synthesis, benzophenone can be reduced to create benzhydrol. In 2012, Kwon and coworkers [62] synthesized the anti-inflammatory activity of a benzophenone derivative by inhibiting IFN- $\gamma$ -induced ICAM-1 expression. Figure 14 depicts the straightforward synthesis of benzoyloxy benzophenone derivatives, which consists of the dicyclohexyl carbodiimide coupling of benzoic acid with phenol, the Fries rearrangement of ester to aryl ketone, and the esterification of benzophenone and acyl chloride to yield benzoyloxy benzophenone. Fries rearrangement and esterification of benzoic acids produced benzoyloxy benzophenones with yields ranging from 24 to 89%.



Figure 14 Synthesis of benzophenone derivatives

#### 2.4.3 Fries rearrangement reaction

The Fries rearrangement is an organic process in which phenolic esters are converted into hydroxyaryl ketones by heating in the presence of a catalyst. Phenolic esters on heating with aluminium trichloride (Lewis acid) give *ortho* and *para* acyl phenol. Solvent is generally not required for the reaction. Utilizing Nitrobenzene, DMF, and Zylene decreases the reaction temperature. Fries rearrangement has also been observed to be catalysed by light (photo Fries rearrangement). Fries rearrangement is well documented. However, the regioselectivity between the formation of 2- or 4-hydroxybenzophenones is still in question because of the versatility of substituents in the structure, which results in a low yield. [63] The *para* isomer is generally generated preferentially at temperatures below 100 C, but above this temperature, the *ortho* isomer predominates. The most commonly employed Lewis acids for the Fries rearrangement reaction are AlCl<sub>3</sub>, FeCl<sub>3</sub>, MsOH, and ZnCl<sub>3</sub> at temperatures between 100°C to 210 °C [62].

Due to the wide ranges of reaction temperatures, Yerande S. and co-workers investigated the various reaction temperatures to obtain the ratio of *ortho-/para*-isomer. Toward this goal, Fries rearrangement was performed using AlCl<sub>3</sub> as a catalyst at various temperatures ranging from 40 °C to 170 °C. They discovered that temperatures below 100°C result in partial conversion, while temperatures above 150°C result in lower isolated yields due to the considerable formation of other side products. Hence all the optimizations were carried out to do Fries rearrangement using 1.5 equiv of AlCl<sub>3</sub> at 120 °C [64].

The precise process of Fries rearrangement remains unclear. Both intermolecular and intramolecular reaction processes are supported by evidence. Therefore, it appears that both mechanisms are acting concurrently. A two-step mechanism may be taken to be Friedel-Crafts acylation in which the acylium ion, (MeCO) is supplied by the substrate, i.e., it is self-acylation. Initially, AlCl<sub>3</sub> binds with the oxygen of the phenoxy group to create the acylium ion. As depicted in Figure 15, the acylium ion attacks the benzene ring. An acyl group of phenol ester migrates to the aryl ring. The reaction is *ortho* and *para* selective, and modifying reaction conditions, such as temperature and solvent, might prefer one of the two products.



Figure 15 Mechanism of Fries rearrangement reaction

In this study, benzhydrol derivatives were synthesized from corresponding phenyl benzoates through Fries rearrangement and reduction reaction. In order to carry out the synthesis using the non-solvent technique, the Fries rearrangement reaction was decided to be used. As Fries rearrangement is an endothermic reaction [65], the reaction mixtures of phenyl benzoates and AlCl<sub>3</sub> heated at constant rates were analyzed by differential scanning calorimetry (DCS) to establish the ideal temperature for phenyl benzoates.

### Differential scanning calorimetry (DSC) analysis

Differential scanning calorimetry (DSC) is a thermal analytical measuring how a sample's physical or chemical properties change, along with temperature, against time. This technique involves the calorimetry of heat released in a chemical process, either a reaction or a conformational alteration. It can be utilized to calculate metrics such as the Heat of Reaction ( $\Delta$ rH), the change in enthalpy associated with a chemical reaction. When  $\Delta$ rH is negative, the process is exothermic and heat is released; when  $\Delta$ rH is positive, the process is endothermic and heat input is required. [66]. In the literature, DSC uses include evaluating structural-phase transition, melting point, and assessing molecular interaction between solid components [67, 68]. Litwinienko and co-workers [69] demonstrated that DSC can be used to investigate the oxidative stability of both simple [linolenic acid (LNA)] and complex (lecithin) lipids. By comparing the kinetics of soy lecithin autoxidation to that of LNA, it is possible to learn about the differences between the thermal oxidative behaviors of simple and complex lipids. Sopade et al. [70] investigated gelatinization in mixtures of sugars offers a promising frontier in clarifying the process. According to DSC research, this process is endothermic, with two stages: breaking of existing hydrogen bonds (endothermic) and creation of new bonds to produce a less-ordered structure (exothermic). DSC techniques could be useful to provide new data about the structural thermodynamics of reaction intermediates.[71] In particular, some reactions, such as isothermal amplification of nucleic acids, can be processed in DSC to obtain new thermodynamic information about polymerizing nucleic acid molecules [72]. A DSC thermograph might represent the complex process involving dehydration of the saccharide rings, depolymerization, and disintegration of the acetylated and deacetylated units of the polymer [73, 74] As a DSC application, to investigate the optimal temperature for the endothermic Fries rearrangement, our research utilized DSC as a promising screening tool for determining the optimal temperature.

### 2.4.4 Reduction

Reduction is the chemical reactions where one or more electrons are gained. When an atom gains one or more electrons during a chemical process, this is referred to as reduction. That suggests its oxidation number is decreasing.

Alcohols can be prepared from carbonyl compounds such as aldehydes, ketones, esters, acid chlorides and even carboxylic acids by hydride reductions. These reductions are the result of two net hydrogen atom additions to the C=O bond. Lithium aluminum hydride (LiAlH<sub>4</sub>) and sodium borohydride are the most prevalent hydride-reducing agents (NaBH<sub>4</sub>). LiAlH<sub>4</sub> is one of the most potent reducing agents, effectively reducing carbonyl. NaBH<sub>4</sub> is less reactive and can be used to convert aldehydes and ketones to alcohols selectively. Since NaBH<sub>4</sub> is not especially reactive, the reaction is often carried out in protic solvents such as ethanol or methanol. [75] the sodium ion is a weaker Lewis acid than the lithium ion, and the hydrogen bonding
between the alcohol and the carbonyl group serves as a catalyst to activate the carbonyl group. This reaction takes place as a result of the activation of the carbonyl group, as shown in Figure 16.



Figure 16 Mechanism of ketone reduction by NaBH<sub>4</sub>

## 2.5 Chemical structure of anti-TB agents

Based on anti-TB drugs, amino aromatic compounds are organic molecules with an amine (-NH<sub>2</sub>) group, which is a significant functional group in drug design. There are many well-known drugs containing amino aromatic structure such as antibiotic isoniazid, ethionamide, prothionamide, *para*-amino salicylic acid. In addition, compounds containing amino groups have been reported in the discovery of leads as new anti-tuberculosis agents (Figure 17). Some compounds containing amide functional group with therapeutic efficacy showed in Figure 18 [76, 77]. Chemically, amide is a powerful electron-withdrawing group that, by attracting electrons nearby, can generate localized or regional electron-deficient zones within molecules. Electrophilic amide groups can react with biological nucleophiles such as proteins, amino acids, enzymes, and nucleic acids in the target sites.



Figure 17 Structure of amino containing anti-TB compounds



Figure 18 Structure of amide containing anti-TB compounds

The addition of a halogen atom in the molecule is commonly used in drug design because halogen is the most electronegative element. Its electron-withdrawing effect can be attributed to changing the acidity, lipophilicity and conformation and modifying the interaction of halogenated compounds with the biological receptor or enzyme, all of which may be influenced by biological and pharmacological properties. As lead modification, halogen atom was incorporated at the different position of phenyl ring and it was observed that antituberculosis activities was reported as shown in Figure 19. [78, 79]



Figure 19 Structure of halogen containing anti-TB compounds

Cinnamic acid has been documented as a conventional anti-tuberculosis drug. [80] The development of 3-(4-cinnamylpiperazinyl iminomethyl) rifamycin SV (**XX**) was the first study to describe the synergistic action of the cinnamyl moiety on the piperazinyl group of the existing antituberculosis rifamycin. This investigation was supported by the enhancement of *trans*-cinnamic acid's activity in medication combinations with rifampin, amikacin, and clofazimine for tuberculosis infection [81]. The molecular hybridization between isoniazid and *trans*-cinnamic acid was described, and the MIC of the produced compound (**XXI**) was 3.12 µg/mL compared to 0.20 µg/mL for isoniazid [82]. Degani et al. developed novel molecular hybrids of cinnamic acid and guanyl hydrazone as compound (**XXII**) that demonstrated MIC of 6.49 µM against MTB H37Rv with an excellent safety profile [83]. The published literature indicates that *trans*-cinnamic acid-containing compounds had synergistic effects on existing antibiotics and promising activities on resistance strains. Therefore, cinnamic derivatives can be potential leads in the design and synthesis of antimycobacterial agents.



Figure 20 Structure of cinnamyl containing anti-TB compounds

The mycobacteria cell wall consists of highly long chain (C60-C90) abranched fatty acids esterified to the arabinogalactan component of the cell wall or trehalose. Its thick cell wall is one of the mechanisms preventing drugs from reaching the mycobacterial cytoplasm. Generally, lipophilic medicines penetrate through the cell membrane more easily and are more active [84]. Therefore, chemical modification of the compound structure to increase the permeability property of cell wall is quite challenging task. Stec J. et al. synthesized the triclosan derivatives to inhibit the enoyl-acyl carrier protein reductase InhA of MTB drug-sensitive and drugresistance strains. It was found that n-butyl attached triazole ring containing compound (XXIII) showed the promising activity with MIC value 0.6 µg/ml when in compare with triclosan (12.5 µg/ml) [85]. It has also been demonstrated that the longchain synthesis compounds XXIV and XXV inhibit M. tuberculosis H37Rv in vitro with a MIC90 value of 29 µM. In contrast, the parent thiolactomycin has a MIC90 value of 125 µM. It suggested that the addition of a long chain could increase activity. In addition, it was discovered that increasing the potency of compounds against M. tuberculosis by modifying the hydrophobicity of the side chain by varying its length and saturation.



Figure 21 Structure of long chain containing anti-TB compounds

According to published data, several functional groups have contributed to the design and development of more selective, drug-like, and pharmacokinetically effective antitubercular drugs for effective tuberculosis treatment.



# CHAPTER 3 MATERIALS AND METHODS

# **3.1 Materials**

# **3.1.1 Instruments**

Name	Source		
Analytical balance	Mettler Toledo, Switzerland		
Infrared spectrophotometer (FT-IR 4100)	Jasco, Japan		
Magnetic stirrer heating plate (IKA <sup>®</sup> C-MAG HS7)	IKA, USA		
Differential Scanning Calorimeter (823e)	Mettler Toledo, Switzerland		
High performance liquid chromatography-	Waters, USA		
Mass spectrometer (UPLC-PDA-QDa)			
Nuclear magnetic resonance spectrophotometer	Bruker, USA		
(300 MHz)			
Rotary evaporator	BÜCHI, Switzerland		
BÜCHI Heating Bath B 490	h		
BÜCHI Rotavapor R-205	50		
	67		

# 3.1.2 Chemical reagents and Instruments

Name	Source
Benzoic acid	Fluka, Switzerland
4-Fluorobenzoic acid	Sigma Aldrich, Germany
3-Bromobenzoic acid	Sigma Aldrich, Germany
4-Bromobenzoic acid	Sigma Aldrich, Germany
4-Butylbenzoic acid	Sigma Aldrich, Germany
3-Acetamidobenzoic acid	Sigma Aldrich, Germany
4-Acetamidobenzoic acid	Sigma Aldrich, Germany
Phenol	Sigma Aldrich, Germany
2-Naphthoic acid	Sigma Aldrich, Germany
4-Aminobenzophenone	Sigma Aldrich, Germany
Butyric acid	Sigma Aldrich, Germany

Name	Source
trans-cinnamic acid	Ajax FineChem Pty Ltd. Australia
N-(3-Dimethylaminopropyl)-N'-	Sigma Aldrich, Germany
ethylcarbodiimide hydrochloride (EDC·HCl)	
4-(Dimethylamino)pyridine (DMAP)	Sigma Aldrich, Germany
Aluminium Chloride	Sigma Aldrich, Germany
Sodium borohydride	Sigma Aldrich, Germany
Benzyl chloroformate	Sigma Aldrich, Germany
Sodium hydroxide	Merck, Germany
Acetic anhydride	J.T.Baker, USA
Sodium sulfate anhydrous	Ajax FineChem Pty Ltd. Australia
Molecular sieve (4A° beads 8-12 mesh)	Sigma Aldrich, Germany
Silica gel 60 (0.063-0.2 mm)	Merck, Germany
Silica gel 60 (0.04-0.06 mm)	Merck, Germany
TLC silica gel 60 F254	Merck, Germany
Dichloromethane AR grade	J.T.Baker, USA
Ethyl acetate AR grade	J.T.Baker, USA
Hexane AR grade	J.T.Baker, USA
Methanol AR grade	J.T.Baker, USA
Ethanol AR grade	J.T.Baker, USA
Toluene	Carlo Erba, France
Tetrahydrofuran	Carlo Erba, France
Methanol HPLC grade	Honeywell, Burdick&Jackson, USA
Acetonitrile HPLC grade	Honeywell, Burdick&Jackson, USA
Nitrogen gas	Masser Speciality Gas Co.Ltd,
	Thailand
Carbon dioxide gas	Masser Speciality Gas Co.Ltd,
	Thailand

