ORIGINAL RESEARCH

Magnitude of Multidrug Resistance and Associated Factors of Pulmonary Tuberculosis Among Adult Smear Positive Patients in Eastern Ethiopia

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¹Harari Health Research and Regional Laboratory, Harar, Ethiopia; ²Department of Medical Laboratory Sciences, College of Health and Medical Science, Haramaya University, Harar, Ethiopia **Background:** In Ethiopia, multidrug resistant tuberculosis is a major public health problem. However, information is scarce regarding MDR-TB and associated factors.

Objective: The study was aimed to assess the magnitude of multidrug resistance and associated factors of pulmonary tuberculosis among adult smear-positive patients in Harari regional state health facilities, eastern Ethiopia.

Methods: A cross-sectional study was conducted among 395 adult smear-positive pulmonary tuberculosis patients attending health facilities from March to October 2019. Smearpositive sputum samples were collected from health facilities, and transported to Harari Health Research and Regional Laboratory, and tested for drug susceptibility using a line probe assay. Data were analyzed using Statistical Package for Social Sciences version 20. Bivariate and multivariable logistic regression analyses with 95% confidence intervals were carried out to identify factors associated with multidrug-resistant tuberculosis.

Results: The overall magnitude of multidrug-resistant tuberculosis was 3.8% (15/395) (95% CI: 2.0–5.8%). Being male (AOR = 4.9; 95% CI: 1.16, 20.5), patients with a previous history of tuberculosis (AOR = 4.9; 95% CI: 1.5, 29.6), treatment failure (AOR = 8.5; 95% CI: 1.61, 45.3), treatment default (AOR = 10.38; 95% CI: 1.86, 58.0), human immunodeficiency virus co-infection (AOR = 9.83.95% CI: 3, 21, 30.1) and a previous history of contact with multidrug-resistant tuberculosis patients (AOR = 14.4; 95% CI: 3.1, 67.6) had higher odds of multidrug-resistant tuberculosis.

Conclusion: The overall magnitude of multidrug-resistant tuberculosis was high. Strengthening the tuberculosis control program by giving special attention to HIV co-infected patients, treatment failure and default, previously infected patients as well as to those individuals who have a history of contact with multidrug-resistant tuberculosis infected patients.

Keywords: magnitude, multidrug resistance, pulmonary tuberculosis, eastern Ethiopia

Background

Resistance to both isonicotinic acid hydrazide (INH) and rifampicin (RIF) with or without resistance to other first-line anti-tuberculosis (TB) drugs is known as multidrug-resistant tuberculosis (MDR-TB).¹ Inappropriate use of antimicrobial drugs, or use of ineffective formulations of drugs (such as the use of single drugs, poor- quality medicines or bad storage conditions), and premature treatment interruption can cause the emergence of MDR-TB. In addition, it can be aggravated by Human Immunodeficiency Virus (HIV), under-nutrition, diabetes, smoking, and alcohol consumption.² Cases of re-treatment drug resistance is frequently detected,

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4493

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because previously treated patients can harbor strains with full or partial drug resistance.³

In 2018, out of the total TB incident cases reported, 484,000 were rifampicin resistant tuberculosis (RR-TB), and of these, 78% had MDR-TB. A total of 214,000 patients died due to MDR/RR-TB.⁴ Amongst all TB deaths an estimated 214,000 are due to MDR-TB.⁵ Even if drug-susceptible TB is reported to be reducing, MDR and extended multidrug-resistant tuberculosis (XDR-TB) are on the upsurge as a result of undiagnosed, untreated, or inappropriately treated MDR-TB cases.⁶

Ethiopia is one of the 30 high MDR-TB burden nations with an estimated prevalence of 2.7% and 14% among new and re-treatment cases, respectively.⁴ The nationwide drug resistance study estimated the rate of MDR-TB to be 1.6% in newly diagnosed TB and 11.8% among previously treated TB cases.⁷

Countries having high MDR-TB are required to undergo nonstop investigation and consistent monitoring of drug resistance to determine the magnitude and distribution of drug-resistant tuberculosis.⁸ Nevertheless, countries like Ethiopia, have a limited capacity of laboratories to carryout culture and drug susceptibility test (DST) of *M. tuberculosis.*⁹

Anti-TB drugs has been in use for a long time in Ethiopia before the implementation of directly observed treatment short courses.¹⁰ However, information is scarce regarding MDR-TB. This study was intended to assess the magnitude of drug resistance and associated factors among adult smear-positive pulmonary tuberculosis patients in Harari regional state health facilities, eastern Ethiopia.