#### **3.1.3 Chemical preparations**

All solvents and starting materials were purchased from Acros Organics, Fluka, Merck KGaA, Ajax Finechem and Sigma-Aldrich. Dichloromethane for esterification was stored over molecular sieves 4Å. The purification of the compounds were isolated by column chromatography on silica gel 60 (0.063-0.2 mm and 0.04-0.06 mm, Merck) by using stated solvent systems and equilibrated with those systems prior to use. The reactions were monitored by thin-layer chromatography (TLC) on precoated Merck 60 F254 silica gel plates and visualized using UV light (254 nm). IR spectra of synthesized compounds were recorded (Neat or KBR pellets or nujol mullets) on a Jasco FT-IR spectrophotometer (Japan) and reported as a wavenumber (cm<sup>-1</sup>). All <sup>1</sup>H and <sup>13</sup>C NMR spectra were recorded on a Bruker-Ultra Shield (300 and 75 MHz for <sup>1</sup>H and <sup>13</sup>C, respectively), using CD<sub>3</sub>COCD<sub>3</sub>, CDCl<sub>3</sub> and CD<sub>3</sub>OD as solvents with trace of tetramethylsilane (TMS) as an internal standard. The chemical shifts ( $\delta$ ) were reported in ppm and coupling constants (J) were reported as Hertz (Hz). Signal multiplicities were represented by s (singlet), d (doublet), t (triplet), q (quartet), quint (quintet), sext (sextet), m (multiplet), app (apparent) and br (broad). Fries rearrangement reaction was determined on a DSC 823e (Mettler Toledo, Switzerland) at heating from 40 °C to 400 °C and was carried out at a heating rate of 5 °C/min under nitrogen atmosphere. หารทยาลัยศิลปาก

# 3.2 Methods3.2.1 General methods for chemistry

Synthetic routes of benzhydrol derivatives are depicted in schemes 1-3. Esterification reaction and Fries rearrangement in scheme 1 were used for synthesis of compounds 4a-f and 9. In the synthesis of aminobenzhydrol derivatives 10 in scheme 2, amino functional groups could be obtained from hydrolysis of acetamide group by aqueous base reaction. In case of amide derivatives in scheme 3, 4-amino benzophenone was coupled with butyric acid, benzoic acid, and cinnamic acid via amide formation to give compounds 13a-c.

All reagents and solvents were used from commercial sources, and all reactions were monitored by TLC using silica plates with fluorescence F254 and UV light visualization. All crude products were confirmed the molar mass of desired compounds with LC-MS. All synthesized compounds were purified by column chromatography Their structures were characterized by spectral data (<sup>1</sup>H, <sup>13</sup>C NMR, IR and MS).



Reactions and conditions : (a) Phenol, EDC, DMAP, CH<sub>2</sub>Cl<sub>2</sub>, rt, overnight; (b) AlCl<sub>3</sub>, 120-140°C, 2h; (c) NaBH<sub>4</sub>, MeOH, rt, 3h; (d) AcO<sub>2</sub>, EDC, DMAP, CH<sub>2</sub>Cl<sub>2</sub>, rt, overnight

Scheme 1. Synthesis of compounds 4a-f, 5a, 5c, 5f and 9

In scheme 1, first, the synthesis of compound 4a-f, benzoic acid and substituted benzoic acid (1a-f) were used as starting materials. Its carboxylic acid group was coupled with phenol in the presence of EDC·HCl and DMAP to form phenyl benzoate. The benzoate derivatives were changed to 4-hydroxy benzophenone (3a-f) by performing Fries rearrangement reaction using AlCl<sub>3</sub>. After that, benzophenones were converted to benzhydrols by reduction using NaBH<sub>4</sub> to give benzhydrol derivatives (4a-f). Due to their significant anti-TB properties, benzhydrol derivatives (4a, 4c, and 4f) were chosen as starting materials for the synthesis of diacetate compounds (5a, 5c, and 5f). Diacetylation was performed by adding acetyl groups into hydroxyl groups of benzhydrol derivatives using acetic anhydride as an acetylating agent. In the synthesis of compound 9, 2-naphthoic acid served as starting material and was converted to compound 9 via reactions a, b and c.



Reaction and conditions: (a) aq. NaOH, EtOH, 50°C, 12 h

Scheme 2. Synthesis of compound 10

In scheme 2, the synthesis of compound 10, 4-acetamido-4'-acetylbenzhydryl acetate (4f) was used as starting material, and the acetyl group at N position was removed by base-catalysed hydrolysis to provide 4-amino-4'acetylbenzhydryl acetate (10).





Reaction and conditions: (a) EDC, DMAP, CH<sub>2</sub>Cl<sub>2</sub>, rt, overnight; (b) NaBH<sub>4</sub>, MeOH, rt, 3h.

# Scheme 3. Proposed synthesis of compound 13a-c

In scheme 3, 4-amino benzophenone was used as a starting material. Its amino group was linked with an alkylcarboxylic acid or an arylcarboxylic acid (**11a-c**) to form amide derivatives (**12a-c**). Then, benzophenone scaffolds were changed to benzhydrol derivatives (**13a-c**) by means of the reduction reaction.



A general procedure for coupling reaction with ester bonds was performed as follows. [86, 87] All glassware were dried in hot-air oven at 120 °C for 4 hours. DCM was dried by storing over freshly dried and cooled 4A molecular sieves for overnight. Aromatic acids (**1a-g**) (1 equiv), EDC·HCl (2 equiv), and DMAP (2 equiv) were weighed separately and dissolved in dried DCM. The mixture was stirred for 1h

or until the mixture was dissolved completely. Then, phenol (1 equiv) in dried DCM was added by stirring to the round-bottomed flask containing acids, EDC and DMAP. Reaction was run overnight at room temperature and monitored by TLC. After complete reaction, the reaction mixture was worked-up with water and evaporated DCM. Water layer was extracted with EtOAc (30mLx3). Organic layer then was washed with water until pH of water layer is neutral and then dried with anhydrous Na<sub>2</sub>SO<sub>4</sub>, filtered, and concentrated under reduced pressure. The desired products (**2a**-**g**) were used directly in the next step without purification.

#### Phenyl benzoate (2a)

To a solution of benzoic acid **1a** (626 mg, 5 mmol, 1 equiv), EDC·HCl (1552 mg, 10 mmol, 2 equiv) and DMAP (1220 mg, 10 mmol, 2 equiv) in dried DCM, stirred for 1h, added phenol (477 mg, 5 mmol, 1 equiv). Reaction was performed according to general procedure for esterification. Compound **2a** as white crystalline powder (849 mg, 78.9%). FT-IR (KBr), *v*: 1729 cm<sup>-1</sup> (C=O stretching in ester).

# 4-Fluorophenyl benzoate (2b)

To a solution of 4-fluoro benzoic acid **1b** (1,405 mg, 10 mmol, 1.4 equiv), EDC·HCl (2,643 mg, 17 mmol, 2.3 equiv) and DMAP (1,865 mg, 15 mmol, 2.1 equiv) in dried DCM, stirred for 1h, added phenol (680 mg, 7.2 mmol, 1 equiv). Reaction was performed according to general procedure for esterification. Compound **2b** as white crystalline powder (1,492 mg, 95.8%). FT-IR (KBr), *v*: 1737 cm<sup>-1</sup> (C=O stretching in ester).

#### 4-Bromophenyl benzoate (2c)

To a solution of 4-bromo benzoic acid **1c** (1,312 mg, 6 mmol, 1.2 equiv), EDC·HCl (4,800 mg, 25 mmol, 5 equiv) and DMAP (3,087 mg, 25 mmol, 5 equiv) in dried DCM, stirred for 1h, added phenol (498 mg, 5 mmol, 1 equiv). Reaction was performed according to general procedure for esterification. Compound **2b** as off-white powder (1,339 mg, 96.7%). FT-IR (KBr), *v*: 1731 cm<sup>-1</sup> (C=O stretching in ester); LC-MS (ESI, m/z) [M+H]+: 278.

#### 4-Butylphenyl benzoate (2d)

To a solution of 4-butyl benzoic acid **1d** (1,785 mg, 10 mmol, 1.25 equiv), EDC·HCl (3,700 mg, 22.5 mmol, 2.8 equiv) and DMAP (2,740 mg, 22.5 mmol, 2.8 equiv) in dried DCM, stirred for 1h, added phenol (770 mg, 8 mmol, 1 equiv). Reaction was performed according to general procedure for esterification. Compound **2d** as pale-yellow liquid (1,900 mg, 93.9%). FT-IR (KBr), *v*: 1713 cm<sup>-1</sup> (C=O stretching in ester).

#### **3-Acetamidophenyl benzoate (2e)**

To a solution of 3-acetamido benzoic acid **1e** (1,865 mg, 10.4 mmol, 1.1 equiv), EDC·HCl (3,135 mg, 20 mmol, 2.1 equiv) and DMAP (2,370 mg, 19.4 mmol, 2 equiv) in dried DCM, stirred for 1h, added phenol (908 mg, 9.6 mmol, 1 equiv). Reaction was performed according to general procedure for esterification. Compound **2e** as white powder (2,048 mg, 83.6%). FT-IR (KBr), *v*: 1724 cm<sup>-1</sup> (C=O stretching in ester).

## 4-Acetamidophenyl benzoate (2f)

To a solution of 4-acetamido benzoic acid **1f** (1,802 mg, 10 mmol, 1 equiv), EDC·HCl (2,765 mg, 17.8 mmol, 1.8 equiv) and DMAP (1,890 mg, 15.5 mmol, 1.6 equiv) in dried DCM, stirred for 1h, added phenol (936 mg, 10 mmol, 1 equiv). Reaction was performed according to general procedure for esterification. Compound **2e** as white powder (2,104 mg, 82.5%). FT-IR (KBr), *v*: 1726 cm<sup>-1</sup> (C=O stretching in ester).

#### **3.2.2.2** Synthesis of 4-hydroxy benzophenone derivatives (3a-f)

#### General procedure for synthesis of Fries rearrangement



Substituted phenyl benzoates (**2a-g**) (1 equiv) and AlCl<sub>3</sub> (10 equiv) were crushed and mixed together in mortar. The mixture was stirred at 120-140 °C for 2h and reaction was monitored by TLC. After 2 hours, reaction was quenched with water and the reaction mixture then was extracted with EtOAc (30mLx3). The combined organic phase was washed with water (50mL) until pH of water is about 7, dried with anhydrous Na<sub>2</sub>SO<sub>4</sub>, filtered and evaporated. Crude products (**3a-g**) were purified by column chromatography (SiO<sub>2</sub>, EtOAc:Hex:MeOH, 5:5:1) [62, 88]

#### 4-Hydroxy benzophenone (3a)

The mixture of phenylbenzoate **2a** (849 mg, 3.9 mmol, 1 equiv) and AlCl<sub>3</sub> (5,070 g, 40 mmol, 10 equiv) was stirred at 140 °C for 45 min. Reaction was performed according to general procedure of Fries rearrangement. Crude products (**3a**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 5:5:1). Compound **3a** as white crystalline powder (284 mg, 33.7%). FT IR (KBr), *v*: 1673 cm<sup>-1</sup> (C=O stretching in ketone).

## 4-Fluoro-4'-hydroxybenzopheone (3b)

The mixture of 4-fluorophenyl benzoate **2b** (864 mg, 4 mmol, 1 equiv) and AlCl<sub>3</sub> (5,300 mg, 40 mmol, 10 equiv) was stirred at 140 °C for 45 min. Reaction was performed according to general procedure of Fries rearrangement. Crude products (**3b**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 2:8:1). Compound **3b** as white crystalline powder (410 mg, 47.5%). FT IR (KBr), *v*: 1604 cm<sup>-1</sup> (C=O stretching in ketone).

#### 4-Bromo-4'-hydroxybenzopheone (3c)

The mixture of 4-bromophenyl benzoate 2c (1,339 mg, 4.8 mmol, 1 equiv) and AlCl<sub>3</sub> (6,445 mg, 48 mmol, 10 equiv) was stirred at 140 °C for 30 min. Reaction was performed according to general procedure of Fries rearrangement. Crude products (**3c**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 2:8:1). Compound **3b** as pale-yellow powder (867 mg, 65.1%). FT IR (KBr), *v*: 1650 cm<sup>-1</sup> (C=O stretching in ketone).

#### 4-Butyl-4'-hydroxybenzopheone (3d)

The mixture of 4-butylphenyl benzoate **2d** (1,900 mg, 7.47 mmol, 1 equiv) and AlCl<sub>3</sub> (9,965 mg, 74.7 mmol, 10 equiv) was stirred at 130 °C for 30 min. Reaction was performed according to general procedure of Fries rearrangement. Crude products (**3d**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 2:8:1). Compound **3b** as pale-yellow liquid (1,502 mg, 78.9%). FT IR (KBr), *v*: 1665 cm<sup>-1</sup> (C=O stretching in ketone) ; LC-MS (ESI, *m/z*) [M+H]+: 255.

#### 3-Acetamido-4'-hydroxybenzopheone (3e)

The mixture of 3-acetamidophenyl benzoate **2e** (517 mg, 2 mmol, 1 equiv) and AlCl<sub>3</sub> (3,242 mg, 24.3 mmol, 12 equiv) was stirred at 140 °C for 30 min. Reaction was performed according to general procedure of Fries rearrangement. Crude products (**3e**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 5:5:1). Compound **3e** as pale yellow liquid (203 mg, 44.9%). FT IR (KBr), *v*: 1665 cm<sup>-1</sup> (C=O stretching in ketone) ; LC-MS (ESI, *m/z*) [M+H]+: 226.

# 4-Acetamido-4'-hydroxybenzopheone (3f)

The mixture of 4-acetamidophenyl benzoate **2f** (1104 mg, 4.3 mmol, 1 equiv) and AlCl<sub>3</sub> (5,53 mg, 38.6 mmol, 9 equiv) was stirred at 140 °C for 60 min. Reaction was performed according to general procedure of Fries rearrangement. Crude products (**3f**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 5:5:1). Compound **3e** as pale yellow liquid (473 mg, 43.1%). FT IR (KBr), *v*: 1665 cm<sup>-1</sup> (C=O stretching in ketone) ; LC-MS (ESI, *m/z*) [M+H]+: 226.

#### **3.2.2.3** Synthesis of (4-Hydroxyphenyl)phenylmethanol derivatives (4a-f)

#### General procedure for reduction



To a solution of substituted benzophenones (**3a-g**, **17a-c**) (1 equiv) in MeOH, NaBH<sub>4</sub> (10 equiv) was added. The mixture was stirred at room temperature for 3h or until starting material disappeared. After that, the solvent was evaporated by rotary evaporator. Water was added to the mixture for working-up and the mixture was extracted with EtOAc (50mLx2). The organic layers were combined and washed with water (50mLx3). After drying over Na<sub>2</sub>SO<sub>4</sub>, EtOAc was removed under reduced pressure to yield desired products (**4a-g**, **9**, **18a-c**) [89]. Crude products (**4a-g**) were purified by column chromatography.

# 4-hydroxy-α-(phenyl)benzyl alcohol (4a)

To a solution of 4-hydroxy benzophenone (**3a**) (284 mg, 1.3 mmol, 1 equiv) in dried MeOH, added NaBH<sub>4</sub> (492 mg, 13 mmol, 10 equiv). Reaction was performed according to general procedure for reduction reaction. Compound **4a** as white crystalline powder (215 mg, 76.2%). IR (cm<sup>-1</sup>): OH str. 3335 (broad); 1H NMR (300 MHz), CD<sub>3</sub>COCD<sub>3</sub>  $\delta$  (ppm): 8.21 (1H, s), 7.40 (2H, d, J = 8.7 Hz), 7.28 (3H, t, J = 7.8 Hz), 7.21 (2H, d, J = 8.7 Hz), 6.76 (2H, d, J = 8.7 Hz), 5.73 (1H, s), 4.64 (1H, s); LC-MS (ESI) [M+H]+ m/z 200

# 4-hydroxy-α-(4'-fluorophenyl)benzyl alcohol (4b)

To a solution of 4-fluoro-4'-hydroxybenzopheone (**3b**) (821 mg, 3.8 mmol, 1 equiv) in dried MeOH, added NaBH<sub>4</sub> (1,436 mg, 38 mmol, 10 equiv). Reaction was performed according to general procedure for reduction reaction. Compound 4b as white crystalline powder (489 mg, 59.3%). FT-IR (cm<sup>-1</sup>): OH str. 3384 (broad), 3180 (broad); <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>OD,  $\delta$  (ppm): 7.35 (2H, dd, J = 5.5, 2.9 Hz, H-Ar), 7.15 (2H, d, J = 8.3 Hz, H-Ar), 7.02 (2H, t, J = 8.9 Hz, H-Ar), 6.73 (2H, d, J = 8.6 Hz, H-Ar), 5.69 (1H, s, CH-OH); <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD),  $\delta$  (ppm): 165.0, 161.8, 157.9, 142.4, 142.4, 136.8, 129.6, 129.5, 129.2, 116.2, 116.0, 115.7, 76.1. LC-MS (ESI, *m/z*): 217.06

#### 4-hydroxy-α-(4'-bromophenyl)benzyl alcohol (4c)

To a solution of 4-bromo-4'-hydroxybenzopheone (3c) (867mg, 3.1 mmol, 1 equiv) in dried MeOH, added NaBH<sub>4</sub> (586 mg, 15.5 mmol, 5 equiv). Reaction was

performed according to general procedure for reduction reaction. Compound **4c** as pale-yellow powder (778 mg, 89.9%). FT-IR (cm<sup>-1</sup>): OH str. 3384 (broad), 3150 (broad); <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>COCD<sub>3</sub>,  $\delta$  (ppm): 8.30 (1H, bs), 7.47 (2H, d, *J* = 8.4 Hz), 7.35 (2H, d, *J* = 8.7 Hz), 7.19 (2H, d, *J* = 8.4 Hz), 6.77 (2H, d, *J* = 8.7 Hz), 5.73 (1H, s), 4.82 (1H, s); LC-MS (ESI, *m/z*): 279

#### 4-hydroxy-α-(4'-butylphenyl)benzyl alcohol (4d)

To a solution of 4-butyl-4'-hydroxybenzopheone (**3d**) (489 mg, 2 mmol, 1 equiv) in dried MeOH, added NaBH<sub>4</sub> (756 mg, 20 mmol, 10 equiv). Reaction was performed according to general procedure for reduction reaction. Compound **4d** as pale-yellow liquid (342 mg, 67.1%). IR (cm<sup>-1</sup>): OH str. 3340 (broad), 3168 (broad); <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>OD,  $\delta$  (ppm): 7.23 (2H, d, J = 8.0 Hz), 7.14 (4H, q), 6.73 (2H, d, J = 8.6 Hz), 5.66 (1H, s, CH-OH), 2.58 (2H, t, J = 7.5 Hz, -CH<sub>2</sub>-), 1.57 (2H, qui, J = 7.6 Hz, -CH<sub>2</sub>-), 1.34 (2H, sext, J = 7.6 Hz, -CH<sub>2</sub>-), 0.92 (3H, t, J = 7.3 Hz, -CH<sub>3</sub>-).

# 4-hydroxy-α-(3'-acetamidophenyl)benzyl alcohol (4e)

To a solution of 3-acetamido-4'-hydroxybenzopheone (**3e**) (203 mg, 0.9 mmol, 1 equiv) in dried MeOH, added NaBH<sub>4</sub> (586 mg, 15.5 mmol, 15 equiv). Reaction was performed according to general procedure for reduction reaction. Compound **4e** as pale yellow liquid (171 mg, 83.7%). IR (cm<sup>-1</sup>): OH str. 3307 (broad), 3189 (broad) 1673 (C=O str.); <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>COCD<sub>3</sub>,  $\delta$  (ppm): 9.08 (NH, br s), 8.22 (OH, br s), 7.60 (1H, d, J = 9.0 Hz, H-Ar), 7.54 (1H, s, H-Ar), 7.21 (3H, m, HAr), 7.08 (1H, d, J = 9.0 Hz, H-Ar), 6.76 (2H, dt, J = 9.0 Hz, H-Ar), 5.69 (1H, s, CH-OH), 4.65 (OH, d), 2.89 (3H, s, CH3). LC-MS (ESI, m/z): 257.03

#### 4-hydroxy-α-(4'-acetamidophenyl)benzyl alcohol (4f)

To a solution of 4-acetamido-4'-hydroxybenzopheone (**3f**) (270 mg, 1.1 mmol, 1 equiv) in dried MeOH, added NaBH<sub>4</sub> (501 mg, 13.3 mmol, 12.1 equiv). Reaction was performed according to general procedure for reduction reaction. Compound **4f** as pale yellow liquid (189 mg, 67.4%). IR (cm<sup>-1</sup>): OH str. 3307 (broad), 1668 (C=O str.); <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>OD,  $\delta$  (ppm): 7.67 (6H, m, J = 8.7 Hz, H-Ar), 6.85

(2H, d, J = 8.7 Hz, H-Ar), 5.74 (1H, s, CH-OH), 4.88 (OH, s), 2.14 (3H, s, CH3). LC-MS (ESI, m/z): 257.06

#### **3.2.2.4** 4-acetoxy-α-(phenyl)benzyl acetate derivatives (5a, 5d, 5g)

General procedure for acetylation



A general procedure for acetylation of dihydroxy-containing compounds was performed as follows. Oven-dried glassware was used for the reaction. Acetic anhydride (1 equiv), EDC·HCl (3 equiv), and DMAP (3 equiv) were dissolved in dry DCM and stirred for 1 hour. Compound **4a**, **4c**, and **4f** in dry DCM was added to the reaction mixture. The mixtures were stirred overnight at room temperature. After that, DCM was evaporated and water was added. Water layer was extracted with EtOAc (50mLx2), EtOAc layer was washed by water (50mL), dried over Na<sub>2</sub>SO<sub>4</sub>, filtered and concentrated under reduced pressure. The desired products (**5a**, **5c**, **5f**) were purified by column chromatography.

#### 4-acetoxy-α-(phenyl)benzyl acetate (5a)

To a solution of acetic anhydride (51 mg, 0.5 mmol, 1 equiv), EDC·HCl (780 mg, 5 mmol, 10 equiv) and DMAP (611 mg, 5 mmol, 10 equiv) in dried DCM, stirred for 1h, added a solution of 4-hydroxy- $\alpha$ -(phenyl)benzyl alcohol (**4a**) (105 mg, 0.49 mmol, 1 equiv) in dried DCM. Reaction was performed according to general procedure for acetylation. Compound **4a** as white crystalline powder (138 mg, 92.9%). FT-IR (KBr), *v*: C=O str. 1735 (sharp); <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>)  $\delta$ : 7.36-7.26 (7H, m), 7.05 (2H, d, J = 8.7 Hz), 6.88 (1H, s), 2.28 (3H, s), 2.15 (3H, s)

#### 4-acetoxy-α-(4'-bromophenyl)benzyl acetate (5c)

To a solution of acetic anhydride (245 mg, 2.4 mmol, 1.2 equiv), EDC·HCl (1,970 mg, 10 mmol, 5 equiv) and DMAP (1,230 mg, 10 mmol, 5 equiv) in dried DCM, stirred for 1h, added a solution of 4-hydroxy- $\alpha$ -(4'-bromophenyl)benzyl alcohol (**4c**) (554 mg, 2 mmol, 1 equiv) in dried DCM. Reaction was performed according to general procedure for acetylation. Compound **5c** as white crystalline powder (502 mg, 69.3%). FT-IR (KBr), *v*: C=O str. 1735.9 (sharp); <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>)  $\delta$ : 7.46 (2H, d, *J* = 8.4 Hz), 7.31 (2H, d, *J* = 8.7 Hz), 7.20 (2H, d, *J* = 8.4 Hz), 7.06 (2H, d, *J* = 8.7 Hz), 6.82 (1H, s), 2.28 (3H, s), 2.15 (3H, s)

# 4-acetoxy-α-(4'-acetamidophenyl)benzyl acetate (5f)

To a solution of acetic anhydride (40 mg, 0.4 mmol, 1 equiv), EDC·HCl (629 mg, 4 mmol, 11.4 equiv) and DMAP (487 mg, 4 mmol, 11.4 equiv) in dried DCM, stirred for 1h, added a solution of 4-hydroxy- $\alpha$ -(4'-acetamidophenyl)benzyl alcohol (**4f**) (90 mg, 0.35 mmol, 1 equiv) in dried DCM. Reaction was performed according to general procedure for acetylation. Compound **5f** as pale-yellow liquid (110 mg, 91.6%). FT-IR (KBr), *v*: C=O str. 1701 (sharp); <sup>1</sup>H NMR (300 MHz, CD<sub>3</sub>OD)  $\delta$ : 7.63 (6H, d, *J* = 8.7 Hz), 6.89 (2H, d, *J* = 8.7 Hz), 6.82 (1H, s), 2.28 (3H, s), 2.15 (6H, s)

#### 3.2.2.5 2-Naphthyl benzoate (7)



The synthesis of 2-Naphthyl benzoate (**7**) using 2-naphthoic acid (**6**) (1,725 mg, 10 mmol, 1 equiv), EDC·HCl (2,511 mg, 15 mmol, 1.5 equiv), and DMAP (1,860 mg, 15 mmol, 1.5 equiv) in dried DCM, stirred for 1h, added phenol (741 mg, 8 mmol, 0.8 equiv), was followed as general procedure of esterification. The obtained product, 2-Naphthyl benzoate (**7**), were used directly in the next step without purification. Compound **7** as white crystalline powder (1,734 mg, 87.4%). FT-IR (KBr), *v*: 1724 cm<sup>-1</sup> (C=O stretching in ester); LC-MS (ESI, *m/z*) [M+H]+: 249.

#### 3.2.2.6 4-hydroxyphenyl(2-naphthyl)methanone (8)



2-Naphthyl benzoate (7) (1,334 mg, 5 mmol, 1 equiv) and AlCl<sub>3</sub> (6,868 mg, 50 mmol, 10 equiv) were crushed and mixed. The reaction mixture was stirred at 130 °C for 1h. The rest of the preparation method was followed as general procedure of Fries rearrangement. Crude products **8** were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 3:7:1). Compound **8** as pale-yellow crystalline powder (740 mg, 59.7%). FT-IR (KBr), *v*: 1600 cm<sup>-1</sup> (C=O stretching in ketone); LC-MS (ESI, *m/z*) [M+H]+: 249.