Methods and Materials Study Seating and Design

This study was conducted in Harari Regional State governmental health facilities from March to October, 2019. The region is located in the eastern part of the county at a distance of 525 km east of Addis Ababa, the capital city of Ethiopia. According to the 2007 census, the total population of the region was 231,870.¹¹

The region consists of six hospitals and eight public health centers that provide TB treatments. It has also one health research and regional referral laboratory in which various diagnostic services including MDR-TB identification and DST are performed.

A cross-sectional study was conducted among adults (≥18 years). New smear-positive pulmonary tuberculosis

patients visiting TB treatment centers' during the study period were included in the study. Smear-positive pulmonary tuberculosis patients who were unable to provide appropriate sputum samples due to different reasons were excluded.

Sample Size Determination and Sampling Technique

The sample size was determined by using a single population proportion formula with the assumption of prevalence of drug resistant tuberculosis (58.6%) from a previous study,¹² with desired precision of 5%, at 95% confidence interval, and 10% non-response rate. A total of 409 participants who fulfilled the inclusion criteria were recruited consecutively.

Data Collection Method

In the data collection, eight laboratory personnel were involved. Data were collected using a structured questionnaire developed after reviewing different literature ^{13–16} and medical records were reviewed to identify participants' HIV and diabetes status. The questionnaire was initially prepared in English then translated into local languages (Amharic and Afan Oromo).

Smear Microscopy

Two consecutive spot-spot sputum specimens of at least 2– 5 mL volume were collected in clean, leak-proof, screw cap, wide-mouth disposable containers, and examined for acid fast bacilli (AFB) in the respective health facility TB laboratory.

Smears for AFB were prepared by taking a small portion of the purulent part of the sputum with an applicator stick and smearing it on a microscope slide which was then air-dried. The standard procedure of auramine staining method was employed to confirm the presence of AFB.¹⁷ Smears were examined using light-emitting diode fluorescence microscopy (LED-FM) using 20x objective for scanning and 40x for confirmation; one length has to be scanned before reporting a smear as negative, corresponding to 300–200 high-power fields.¹⁷

All smear-positive samples for AFB were collected and transported in a cold box to the Health Research and Regional Laboratory of Harari Regional State for molecular tests by line probe assay (LPA).

Molecular Detection of Drug Resistance

The GenoType MTBDRplus Ver 2.0 assay was performed according to the manufacturer's instructions (Germany: Hain Life Sciences). The three assay-based steps are involved in the whole procedure namely: deoxyribonucleic acid (DNA) extraction, multiplex polymerase chain reaction (PCR) amplification and reverse hybridization.

DNA Extraction

Deoxyribonucleic acid (DNA) was extracted from 500 μ L of the decontaminated specimen in a 1.5 mL microcentrifuge tube and centrifuged at 10,000 g for 15 minutes. The pellet was resuspended in 100 μ L genolyse A and incubated at 95 °C for 5 minutes, and then 100 μ L of genolyse B was added. After centrifugation, the supernatant was taken as DNA extract.

Master Mix Preparation

The master mix was prepared in a cleanroom to prevent contamination of the molecular laboratory. All the reagents needed for amplification are included in the amplification mix-A (AM-A) and amplification mix-B (AM-B) for master mix preparation. Master mixes were prepared according to the instruction. In template addition room, 5 μ L of DNA extracts was added to the corresponding PCR master mix tubes. For positive control, 5 µL of DNA extract from H37Rv quality control strain was added to the mixture, and for negative control, 5 µL of sterile distilled water was used. After addition, polymerase chain reaction (PCR) was performed using pre-made amplification mixes (amplification mix-A and amplification mix-B) that contained all the necessary components. After completion of the PCR process, the amplicon was detected with a series of procedures by adding different reagents to the strip. The strips were formed color bands after the addition of the final substrate reagent. Finally, amplification and hybridization were carried out to detect the drug-resistant pattern according to the instructions.¹⁸ The master mix was prepared in master mix room by adding Mix-A (AM-A), and Mix-B (AM-B) solutions according to the manufacturer's instruction. In the template addition room, 5 µl of DNA extracts was added into a 96 PCR wells containing master mix solution. For positive control, 5 µl of DNA extract from H37Rv strain was added to the mixture, and for the negative control 5 µl of sterile nucleic acid free water was used. After addition of the substrate, amplification, hybridization