3.2.2.7 4-hydroxyphenyl(2-naphthyl)methanol (9)



The synthesis of 4-hydroxyphenyl(2-naphthyl)methanol (**9**) using 2-naphthyl ketone (**8**) (380 mg, 1.5 mmol, 1 equiv) in MeOH, NaBH<sub>4</sub> (587 mg, 15 mmol, 10 equiv) was followed as general procedure of reduction. The desired product (**9**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 2:8:1). Compound **9** as off-white crystalline powder (289 mg, 77.7%). <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>OD,  $\delta$  (ppm): 7.86-7.75 (4H, m, Ar-H), 7.43 (3H, m, Ar-H), 7.21 (2H, d, J = 8.4 Hz), 6.75 (2H, d, J = 8.6 Hz), 5.56 (1H, s, CH-OH). <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD),  $\delta$  (ppm): 116.18-157.90 (Ar-C), 76.86 (C-OH); LC-MS (ESI, m/z) [M-OH]+: 233.

#### 3.2.2.8 4-((4-aminophenyl)(hydroxy)methyl)phenol (10)



Amide hydrolysis was performed under alkaline condition using aqueous NaOH as catalyst. The solution of NaOH (3,086 mg, 77 mmol, 10 equiv) was added by stirring to the round-bottomed flask containing solution of compound **4f** (1,950 mg, 7.6 mmol, 1 equiv) in EtOH at 50 °C for 12h (Shimizu et al. 2014). The reaction was monitored by TLC. After complete reaction, EtOH was evaporated and then water was added for working-up. EtOAc (50mlx2) was used for extraction. The combined organic layer was washed with water until pH of water phase is neutral. Then, organic layer was dried over anhydrous Na<sub>2</sub>SO<sub>4</sub>, filtered, and dried by evaporation of solvent. Crude product (**10**) was purified by column chromatography (SiO<sub>2</sub>, EtOAc:Hex:MeOH, 3:7:1). Compound **10** as pale yellow powder (446 mg, 27.4%). FT-IR (cm<sup>-1</sup>), *v*: N-H str. 3328 (sharp), O-H str. about 3300 (board), C-H str. 3023, N-H bend 1618, C=C str. 1617-1450, C-O str. 1176, 1122, C-N str. 1018. LC-MS (ESI, *m/z*) [M]+: 214.

# 3.2.2.9 4-amido benzophenone derivatives (12a-c)

General procedure for amide formation



A general procedure for coupling reaction with amide bonds is performed as follows. All glassware were dried in hot-air oven at 120 °C for 4 hours. DCM was dried by storing over freshly dried and cooled 4A molecular sieves for overnight.

Aliphatic acids or aromatic acids (**11a-c**) (1 equiv), EDC·HCl (3 equiv), and DMAP (1 equiv) were weighed separately and dissolved in dried DCM. The mixture was stirred for 1h or until the mixture is dissolved completely. Then, 4-aminobenzophenone (1 equiv) in dried DCM was added by stirring to the round-bottomed flask containing acids, EDC and DMAP. Reaction was run overnight at room temperature and monitored by TLC. After complete reaction, the reaction mixture was worked-up with water and evaporated DCM. Water layer was extracted with EtOAc (30mLx3). Organic layer then was washed with water until pH of water layer is neutral and then dried with anhydrous Na<sub>2</sub>SO<sub>4</sub>, filtered, and concentrated under reduced pressure. The desired products (**12a-c**) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 3:7:1).

#### 4-butyramido benzophenone (12a)

To a solution of butyric acid **11a** (881 mg, 10 mmol, 1.25 equiv), EDC·HCl (2,376 mg, 15 mmol, 1.88 equiv) and DMAP (1,859 mg, 15 mmol, 1.88 equiv) in dried DCM, stirred for 1h, added a solution of 4-aminobenzophenone (1,577 mg, 8 mmol, 1 equiv) in dried DCM. Reaction was performed according to general procedure for amide formation. Compound **12a** as off-white powder (1,538 mg, 72.0%). FT-IR (cm<sup>-1</sup>), *v*: 3305 (N-H str.), 1658(C=O str. in amide), 1600 (C=O str. in ketone). LC-MS (ESI, m/z) [M]+: 268.

## 4-benzamido benzophenone (12b)

To a solution of benzoic acid **11b** (1,220 mg, 10 mmol, 2 equiv), EDC·HCl (1,563 mg, 10 mmol, 2 equiv) and DMAP (1,230 mg, 10 mmol, 2 equiv) in dried DCM, stirred for 1h, added a solution of 4-aminobenzophenone (993 mg, 5 mmol, 1 equiv) in dried DCM. Reaction was performed according to general procedure for amide formation. Compound **12b** as off-white powder (897 mg, 59.6%). FT-IR (cm<sup>-1</sup>), *v*: 3312 (N-H str.), 1650 (C=O str. in amide), 1590 (C=O str. in ketone).

#### 4-cinnamido benzophenone (12c)

To a solution of *trans*-cinnamic acid **11c** (459 mg, 3 mmol, 1.5 equiv), EDC·HCl (623 mg, 4 mmol, 2 equiv) and DMAP (494 mg, 4 mmol, 2 equiv) in dried DCM, stirred for 1h, added a solution of 4-aminobenzophenone (405 mg, 2 mmol, 1 equiv) in dried DCM. Reaction was performed according to general procedure for amide formation. Compound **12c** as off-white powder (206 mg, 31.5%). FT-IR (cm<sup>-1</sup>), *v*: 3323 (N-H str.), 1658 (C=O str. in amide), 1660 (C=O str. in ketone).

#### 3.2.2.10 4-amido benzhydrol derivatives (13a-c)



The synthesis of benzhydrol compounds (13a-c) using benzophenone compounds (12a-c) (1 equiv) dissolved in MeOH, NaBH<sub>4</sub> (10 equiv) was followed as general procedure of reduction. The desired product (13a-c) were purified by column chromatography (SiO2, EtOAc:Hex:MeOH, 3:7:1)

# 4-butyramido-α-(phenyl)benzyl alcohol (13a)

To a solution of 4-butyramido benzophenone (**12a**) (249 mg, 0.9 mmol, 1 equiv) and NaBH<sub>4</sub> (412 mg, 9 mmol, 10 equiv) in MeOH, stirred at room temperature for 1h, was followed as general procedure of reduction. Compound **13a** as off-white powder (192 mg, 79.9%). FT-IR (cm<sup>-1</sup>), *v*: 3303 (N-H str.), 3328 (O-H str.), 3124 (sp2 C-H), 3057 (sp2 C-H), 2962-2871 (sp3 C-H str.), 1660 (C=O str. in amide); <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>OD,  $\delta$  (ppm): 7.50 (2H, d, Ar-H), 7.37 -7.18 (7H, m, Ar-H), 5.78 (1H, s, CH-OH), 2.32 (2H, t, J = 7.4 Hz), 1.70 (2H, sext, J = 7.4 Hz), 0.98 (3H, t, J = 7.4 Hz). <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD),  $\delta$  (ppm): 174.68 (C=O), 145.98-121.25 (Ar-C), 76.68 (C-OH); LC-MS (ESI, *m/z*) [M-OH]+: 252.

#### 4-benzamido-α-(phenyl)benzyl alcohol (13b)

To a solution of 4-benzamido benzophenone (**12b**) (897 mg, 3 mmol, 1 equiv) and NaBH<sub>4</sub> (1134 mg, 30 mmol, 10 equiv) in MeOH, stirred at room temperature for 1h, was followed as general procedure of reduction. Compound **13a** as off-white powder (451 mg, 49.9%). FT-IR (cm<sup>-1</sup>), *v*: 3318 (N-H str.), 3300 board (O-H str.), 3060 (sp2 C-H ), 3030 (sp2 C-H ), 1651 (C=O str. in amide), 1182 (C-O str.) ; <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>OD,  $\delta$  (ppm): 7.92 (2H, d, J = 8.1 Hz, Ar-H), 7.64 (2H, d, J = 11.0 Hz, Ar-H), 7.60 -7.47 (3H, m, Ar-H), 7.40 -7.20 (7H, m, Ar-H), 5.78 (1H, s, CH-OH); <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD),  $\delta$  (ppm): 169.04 (C=O), 146.04 -122.32 (Ar-C), 76.73 (C-OH).

# 4-cinnamido-α-(phenyl)benzyl alcohol (13c)

To a solution of 4-cinnamido benzophenone (**12c**) (200 mg, 0.6 mmol, 1 equiv) and NaBH<sub>4</sub> (287 mg, 7.6 mmol, 12.6 equiv) in MeOH, stirred at room temperature for 1h, was followed as general procedure of reduction. Compound **13c** as off-white powder (151 mg, 77.0%). FT-IR (cm<sup>-1</sup>), *v*: 3313 (N-H str.), 3311 board (O-H str.), 3058 (sp2 C-H ), 3029 (sp2 C-H ), 1658 (C=O str. in amide), 1184, 1112(C-O str.) ; <sup>1</sup>H NMR (300 MHz), CD<sub>3</sub>OD,  $\delta$  (ppm): 7.68 -7.55 (5H, m, Ar-H), 7.44 -7.19 (10H, m, Ar-H), 6.78 (1H, d, CH=CH), 5.74 (1H, s, CH-OH); <sup>13</sup>C NMR (75 MHz, CD<sub>3</sub>OD),  $\delta$  (ppm): 166.78 (C=O), 146.02 -121.23 (Ar-C), 76.71 (C-OH).

# 3.3 Evaluation of antitubercular activities

#### Determination of minimum inhibitory concentration by agar dilution method

Synthesized chemical compounds were tested for their antitubercular activities against *M.tuberculosis* H37Rv and 20 MTB clinical isolates at National Center for Genetic Engineering and Biotechnology (BIOTEC), using agar dilution method. [90] This is standard method to find minimum inhibitory concentration (MIC) for antimycobacterial drug-susceptibility testing. In agar dilution method, MIC means the lowest drug concentration inhibiting visible growth of a bacterial population of *M.tuberculosis* on solid Middlebrook agar medium within 21 days of incubation at 37°C. The *in vitro* bioassay method for investigating MIC-based structure activity relationship of different synthesized compounds is one of the promising methods in

the early stage of drug discovery process. Proposed general protocol for agar dilution method is presented as follows:

#### (1) Preparation of inoculums

Inoculums were prepared by scraping colonies of exponential growth *M.tuberculosis* H37Rv from solid medium. Colonies were dispersed in a 20x150 mm screw-cap tube containing 2-3 ml of Middlebrook7*H9* broth supplemented with 10% oleic acid-albumin-dextrose-catalase (OADC) and 0.05% Tween 80 and 5-8 glass beads (diameter of 6 mm). The mixture was vortexed for 2-3 min, and standed for 20 min to allow large clumps of cells to settle. Cell suspension was transferred to a new sterile tube and adjusted to a turbidity equivalent to a McFarland Standard No.1 (ca. 3 x  $10^7$  cells/mL) with *M7H9* broth. Then, it was diluted to reach standard inoculums (ca. 1.5 x  $10^5$  cells/mL).

# (2) Preparation of test compounds and antibiotic-containing Middlebrook7H10 agar dilution plates

Middlebrook 7H10-10% oleic acid-albumin-dextrose-catalase (OADC) (Difco, USA) agar plates with different concentrations of test compounds (Final concentration: 20, 40 or 80 µg/mL) and kanamycin was used as positive controls.

#### (3) Inoculation of 7H10 agar plates

McFarland (1.5 x  $10^5$  cells) of each *M.tuberculosis* strain standard inoculums 5 µl was spotted onto both test compound, antibiotic containing and drug-free control *7H10* agar plates. Plates were sealed in CO<sub>2</sub>-permeable polyethylene bags and incubated at 37 °C for 3 weeks.

# (4) Determination of MIC

MIC was determined as the concentration of not detecting any colony (% resistance bacteria) compared with the control plate without drug.

% Resistance bacteria = Number of colonies on drug containing medium x 100

Number of colonies on drug free medium

# CHAPTER 4 RESULTS AND DISCUSSION

This research was included two experimental portions i.e. chemical synthesis and activity testing for antituberculosis activity. The scope of the research project was illustrated in Figure 22. Total fourteen benzhydrol derivatives compounds were designed and synthesized. The chemical structures, molecular weights and formulae of proposed benzhydrol compounds are displayed in Table 1. In order to synthesize those compounds, firstly, substituted benzoic acids were used as starting materials and their carboxylic acids were coupled with phenol to form esters by using EDC·HCl and DMAP as catalysts. The benzoate derivatives were changed to 4-hydroxy benzophenone by performing Fries rearrangement reaction. In order to determine the heating energy of the combustion in Fries rearrangement reaction using AlCl<sub>3</sub> as catalyst for synthesis of benzophenone, DSC was used to investigate the optimum temperatures for progression of the reaction and show how the products evolve with respect to temperature. After that, 4-hydroxy benzophenones were converted to benzhydrols by reduction using NaBH<sub>4</sub> to give benzhydrol derivatives. Based on their significant anti-TB properties, benzhydrol derivatives (4a, 4c, and 4f) were chosen as starting materials for the synthesis of diacetate compounds (5a, 5c, and 5f) Diacetylation was performed by acetylating hydroxyl groups of benzhydrol derivatives using acetic anhydride as an acetylating agent. All synthesized target compounds were purified by column chromatography and elucidated by means of FT IR, <sup>1</sup>H NMR, <sup>13</sup>C NMR and MS spectroscopic methods. General reaction pathways for synthesizing compounds 4a-f, 5a, 5c, 5f, 9, 10 and 13a-c were shown in Scheme 1-3. Then, the synthesized target compounds 3a-b, 4a-f, 5a, 5c, 5f, 9, 10 and 13a-c were tested for their antituberculosis activities.



Scheme 1 : esterification + Fires rearrangement + reduction Scheme 2 : base-catalyzed hydrolysis

Scheme 3 : amide formation + reduction

Figure 22 The scope of research project.

#### **4.1 Chemical synthesis**

#### 4.1.1 Phenyl benzoate compounds (2a-f)



For this reaction, substituted benzoic acids (1a-f) were used as starting materials to afford compounds 2a-f. The aim of this step was to preparing esters before the next step i.e. Fries Rearrangement reaction. The reaction was accelerated by the presence of carbodiimide activator (EDC·HCl) with catalytic amount of DMAP in dried dichloromethane. The reaction was completed at room temperature after 12-24 h to afford compound 2a-f in 78.9 to 96.7 % yield. The reaction mechanism for esterification was described in Figure 23.[91] The ester compounds were identified by IR spectrum to confirm ester functional groups.



IR spectrum of phenyl benzoate compounds showed characteristic vibration bands of stretching of functional groups. The bands of ester moiety were displayed carbonyl C=O stretching at 1735 cm<sup>-1</sup>, C-O stretching at 1269 cm<sup>-1</sup> and 1159 cm<sup>-1</sup>.



Figure 23 The reaction mechanism of the esterification.

#### 4.1.2 Benzophenone compounds (3a-m)



This step is one of the critical reaction steps of the chemical synthesis in this study. Fries rearrangement reaction was chosen to create the formation of benzophenone compounds. In the previous experiment, benzhydrol compounds were prepared by coupling reaction between aryl bromide and aryl aldehyde using organolithium reagents as reported in [92] with overall yield of 52.7%. However, it was difficult to control the reaction to acquire the desirable product since the reaction needed to be done at very low temperature and the reactive aryl anion intermediate was easily decomposed when exposed to moisture. Using Fries rearrangement, a more straightforward synthesis method for synthesizing benzhydrol compounds derived from benzophenones with a higher yield has been developed in order to circumvent the aforementioned synthetic difficulty.

# 4.1.2.1 Fries rearrangement reaction investigation by DSC analysis

The Fries rearrangement reaction was convenient to use as a nonsolvent method for synthesis. In order to specify the appropriate temperature for Fries rearrangement reaction, the reaction mixtures of selected phenyl benzoates **2b**, **2e**, **2f** and AlCl<sub>3</sub> heated at a heating rate of 5 °C/min under nitrogen atmosphere were studied using a DSC 823e (Mettler Toledo, Switzerland). Phenyl benzoates **2b**, **2e**, and **2f** were representatively phenyl benzoate substrates for the Fries rearrangement reaction because they were synthesized with high yields. The mixtures of compounds **2b**, **2e**, **2f** and AlCl<sub>3</sub> were inspected by heating from 40 °C to 400 °C which cover endothermic and exothermic temperatures of these mixtures during the progress of the Fries rearrangement reaction. The obtained products were assayed by employing LC-

MS to determine the optimal temperatures in order to get the highest yields of compounds **3b**, **3e** and **3f**.

# DSC analysis method used for investigation of temperature on the Fries rearrangement reaction.

On the basic of calorimetry technique, DSC is a thermal analysis equipment examining the changes in physical properties of a sample upon increasing temperature against time. This technique has been used to gain insight the thermodynamics of reaction intermediates. [93] In this study, phenyl benzoates **2b**, **2e** and **2f** were subjected to react with AlCl<sub>3</sub> and determined by using DSC at heating from 40 °C to 400 °C. The range of temperatures used in the Fries rearrangement reaction covered endothermic and exothermic changes of these compounds. The resulted products were subjected to LC-MS analysis to determine the optimal temperatures for the preparation of compounds **3b**, **3e** and **3f**.



Figure 24 DSC curves of the mixtures of compounds 2b, 2e, 2f and AlCl<sub>3</sub>

Figure 24 illustrates the DSC curves of reaction mixtures of compounds 2b, 2e, 2f and AlCl<sub>3</sub>. The sharp endothermic peak shown at 64.0 °C is the melting point of compound 2b. The thermograph of compound 2b also includes two broad endothermic peaks about 79-95 °C and 126-157 °C. The DSC curves of compounds 2e and 2f show the endothermic peaks at about 121-137°C and 146-156 °C and at about 133-148 °C and 173-180 °C, respectively. The endothermic peaks may correspond to the interaction between phenyl benzoates and AlCl<sub>3</sub>. The exothermic peaks at about 177-183 °C and 184-205 °C of compounds 2e-f, consecutively, are related to their thermal decomposition [74].

 Table 3 % Compositions of Fries rearrangement products by LC-MS analysis at variable temperatures

Starting		% Compositions <sup>a</sup>					
materials	Compounds	Temperature of Fries rearrangement reaction (°C)					
(Compounds)		60	-95	130	140	180	190
2b	3b	$4.24\pm0.76$	55.77± 8.55	$60.80\pm3.18$	$44.39\pm0.72$	$46.89 \pm 7.72$	$43.34 \pm 1.86$
	2b	$90.90 \pm 4.40$	$1.32\pm0.50$	$1.93\pm0.19$	$12.63\pm0.67$	$4.68\pm3.83$	$4.69\pm0.25$
2e	3e		$21.83 \pm 1.71$	$69.70\pm4.70$	$84.28 \pm 4.75$	$24.72\pm3.41$	$17.18 \pm 1.18$
	2e	$92.80\pm2.76$	$32.07\pm2.97$	25	Ŋ.	-	$6.45\pm0.79$
2f	3f		21.23 ± 1.95	27.21 ± 2.60	$17.42 \pm 1.67$	$2.93 \pm 1.66$	$10.54\pm3.15$
	2f	$93.23 \pm 5.94$	$30.56\pm3.53$	$18.42\pm6.45$	$36.03 \pm 4.48$	$64.65\pm2.81$	$14.44\pm2.05$

<sup>a</sup> % Compositions represent means ± SD of triplicate samples from three independent experiments.

As displayed in Table 3, raising the reaction temperature from 140 °C to 180 °C resulted in extremely lower yields of products **3b**, **3e**, **3f** while yield of product **3b** increased a few percent. This could have been percieved from the presupposition that amide functional groups existed in compounds **2e-f** and **3e-f** might reacted with the excess chloride of AlCl<sub>3</sub> and such changes in enthalpy leading to exothermic reactions.[94] Therefore, the exothermic peaks in the DSC curve of amido compounds **2e-f** and **3e-f** could be evidences of the product decomposition. [94-96] This supposition was attested by the fact that fluoro compounds **2b** and **3b** with no amido group did not show an exothermic process in the DSC thermograph.

According to DSC and LC-MS results, DSC curves could inform that Fries rearrangement reactions between ester compounds and AlCl<sub>3</sub> commenced at endothermic temperature, followed by the decomposition of products occurring at exothermic temperature. As published works, the effective temperature for running Fries rearrangement of phenyl benzoate compounds (**2b**, **2e**, **2f**) was assumed around 130-140°C for 1 hour. Benzophenone compounds (**3b**, **3e**, **3f**) were generated from this condition with high to moderate percent yield (33.7 - 78.9 %).[97] The reaction mechanism of this step was described in Figure 25. As the results of this study, we assumed the optimal reaction temperatures of Fries rearrangement of phenylbenzoate compounds to attain high percent yields for the preparation of benzophenone derivatives (3a-f, 8) were 130-140 °C for 30-60 min as shown in Table 4.

The mechanism of the Fries rearrangement reaction is illustrated in Figure 25. The acylium carbocation starts to attack the aromatic ring via an electrophilic aromatic substitution reaction. It is noted that the orientation of this electrophilic aromatic substitution is temperature-dependent. Low reaction temperatures favor para position substitutions while relatively high temperatures favor ortho position substitutions. [98-100] The obtained benzophenone derivatives were identified as *para*- isomers using the <sup>1</sup>H-NMR spectrum.



There are two signals of proton on aromatic ring A showed as representative *para-* substitution of hydroxyl group. The signal of the symmetry protons at 2 and 6 positions showed at the chemical shift of 7.03 ppm as a doublet with a coupling constant of 8.8 Hz, the same as the coupling constant of the proton signal at 3 and 5 positions was 8.8 Hz at the chemical shift of 6.89 ppm. The appearance of two symmetry signals of protons on aromatic ring A implies that the obtained product was a para-isomer.



Figure 25 The reaction mechanism of Fries rearrangement.

Entry	Compd	Product	Temp	Time	Yield	Physical
			(°C)	(min)	(%)	state
1	2a	3a	140	45	33.7	white powder
2	2b	3b	140	45	47.5	white powder
3	2c	3c	140	30	65.1	pale-yellow powder
4	2d	3d	130	30	78.9	white powder
5	2e	3e	140	30	44.9	pale yellow liquid
6	2f	3f	140	60	43.1	pale yellow liquid
7	7	8	130	60	59.7	Off-white powder

**Table 4** The reaction conditions and percent yields of compounds 3a-f and 8.

### 4.1.3 Benzhydrol compounds (4a-f) and 9

The aim of this step is to convert from ketone functional group of compounds **3a-f** and **8** to hydroxyl group of compounds **4a-f** and **9**. The ketone compounds can be reduced by the hydride nucleophile of sodium borohydride (NaBH<sub>4</sub>). First, hydride anion attacked carbonyl to generate an alkoxide anion intermediate. Then, the alkoxide was protonated by methanol as a proton source to provide secondary alcohol. In the reduction by sodium borohydride hydrolysis occurs automatically when methanol is used as a solvent. [101] The reaction mechanism of reduction step was described in Figure 26. This reaction created the stereogenic carbon, and the alcohol products were assumed to be a racemic mixture due to the low effect of steric hindrance surrounding carbonyl carbon. In a further study, chiral benzhydrol derivatives can be determined the degree of enantiomers excess by using a polarimeter.



Figure 26 The reaction mechanism of sodium borohydride reduction

## 4.1.4 Diacetyl benzhydrol compounds (5a, 5c, 5f)

The aim of this step was to change the active hydroxyl group of compound 4hydroxy- $\alpha$ -(phenyl)benzyl alcohol (**4a**), 4-hydroxy- $\alpha$ -(4'-bromophenyl)benzyl alcohol (**4c**) and 4-hydroxy- $\alpha$ -(4'-acetamido phenyl)benzyl alcohol (**4f**) to ester group in order to mimic the structure of ACA (I), chemical constituent of *Alpinia galanga*. For acetylation, alcohol was acylated using acetic anhydride as a source of acyl groups. Carbodiimide activator (EDC) with a catalytic quantity of DMAP in dichloromethane enhanced the process. The reaction was completed at room temperature for 12 hours to obtain compound **5a**, **5c**, **5f** with yields of 69.3 to 92.9 percent.

## 4.1.5 4-hydroxy-α-(4'-amino phenyl)benzyl alcohol (10)

The aim of this step is to cleave the acetyl functional group from compound **4f** to afford the amino compound **10**. The carbonyl carbon of the amide can be removed by acid or base hydrolysis. Normally, amide is difficult to be hydrolyzed; even heating for a long time, amide does not react with water molecules to decompose. However, amide can be interacted with water molecules either in an acidic or a basic medium. In a basic medium, amide is hydrolyzed to attain a carboxylic acid and the salt of ammonia or amine salt. The mechanism of base-catalyzed hydrolysis is shown in Figure 27. The reaction mechanism of amide hydrolysis in basic medium starts with the nucleophilic attack at the carbonyl carbon by a hydroxide ion to form a tetrahedral alkoxide intermediate. This step is reversible intermediate. The carbonyl
bond is reformed along with the elimination of an amide ion (-NHR) leaving group, forming a carboxylic acid. In the final step, the amide ion is protonated by a proton from the carboxylic acid to form a carboxylate salt and an amine as products.[102]



Figure 27 The reaction mechanism of the base-catalyzed hydrolysis

#### 4.1.6 Amidobenzhydrol compounds (13a-c)

The aim of this step was to link the various types of functional groups (i.e. alkyl, aryl and long chain) to amine group by amide bond in order to explore the binding target of anti-TB. For amide formation, the substituted acids (**11a-c**) were used as reagents for coupling with 4-aminobenzopheone. The reaction was accelerated in the presence of carbodiimide activator (EDC) with catalytic amount of DMAP in CH<sub>2</sub>Cl<sub>2</sub> [103]. The reaction was completed at room temperature after 12 h to afford compound **12a-c** as off-white solid in 31.5 to 72.0 % yield. The reaction mechanism for amide formation was described in Figure 28 [91]. Next, converting the ketone group to benzhydrol compounds (13a-c) was done by reduction reaction using NaBH<sub>4</sub> as hydride nucleophile.



Figure 28 The reaction mechanism of the amide formation

Entry	Com-	Product	<b>Overall Yield</b>	Physical	
	pound		(%) <sup>a</sup>	state	
1	4a	но	20.26	white solid	
2	4b	HO	26.98	white solid	
3	4c	HO	56.59	pale yellow solid	
4	4d	но	49.71	pale yellow liquid	
5	4e	HO HH HVAC	31.41	pale yellow liquid	
6	4f	HO HO AC	23.97	pale yellow liquid	
7	5a	Aco	18.82	white solid	
8	5c	Acto Br	- 39.22	pale yellow solid	
9	5f	Aco Aco	21.96	pale yellow liquid	
10	9	но	40.54	off-white solid	
11	10	HO NH <sub>2</sub>	6.57	pale yellow solid	
12	13a	OH NH NH	57.53	off-white solid	
13	13b	OH N N N	29.74	off-white solid	
14	13c		24.26	off-white solid	

Table 5 The physical properties and percent yield of benzhydrol derivatives

<sup>a</sup> The overall yield of this convergent synthesis was calculated based on overall steps starting from compound 1 or 6 or 11 to afford final products

#### 4.2 Structural characterizations

#### **4.2.1 IR spectroscopic characterization**

IR spectroscopy was used to determine the functional group and possibly identify components of benzhydrol derivatives. IR spectra of some benzhydrol derivatives were recorded (Neat or KBR pellets or nujol mullets) on a Jasco FT-IR spectrophotometer (Japan) and a Thermo Scientific Nicolet FI IR spectrophotometer (USA) reported as a wavenumber (cm<sup>-1</sup>).

#### 4.2.1.1 IR Characterization of 4-hydroxy-α-(phenyl)benzyl alcohol (4a)

Characteristic vibration bands of functional groups were displayed in IR spectrum of 4-hydroxy- $\alpha$ -(phenyl)benzyl alcohol (4a). A spectrum of 4a was used as a representative spectrum to explain the compound's significant vibration bands of functional groups. The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed at 3399 and 3165 cm<sup>-1</sup> (strong). The strong aromatic C-O stretching bands was 1611 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C=C stretching band at 1513 and 1448 cm<sup>-1</sup>, C-O stretching band of alcohol at 1005 cm<sup>-1</sup>, =C-H out of plane bending at 698 cm<sup>-1</sup>.

### 4.2.1.2 IR Characterization of 4-hydroxy-α-(4'-fluorophenyl)benzyl alcohol (4b)

The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed at 3386 and 3180 cm<sup>-1</sup> (broad). The strong C-O stretching band of aromatic was 1600 cm<sup>-1</sup>. The bands of the aromatic ring included aromatic C=C stretching band at 1510 and 1454 cm<sup>-1</sup>. The strong C-F stretching band was indicated at 1230 cm<sup>-1</sup>.

# 4.2.1.3 IR Characterization of 4-hydroxy-α-(4'-bromophenyl)benzyl alcohol (4c)

The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed at 3385 and 3153 cm<sup>-1</sup> (strong). The bands of aromatic ring included aromatic C=C stretching band at 1510 and 1453 cm<sup>-1</sup>. The strong C-O stretching bands of alcohol and strong C-Br stretching bands were indicated in the region of 1002 and 554 cm<sup>-1</sup>.

# 4.2.1.4 IR Characterization of 4-hydroxy-α-(4'-butylphenyl)benzyl alcohol (4d)

The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed at 3400 and 3168 cm<sup>-1</sup> (broad). The aliphatic C-H stretching bands in butyl were seen at 2958, 2925 and 2856 cm<sup>-1</sup>. The aliphatic C-H bending vibration was observed at 1362 and 1225 cm<sup>-1</sup>. The strong C-O stretching bands of alcohol showed at 1001 cm<sup>-1</sup>.

### 4.2.1.5 IR Characterization of 4-hydroxy-α-(3'-acetamido phenyl)benzyl alcohol (4e)

The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed at 3317 and 3190 cm<sup>-1</sup> (broad). The strong C-O stretching bands of alcohol showed at 995 cm<sup>-1</sup>. The strong C=O stretching at 1674 cm<sup>-1</sup>, N-H stretching in the region of approximate 3300 cm<sup>-1</sup> and N-H bending at 1605 cm<sup>-1</sup> indicated the presence of amide functional group.

### 4.2.1.6 IR Characterization of 4-hydroxy-α-(4'-acetamido phenyl)benzyl alcohol (4f)

The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed in the region at 3296 and 3103 cm<sup>-1</sup> (broad). The strong C-O stretching bands of alcohol showed at 1016 cm<sup>-1</sup>. The strong C=O stretching at 1666 cm<sup>-1</sup>, N-H stretching in the region of approximately 3300 cm<sup>-1</sup> and N-H bending at 1627 cm<sup>-1</sup> indicated the presence of an amide functional group.

#### 4.2.1.7 IR Characterization of 4-acetoxy-α-(phenyl)benzyl acetate (5a)

The characteristic C=O stretching of acetyl groups was displayed at 1824, 1739 cm<sup>-1</sup> (strong). The aliphatic C-H stretching bands of methyl at 2935 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C-H stretching at 3064, 3034 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C=C stretching band at 1505 and 1455 cm<sup>-1</sup>.

# 4.2.1.8 IR Characterization of 4-acetoxy-α-(4'-bromo phenyl)benzyl acetate (5c)

The characteristic C=O stretching of acetyl groups was displayed at 1761, 1741 cm<sup>-1</sup> (strong). The aliphatic C-H stretching bands of methyl at 2922, 2895 and 2804 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C-H stretching at about

 $3000 \text{ cm}^{-1}$ . The bands of aromatic ring included aromatic C=C stretching band at 1491 and 1448 cm<sup>-1</sup>. The C-Br stretching bands were indicated at 559 cm<sup>-1</sup>.

# 4.2.1.9 IR Characterization of 4-acetoxy-α-(4'-acetamido phenyl)benzyl acetate (5f)

The characteristic C=O stretching of acetyl groups was displayed at 1726, 1676 cm<sup>-1</sup> (strong). The aliphatic C-H stretching bands of methyl at 2983 and 2771 cm<sup>-1</sup>. The strong C=O stretching at about 1676 cm<sup>-1</sup>, N-H stretching at 3324 cm<sup>-1</sup> and N-H bending in the region of approximate 1600 cm-1 indicated the presence of amide functional group. The bands of aromatic ring included aromatic C-H stretching at 3068 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C=C stretching band at 1520 and 1408 cm<sup>-1</sup>.

#### 4.2.1.10 IR Characterization of 4-hydroxyphenyl(2-naphthyl)methanol (9)

The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed at 3386 and 3164 cm<sup>-1</sup> (broad). The strong C-O stretching bands of alcohol showed at 1024 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C-H stretching bend at about 3000 cm<sup>-1</sup>, skeletal C=C stretching at 1511 and 1456 cm<sup>-1</sup>, =C-H out of plane bending at 785 cm<sup>-1</sup>.

#### 4.2.1.11 IR Characterization of 4-hydroxy-α-(4'-amino phenyl)benzyl alcohol (10)

The bands of amine functional group included bands of N-H stretching at about 3300 cm<sup>-1</sup>, C-N stretching at 1081 cm<sup>-1</sup> with medium-weak intensity, medium intensity of N-H bending at 1618 cm<sup>-1</sup>. The characteristic O-H stretching for intramolecular hydrogen bonding of alcohol was displayed around 3329 cm<sup>-1</sup> (broad). The strong C-O stretching bands of alcohol showed at 1022 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C-H stretching bend at about 3024 cm<sup>-1</sup>, skeletal C=C stretching at 1515 and 1450 cm<sup>-1</sup>. =C-H out of plane bending at 748 cm<sup>-1</sup>.

# 4.2.1.12 IR Characterization of 4-butyramido-α-(phenyl)benzyl alcohol (13a)

The bands of amide functional group included N-H stretching (strong) at about 3300 cm<sup>-1</sup>, C=O stretching (strong) at 1660 cm<sup>-1</sup>, N-H bending at 1601 cm<sup>-1</sup>. The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was

displayed around 3303 cm<sup>-1</sup> (broad). The strong C-O stretching bands of alcohol showed at 1016 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C-H stretching (medium) at 3057 cm<sup>-1</sup>, C-H out of plane bending at about 727 cm<sup>-1</sup>, skeletal C=C stretching at 1533 and 1448 cm<sup>-1</sup>. The aliphatic sp<sup>3</sup> C-H stretching bands at 2962, 2873 cm<sup>-1</sup> and aliphatic C-H bending vibration at 1309, 1271 cm<sup>-1</sup> indicated the presence of butyl groups.

## 4.2.1.13 IR Characterization of 4-benzamido-α-(phenyl)benzyl alcohol (13b)

The bands of amide functional group included N-H stretching (strong) at about 3300 cm<sup>-1</sup>, C=O stretching (strong) at 1651 cm<sup>-1</sup>, N-H bending at 1599 cm<sup>-1</sup>. The characteristic O-H stretching for intra-molecular hydrogen bonding of alcohol was displayed around 3319 cm<sup>-1</sup> (broad). The strong C-O stretching bands of alcohol showed at 1012 cm<sup>-1</sup>. The bands of aromatic ring included aromatic C-H stretching (medium) at 3060, 3030 cm<sup>-1</sup>, skeletal C=C stretching at 1523-1446 cm<sup>-1</sup>, =C-H out of plane bending at about 731 cm<sup>-1</sup>

#### 4.2.1.14 IR Characterization of 4-cinnamido-α-(phenyl)benzyl alcohol (13c)

The bands of amide functional group included N-H stretching (strong) at about 3300 cm<sup>-1</sup>, C=O stretching (strong) at 1658 cm<sup>-1</sup>, N-H bending at 1601 cm<sup>-1</sup>. The O-H stretching at about 3313 cm<sup>-1</sup> (broad), strong C-O stretching bands at 1012 cm<sup>-1</sup> revealed the presence of alcohol group. In the spectrum, unsaturated aliphatic C=C stretching at 1533 cm<sup>-1</sup>, the band of =C-H bending (strong) at 729 cm<sup>-1</sup>, the bands of aromatic ring included aromatic C-H stretching (medium) at 3058, 3029 cm<sup>-1</sup> were indicated the existence of cinnamic group.

					Way	venumber	: (cm <sup>-1</sup> )					
Compd	N-H st.	H-O	Aromatic C-H st.	Aliphatic C-H st.	C=0	N-H hend	Aromatic C-O st.	Aromatic C=C st.	Aliphatic C-H bend	C-O	=C-H bend	etc.
4a		3399, 3165	app 3000	)   			1611	1513, 1448		1005	698	
4b		3386, 3180	app 3000		ı		1600	1510, 1454		1007	783	1230 (C-F st.)
4c		3385, 3153	app 3000	I	I	ı	1600	1510, 1453	I	1002	785	554 (C-Br st.)
4d		3400, 3168	app 3000	2958, 2925 , 2856	I	I	1597	1512, 1452	1362, 1225	1001	775	
4e	app 3300	3317, 3190	app 3000	2810	1674	1605	1587	1512, 1441	1331, 1284	995	764	
4f	app 3300	3296, 3103	app 3000	2910	1666	1627	1599	1525, 1404	1371, 1288	1016	771	
Sa		1	3064, 3034	2935	1824,1 739	I	1607	1505, 1455	1370, app 1200	1016	747	
Sc		1	app 3000	2922, 2895 , 2804	1761,1 741	I	1620	1491, 1448	1369, 1234	1009	775	559 (C-Br st.)
Sf	3324	1	3068	2983, 2771	1726, 1676	app 1600	1592	1520, 1408	1369, 1268	1000	767	
6		3386, 3164	app 3000	I	ı	ı	1599	1511, 1456	I	1024	785	
10	app 3000	3329	3024	I	ı	app 1600	1617	1515, 1450	I	1022	748	1081 (C-N st.)
13a	app 3300	3303	3057	2962, 2873	1660	1601	I	1533, 1448	1309, 1271	1016	727	
13b	app 3300	3319	3060, 3030	I	1651	1599	I	1523, 1446	I	1012	731	ı
<b>13</b> c	app 3300	3313	3058, 3029	I	1658	1601	ı	1495, 1448	I	1012	729	1533 (C=C st. cinnamic)

Table 6 Characteristic IR bands of important functional groups of synthesized compounds

68

#### 4.2.2 NMR spectroscopic characterization

NMR spectroscopy was used to determine the structure of benzhydrol derivatives. All of the <sup>1</sup> H, <sup>13</sup>C NMR spectra were recorded on a Bruker-Ultra Shield (300 and 100 MHz for <sup>1</sup> H and <sup>13</sup>C, respectively), using CD<sub>3</sub>OH or CD<sub>3</sub>COCD<sub>3</sub> as solvents with trace of CH<sub>3</sub>OH or TMS as internal standards.

#### 4.2.2.1 NMR Characterization of 4-hydroxy-α-(phenyl)benzyl alcohol (4a)

The <sup>1</sup>H NMR spectroscopy was used to determine the structure of benzhydrol (**4a**) using CD<sub>3</sub>COCD<sub>3</sub> as a solvent with TMS as an internal standard. The C-H proton  $\alpha$  to the alcohol showed singlet signal at 5.73 ppm. The resonances of aromatic protons at positions of 3, 5 and 2, 6 showed at 6.76 ppm and 7.40 ppm as doublet splitting patterns with coupling constant 8.7 Hz, respectively. The chemical shifts of protons at 2', 6' appeared in the aromatic region of 7.26 to 7.32 ppm as multiplet. The resonances of 3', 4 ' and 5' protons appeared in the more upfield region between 7.16 to 7.22 ppm as multiplet.

#### 4.2.2.2 NMR Characterization of 4-hydroxy-α-(4'-fluorophenyl)benzyl alcohol (4b)

The C-H proton  $\alpha$  to the alcohol showed singlet signal at 5.69 ppm. The resonances of aromatic protons at positions of 3, 5 and 2, 6 showed at 6.74 ppm and 7.15 ppm as doublet splitting patterns with coupling constant 8.6 Hz and 8.3 Hz, respectively. The chemical shifts of protons at 2', 6' appeared in the aromatic region of 6.98 to 7.05 ppm as triplet. The resonances of 3' and 5' protons appeared in the more downfield region between 7.32 to 7.37 ppm as multiplet due to electron withdrawing effect of fluorine.

The <sup>13</sup>C NMR spectra showed the carbon of aromatic ring in the region of 115.71 - 165.01 ppm. The  $\alpha$ -carbon to OH group appeared at 76.07 ppm

### 4.2.2.3 NMR Characterization of 4-hydroxy-α-(4'-bromophenyl)benzyl alcohol (4c)

The C-H proton  $\alpha$  to the alcohol showed singlet signal at 5.73 ppm. The resonances of aromatic protons at positions of 3, 5 and 2, 6 showed at 6.77 ppm and 7.20 ppm as doublet splitting patterns with coupling constant 8.4 Hz. The chemical

shifts of protons at 2', 6' appeared in the aromatic region of 7.35 ppm as doublet. The resonances of 3' and 5' protons appeared in the more downfield region between 7.47 ppm as doublet due to electron withdrawing effect of bromine with the same coupling constants of 8.4 Hz.

### 4.2.2.4 NMR Characterization of 4-hydroxy- $\alpha$ -(4'-butylphenyl)benzyl alcohol (4d)

The signal of 4" proton at butyl group showed at the chemical shift of 0.92 ppm as triplet with coupling constant 7.3 Hz. Two protons at position 3" appeared resonance at 1.34 ppm as sextet resulting from the coupling of neighboring protons with coupling constant 7.6 Hz. The splitting pattern of 2" proton was quintet due to coupling of 1" and 3" protons and its signal appeared at 1.57 ppm with coupling constant 7.6 Hz. The proton of 1" showed chemical shift of 2.58 ppm as triplet with coupling constant 7.5 Hz. The C-H proton at  $\alpha$ -position to the alcohol showed singlet signal at 5.66 ppm. The resonances of two aromatic protons at positions of 3 and 5 showed at 6.73 ppm as doublet. Two protons at position 2' and 6' appeared signals at 7.11 ppm as doublet. The resonances of two aromatic protons at positions of 3' and 5' attached to butyl group showed at 7.23 ppm as doublet with coupling constant 8.0 Hz.

The <sup>13</sup>C NMR spectrum showed that the signals of carbons from butyl side chain appeared in the region of 14.42 ppm to 36.38 ppm. The signals of CH-OH carbon at  $\alpha$  to the alcohol showed at 76.66 ppm and aromatic ring carbons in the region of 116.04 to 157.70 ppm. The signals of CH-OH at aromatic rings appeared at 157.70 ppm

### 4.2.2.5 NMR Characterization of 4-hydroxy- $\alpha$ -(3'-acetamido phenyl) benzyl alcohol (4e)

The signal of  $CH_3$  proton at acetamido group showed at the chemical shift of 2.90 ppm as a singlet. The resonance of NH proton in acetamido group appeared at 4.65 ppm. The C-H proton at  $\alpha$ -position to the alcohol showed a singlet signal at 5.69 ppm. The resonances of two aromatic protons at positions 3 and 5 showed 6.76 ppm as a doublet with a coupling constant of 9.0 Hz. Two protons at positions 2 and 6 appeared signals at 7.58 ppm. The characteristic chemical shift for 2' proton in

aromatic ring attached to acetamido side chain showed at 7.08 ppm as doublet due to coupling with NH proton. The chemical shift of three protons of the aromatic ring at 4', 5' and 6' positions appeared in the region of 7.17 to 7.22 ppm.

### 4.2.2.6 NMR Characterization of 4-hydroxy- $\alpha$ -(4'-acetamido phenyl) benzyl alcohol (4f)

The signal of CH<sub>3</sub> proton at acetamido group showed the chemical shift of 2.94 ppm as singlet. The resonance of NH proton in acetamido group appeared at 4.82 ppm as a singlet. The C-H proton at  $\alpha$ -position to the alcohol showed a singlet signal at 5.73 ppm. The resonances of the symmetry protons at positions 3, 5 and 2, 6 on aromatic ring A showed at 6.77 ppm and 7.47 ppm as doublet splitting patterns with coupling constant 8.7 Hz. As doublet, the chemical shifts of protons at 3', 5' appeared in the aromatic region of 7.19 ppm. The resonances of 2' and 6' protons appeared in region 7.35 ppm as a doublet with the same coupling constants of 8.4 Hz.

#### 4.2.2.7 NMR Characterization of 4-acetoxy-α-(phenyl)benzyl acetate (5a)

The signal of methyl proton in CH-OCOCH<sub>3</sub> and Ar-OCOCH<sub>3</sub> showed at the chemical shift of 2.15 ppm and 2.28 ppm as singlets, respectively. The C-H proton at  $\alpha$ -position to the alcohol showed a singlet signal at 6.88 ppm. The characteristic resonances for aromatic ring A in 3,5-positions appeared at 7.05 ppm as doublet splitting patterns with coupling constant 8.7 Hz. The splitting pattern resulting from the coupling with neighboring protons H-2, H-6, H-2', H-3', H-4', H-5' and H-6' was multiplet, and chemical shift appeared in the region of 7.26-7.36 ppm.

### 4.2.2.8 NMR Characterization of 4-acetoxy- $\alpha$ -(4'-bromophenyl)benzyl acetate (5c)

The signal of methyl proton in CH-OCOC<u>H</u><sup>3</sup> and Ar-OCOC<u>H</u><sup>3</sup> showed the chemical shift of 2.15 ppm and 2.28 ppm as singlets. The C-H proton at  $\alpha$ -position to the alcohol showed a singlet signal at 6.82 ppm. The resonances of aromatic protons at positions 3, 5 and 2, 6 showed at 7.06 ppm and 7.31 ppm as doublet splitting patterns with the same coupling constants of 8.7 Hz. The chemical shifts of protons at 2', 6' and 3', 5' appeared in the aromatic region of 7.20 ppm and 7.46 ppm as doublets with the same coupling constants of 8.4 Hz. The resonances of 3' and 5' protons

appeared in the more downfield region due to the electron-withdrawing effect of bromine.

#### **4.2.2.9 NMR Characterization of 4-hydroxyphenyl (2-naphthyl)** methanol (9)

The <sup>1</sup>H NMR spectroscopic data showed that the characteristic resonances for C-H proton at  $\alpha$ -position to the alcohol appeared singlet signal at 5.86 ppm. The resonances of aromatic protons at positions of 3, 5 and 2, 6 showed at 6.75 ppm and 7.21 ppm as doublet splitting patterns with the coupling constants of 8.6 and 8.4 Hz, respectively. The resonances of three naphthalene protons at the positions of 2', 9' and 10' showed at 7.43 ppm as multiplet. Four protons of naphthalene ring at position 4', 5', 6', and 7' appeared signals at the region between 7.75 to 7.86 ppm as multiplet.

The <sup>13</sup>C NMR spectrum showed that the signals  $\alpha$ -carbon to OH group appeared at 76.86 ppm. The carbons of aromatic rings were found in the region of 116.18 -157.90 ppm.

4.2.2.10 NMR Characterization of 4-butyramido-α-(phenyl)benzyl alcohol (13a)

The signal of CH<sub>3</sub> proton at 3" position of butyramide group showed at the chemical shift of 0.98 ppm as triplet with coupling constant 7.4 Hz. The splitting pattern of 2" proton (CH<sub>2</sub>) was sextet due to coupling of 1" and 3" protons and its signal appeared at 1.70 ppm with coupling constant 7.4 Hz. The proton of 1" showed chemical shift of 2.32 ppm as triplet with coupling constant 7.4 Hz. The proton of  $\alpha$ -carbon to OH group appeared at 5.74 ppm as a singlet. The resonances of five protons at aromatic ring-A and two protons at 2', 6' positions of aromatic ring-B showed in the region of 7.18-7.37 ppm as multiplet. The chemical shifts of protons at 3', 5' appeared in the aromatic region of 7.50 ppm as doublet. The differences of chemical shift of aromatic protons at 3', 5' positions were affected by inductive effect of amide group at *para*- positon on aromatic ring.

The <sup>13</sup>C NMR spectrum showed that the signals  $\alpha$ -carbon to OH group appeared at 76.68 ppm. The carbons of aromatic rings were found in the region of 121.25 -145.98 ppm. The carbon of amide group showed C=O signal in the most downfield at 174.68 ppm

## 4.2.2.11 NMR Characterization of 4-benzamido-α-(phenyl)benzyl alcohol (13b)

The C-H proton  $\alpha$  to the alcohol showed singlet signal at 5.78 ppm. The resonances of seven protons at aromatic ring A (H-2, H-3, H-4, H-5, H-6) and ring B (H-2', H-6') showed in the region of 7.20 – 7.40 ppm as multiplet splitting patterns. The doublet signal at 7.64 ppm was the chemical shifts of protons at 3', 5' position of aromatic ring connected to amine group. The chemical shifts of protons at 3'', 4'', 5'' appeared in the aromatic region of 7.50 ppm as multiplet. The resonances of 2'' and 6'' protons appeared in the more downfield region of 7.92 ppm as doublet due to electron withdrawing effect of carbonyl group.

The <sup>13</sup>C NMR spectrum showed that the signals  $\alpha$ -carbon to OH group appeared at 76.73 ppm. The carbons of aromatic rings were found in the region of 122.32 -146.04 ppm. The carbon of amide group showed C=O signal in the most downfield at 169.04 ppm

4.2.2.12 NMR Characterization of 4-cinnamido-α-(phenyl)benzyl alcohol (13c)

The <sup>1</sup>H NMR spectroscopic data showed that the characteristic resonances for C-H proton at  $\alpha$ -position to the alcohol appeared singlet signal at 5.74 ppm. The resonance of H-a position of CH=CH of cinnamide group appeared at 6.78 ppm as two broad singlets from the unlabelled isotopomer. The signal of H-b position of CH=CH group appeared in the region of 7.55 – 7.63 ppm. The resonances of aromatic protons at ring A and ring C showed in the region of 7.20 – 7.40 ppm as multiplet splitting patterns. The chemical shifts of protons at 2', 3', 4', 5' at ring B appeared in the aromatic region of 7.55 - 7.63 ppm as multiplet.

The <sup>13</sup>C NMR spectrum showed that the signals  $\alpha$ -carbon to OH group appeared at 76.71 ppm. The unsaturated carbon including aromatics and alkenes showed in the region of 121.23 -146.02 ppm. The carbon of amide group showed C=O signal in the most downfield at 166.78 ppm

#### **4.3 Structure-activity relationship of synthesized compounds and antituberculosis activity**

The synthesized compounds **4a-f**, **5a**, **5c**, **5f**, **9**, **10** and compound **13a-c** were investigated for antituberculosis activity on 20 clinical isolates and MTB H37Rv reference strains (ATCC 27294) according to the procedure described in section 3.3. The results of *in vitro* antituberculosis activities are summarized in Table 7.

It could be concluded that the inductive effects of substituents (fluorine or bromine atom) on a B-ring which significantly enhanced the anti-TB activity compared with unsubstituted 4-hydroxy benzhydrol **4a**. Firstly, we synthesized and investigated the high electronegativity of fluorine or bromine atom on aromatic benzene ring for the structure activity relationship of benzhydrol derivatives. It was found that fluorobenzhydrol **4b** and bromobenzhydrol **4c** had greater anti-TB efficacy than compound **4a** against all 20 MTB clinical isolates and MTB H37Rv reference strains with a MIC of 40  $\mu$ g/ml.

Secondly, we examined the significance of the hydroxy group at the  $\alpha$ -carbon for anti-TB activity by substituting the hydroxy group with a ketone group. It was found that ketone derivatives **3a** and **3b** were less active than their hydroxy derivatives **4a** and **4b**, respectively. Thus, the hydroxy group at the  $\alpha$ -carbon was essential for biological activity.

Thirdly, in our previous work, we synthesized 4-hydroxy- $\alpha$ -(3'-bromophenyl)benzyl alcohol, **14**, and examined its anti-tuberculosis (TB) activity for the *meta*-substitution of the electron-withdrawing bromo group with a MIC of 80 µg/ml [92]. It showed lower anti-TB activity than compound **4c** with *para*-substituted bromine on B-ring (MIC of 40 µg/ml). As a result, the electronegativity effect of the bromine atom at *para* substitution on the B-ring could enhance the activities. In addition, the insertion of the acetamido group to either the *meta* or *para* position of the B-ring led to diminishing activity with MIC values >80 µg/ml.

Fourthly, we studied the effects of substitution of various functional groups on the B-ring on anti-TB activity. The compound with the highest activity was compound **4d**, which possessed an alkyl long chain substitution at *para*-position of Bring with MIC value of 20-40  $\mu$ g/ml. Furthermore, compound **13c** containing *trans*cinnamyl group at the *para*-position of ring B showed effective anti-TB activity with a MIC of 40  $\mu$ g/ml. These results showed that phenyl substitution with an alkyl linker could enhance the activity. However, the presence of bulky groups (i.e., *p*-benzamide substitution, naphthalene) of compounds **13b** and **9** decreased anti-TB effectiveness. These may be the results of the steric hindrance of compounds to access the target site.

Finally, we examined the effect of acetylation on hydroxyl groups of benzhydrol derivatives. The result revealed that diacetylated derivatives **5c** and **5f** failed to inhibit MTB. The MIC values of dihydroxy benzhydrols **4a**, **4c**, and **4f** were 40-80  $\mu$ g/ml; however, diacetylated analogue of **4a** (compound **5a**) showed better anti-TB activity than its dihydroxy compound, with a MIC value < 40  $\mu$ g/ml. The result of acetoxy substitution was reported in previous presentation [104].

The effects of the structural modification of benzhydrol derivatives on anti-TB activity could be concluded to the structure-activity relationship (SAR) as follows.

- 2. Substituting the hydroxyl group or leaving it unsubstituted (H atom) at the *para-position* of A-ring exhibited anti-TB activity.
- 3. The *para*-substitution of halide, such as fluoride or bromide atom at the Bring, could improve anti-TB activity.
- 4. The *para*-substitution of bromide showed higher anti-TB activity than the *meta*-substitution.
- 5. The presence of acetamido groups (NHCOCH<sub>3</sub>) at the B-ring reduced the activity.
- 6. Enlarging with the bulky aromatic ring scaffold on the B-ring caused low activities.
- 7. Elongating the benzhydrol core structure attached to the alkyl or phenyl long chains at the *para*-position led to higher activities.

In this series, the *n*-butyl compound (**4d**) exhibited the highest anti-TB activity with a MIC of 20-40  $\mu$ g/ml. A long chain alkyl substituents of compound **4d** may have more appropriate physical properties than other moieties for accessing the target site. The butyl elongation chains with flexible conformation were added to the benzhydrol scaffold which might reach into the binding pocket with a high or long cavity. Furthermore, the similarity between the lipophilic nature of the alkyl long chain in molecule **4d** and the high fatty acid content (mycolic acid) of the cell wall of tuberculosis strains might hint at the target of action. Therefore, one promising mechanism of action for benzhydrol **4d** might be the inhibition of mycolic acid biosynthesis. Substrates of mycolic acid biosynthesis containing a large number of carbon units interact with an enzyme involved in the elongation of long-chain fatty acids with hydrophobic interaction.[105] The lipophilic nature of alkyl in the proposed compound **4d** might interact with the target in mycolic acid synthesis.

The chemical constituent of ACA from natural source responsible for antituberculosis activity is S-configuration i.e., 1'-S-1'-Acetoxychavicol acetate. According to the literature, MIC of synthetic ACA was higher than natural S-form of ACA on *M. tuberculosis* stains (63). Therefore, it can be postulated that the racemic form of synthetic 4-hydroxy- $\alpha$ -(4'-butylphenyl)benzyl alcohol (4d) contributed to lower antituberculosis activity and chirality may play an important role in the intermolecular interaction with binding sites of biological targets.

In order to develop a new candidate for an anti-Tb drug, the logarithm of the partition coefficient between n-octanol and water (cLogP) calculation and molecular weight (MW) were used in defining drug ability.[106] Candidate **4d** was a small molecule with an MW value of 256.34 and showed hydrophobic property with a cLogP value of 3.871. The cLogP values of **4d** and **13c** were higher than clogP values of other benzhydrols due to their greater hydrophobicity. In addition, ACA modification data [43] indicated that compounds of greater lipophilicity were much more likely to have high antituberculosis activity. However, since their cLogP values are more than three, it is concerned that aqueous solubility would be limited.

According to the results, it is concluded that compound **4d** showed the highest potential in the series to be a new lead for further structural modification in an attempt to improve the antituberculosis activity and drug-like properties.



R В A R₁O 4a-f, 5a, 5c, 5f, 10,14  $\overline{QR}_1$ 

QR₁



Cpds	<b>R</b> 1	R <sub>2</sub>	MIC (µg/mL) <sup>a</sup>	MW	cLogP
3a	Н	H	80	199.23	3.054
3b	Н	F	80	217.22	3.272
3f	Н	NH-CO-CH <sub>3</sub>	> 80	256.28	2.213
4a	Н	4-н	40 - 80	200.23	1.785
4b	Н	4-F	40	218.22	1.928
4c	Н	4-Br	40	279.13	2.648
4d	H	4-C4H9	20 - 40	256.34	3.871
4e	H	3-NH-CO-CH <sub>3</sub>	>80	257.28	0.804
4f	Н	4-NH-CO-CH <sub>3</sub>	>80	257.28	0.804
5a	-CO-CH <sub>3</sub>	Н	< 40	284.31	2.657
5c	-CO-CH <sub>3</sub>	4-Br	Inactive	363.20	3.520
5f	-CO-CH <sub>3</sub>	4-NH-CO-CH <sub>3</sub>	> 80	341.36	1.676
9	Н	-	80	250.29	2.959
10	Н	4-NH2	-	215.25	0.558
13a	Н	C <sub>3</sub> H <sub>7</sub>	80	240.28	2.529
13b	Н	C <sub>6</sub> H <sub>6</sub>	80	303.35	2.960
13c	Н	CH=CH-C <sub>6</sub> H <sub>6</sub>	40 - 80	329.39	3.974
14	Н	3-Br	80	279.13	2.648
Kanamycin (control)			10	484.50	-3.88 <sup>b</sup>

Table 7 In vitro antitubercular activities of synthesized compounds against MTB<sup>a</sup>

<sup>a</sup> 20 MTB clinical isolates and MTB H37Rv reference strains (ATCC 27294). <sup>b</sup> Data is available on the website (http://drugcentral.org).

#### CHAPTER 5 CONCLUSION

The structure-based design and synthesis of benzhydrol derivatives were developed and assessed for their *in vitro* antituberculosis activity. The various substituents, such as halide atoms, alkyl, aryl, acetamido, and amide derivatives, were connected to the B-ring of benzhydrol structure. Three main structural groups were designed for this series. First, designing 4-hydroxybenzhydrols (**4a-f**, **9**, **10**) contained the hydroxy group at ring A, focusing on varying substituents to explore the target binding site of *M. tuberculosis* inhibition mainly on the B-ring. The second group was benzhydrol diacetates (**5a**, **5c**, **5f**) designed to mimic ACA structure containing acetoxy groups. The last group, 4-amidobenzhydrols (**13a-c**), were generated from connecting benzhydrol to various alkyl, aryl, or cinnamyl groups that might interact on binding pockets of target sites.

The synthesis pathways to afford designed benzhydrol derivatives 4a-f, 9 included three steps, the benzoic acid and substituted benzoic acid compounds were coupled with phenol by esterification reaction to form benzoate derivatives 2a-f and 7 provided 78.9 to 96.7 percent yields. Next, the connection between two benzene rings by a ketone group to form benzophenone derivatives was performed via the Fries rearrangement reaction. The optimal temperatures for the synthesis of Fries rearrangement products with moderate to good yields (33.7 - 78.9 %) were 120 to 140 °C investigated by DSC analysis. The DSC data were effectively applied to estimate the optimal reaction temperatures to attain high percent yields for the Fries rearrangement reaction. The reduction step was achieved by sodium borohydride in methanol within one hour with high percent yields, 59.3% to 89.9%. The overall percent yields of benzhydrol dervivatives 4a-f and compound 9 were 20.3% to 56.6%. Compound 10 was obtained from amide hydrolysis of 4f and provided low percent yields (6.57%). The synthesis of diacetyl products 5a, 5c and 5f was completed by acetylation of the compound 4a, 4c and 4f by EDC, DMAP and acetic anhydride with the overall yields of 18.8% to 39.2%. For the synthesis of acetamido products 13a-c, 4-aminobenzophenone was coupled with various acids by amide bond linkage, followed by NaBH<sub>4</sub> reduction with overall percent yields of 24.3% to 57.5%. All the structures of synthesized designed compounds were confirmed by FT-IR, <sup>1</sup>H NMR, and <sup>13</sup>C NMR methods and tested for antituberculosis activities.

According to MIC results, five compounds (**3a**, **3b**, **9**, **13a** and **13b**) had low activity against M. *tuberculosis* with MICs of 80 µg/ml. Compounds **4a**, **4b**, **4c**, **5a**, and **13c** with MICs between 40 and 80 µg/ml exhibited moderate activity. The highest efficacy of this series was compound **4d** with a MIC of 20-40 µg/ml. The chemical structures of active benzhydrol derivatives of this series are shown in Figure 29. The structure-activity relationship study revealed that the most potent compound for anti-TB activity was compound **4d** with alkyl long chain substituent at the *para*-position, at which alkyl side chain was similar to the fatty acid content of cell walls of tuberculosis strains. At the *para* position of the B-ring, electron-withdrawing groups like fluorine and bromine could provide electron-withdrawing groups that might interact with complementary of binding sites. Halide derivatives showed moderate to good potential to inhibit mycobacterial growth. The active 11 compounds could imply that the benzhydrol derivatives offered promising new leads for further development. Significantly, compound **4d** exhibited the highest potential for further structural modification as a new antituberculosis agent.

For additional research on benzhydrol's antituberculosis activity, the expansion of the alkyl/aryl chain could influence the activity. As previously stated, ACA data showed high activity in *S*-form. The chirality effect at the  $\alpha$ -carbon of benzhydrol is an intriguing topic for further study. Upon chiral separation of the products, we expect that one of the enantiomers could achieve higher activity than their racemates. According to the activities of benzhydrol derivatives against 20 clinical MTB strains, it is possible to combat MDR-TB; however, future research suggests modifying the substituent's chemical structure at the *para*-position on the B-ring. In order to obtain information on how benzhydrol derivatives bind to their targets, target enzymes involved in mycolic acid biosynthesis should be further investigated. Once the target enzyme is known further design on modification of benzhydrol derivatives based on their binding affinities a target enzyme could be done.

Although, benzhydrol derivatives showed good activity against mycobacterial tuberculosis *in vitro*, inhibitory properties may not correlate with antimycobacterial *in* 

vivo results. Next step of study should include in vivo testing since in vivo studies are essential to drug development to provide the information on drug's properties, including in pharmacokinetic and pharmacodynamic processes of drug action.



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#### **4-hydroxy-***α***-(phenyl)benzyl alcohol (4a)**



#### **IR** spectrum



<sup>1</sup>H NMR spectrum (300 MHz, CDCl3) of 4-hydroxy-α-(phenyl)benzyl alcohol (4a)



#### 4-hydroxy-α-(4'-fluorophenyl)benzyl alcohol (4b)



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<sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>OD) of 4-hydroxy-α-(4'-fluorophenyl)benzyl alcohol (4b)



<sup>13</sup>C NMR spectrum (75 MHz, CD<sub>3</sub>OD) of 4-hydroxy-α-(4'-fluorophenyl)benzyl alcohol (4b)



<sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>COCD<sub>3</sub>) of 4-hydroxy-α-(4'bromophenyl)benzyl alcohol (4c)





#### 4-hydroxy-α-(4'-butylphenyl)benzyl alcohol (4d)



<sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>OD) of 4-hydroxy-α-(4'-butylphenyl)benzyl alcohol (4d)
<sup>13</sup>C NMR spectrum (75 MHz, CD<sub>3</sub>OD) of 4-hydroxy-α-(4'-butylphenyl)benzyl alcohol (4d)





<sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>COCD<sub>3</sub>) of 4-hydroxy-α-(3'-acetamidophenyl)benzyl alcohol (4e)

## 4-hydroxy-α-(4'-acetamidophenyl)benzyl alcohol (4f)





## <sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>COCD<sub>3</sub>) of 4-hydroxy-α-(4'-acetamidophenyl)benzyl alcohol (4f)



#### 4-acetoxy-α-(phenyl)benzyl acetate (5a)





<sup>1</sup>H NMR spectrum (300 MHz, CDCl<sub>3</sub>) of 4-acetoxy-α-(phenyl)benzyl acetate (5a)

### 4-acetoxy-α-(4'-bromo phenyl)benzyl acetate (5c)









## 4-acetoxy-α-(4'-acetamido phenyl)benzyl acetate (5f)



## 4-hydroxyphenyl(2-naphthyl)methanol (9)





<sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>OD) of 4-hydroxyphenyl(2-naphthyl)methanol (9)



# <sup>13</sup>C NMR spectrum (75 MHz, CD<sub>3</sub>OD) of 4-hydroxyphenyl(2-naphthyl)methanol (9)



## 4-butyramido-α-(phenyl)benzyl alcohol (13a)









<sup>13</sup>C NMR spectrum (75 MHz, CD<sub>3</sub>OD) of 4-butyramido-α-(phenyl)benzyl alcohol (13a)



### 4-benzamido-α-(phenyl)benzyl alcohol (13b)





<sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>OD) of 4-benzamido-α-(phenyl)benzyl alcohol (13b)



<sup>13</sup>C NMR spectrum (75 MHz, CD<sub>3</sub>OD) of 4-benzamido-α-(phenyl)benzyl alcohol (13b)



<sup>1</sup>H NMR spectrum (300 MHz, CD<sub>3</sub>OD) of 4-cinnamido-α-(phenyl)benzyl alcohol (13c)



<sup>13</sup>C NMR spectrum (75 MHz, CD<sub>3</sub>OD) of 4-benzamido-α-(phenyl)benzyl alcohol (13b)



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