and detection of drug resistance were carried out by using line probe assay.¹⁹

Data Quality Control

Three days of training was given for data collectors on data collection procedures and data quality management. The questionnaire was pretested on 5% of TB infected patients at Haramaya District Hospital, eastern Ethiopia to check its consistency and simplicity before the actual data collection. Instruments and reagents were checked for reliability and reproducibility of the test before any test started. All new lots of reagents were tested with a known positive and negative controls.

Method of Data Analysis

The data were cleaned, coded, and double-entered using Epi-Data version 3.1 software and then exported to Statistical Package for Social Sciences version 20 software for analysis. Descriptive statistical tools were used to summarize the findings. Bivariate and multivariable logistic regression analyses were performed to assess the association between dependent and independent variables. Crude and adjusted odds ratio at 95% confidence interval were used to identify the predictors. Those variables with a p-value < 0.25 at bivariate logistic regression model. A p-value less than 0.05 in the multivariate analysis was used as a cutoff value to declare statistical significance.

Results

Socio-Demographic Characteristics

A total of 395 smear-positive pulmonary tuberculosis patients were enrolled, with a 96.6% response rate. Among them, about half of the respondents (n = 203, 51.4%) were at the age of 26–45 years with the mean age of 32.7 (standard deviation(SD) \pm 12.7) years and the majority of participants (n = 295, 74.4%) were females. Most of the respondents (n = 311, 78.7%) were from rural areas and (n = 133, 33.7%) attended secondary school (Table 1).

Behavioral Characteristics of Study Participants

Of all the participants, those who had reported a smoking habits, chewing khat, and drinking alcohol were 25.3% (100/395), 39% (154/395), and 21.3% (84/395), respectively.

Sex

Residence

Occupation

Educational

Family Size

status

>46

Male

Female

Rural

Urban

Farmer

Student

Merchant Unemployed

Daily laborer

Government

Unable to read and

Read and write

College and above

Elementary

Secondary

≥5

<5

Prisoners

Write

Housewife

Table I Socio-Demographic Characteristics of Smear-PositivePulmonary Tuberculosis Patients Who Attended HealthFacilities in Harari Regional State, Eastern Ethiopia, 2020(n = 395)

53

100

295

311

84

58

40

67

80

6

45

52

47

51

52

57

133

102

308

87

134

25.3

74.7

68.0

32.0

14.7

10.1

17.0

20.3

1.5

11.4

13.2

11.9

12.9

13.2

14.4

33.7

25.8

78

22

 Variable
 Frequency
 Percentage (%)

 Age
 <25</td>
 139
 35.2

 26-45
 203
 51.4

Clinical	Characteristics	
Clinical	Characteristics	

Among 395 participants, 113 (28.6%) had a previous history of anti-TB treatment but 282 (71.4%) were new cases.

Variable		Frequency	Percentage(%)	
Previous history of TB	Yes	113	28.6	
	No	282	71.4	
History of contact with MDR	Yes	106	26.8	
TB/TB patients	No	289	73.2	
HIV status	Positive	58	14.7	
	Negative	337	85.3	
Diabetic	Yes	30	7.6	
	No	365	92.4	
Treatment category	Failure	63	15.9	
	Relapse	26	6.6	
	Default	24	6.1	
	New	282	71.4	

One hundred and six (26.8%) participants had a contact history with TB patients and those who were co-infected with HIV and diabetes were 58 (14.7%) and 30 (7.6%), respectively (Table 2).

Magnitude of MDR TB

The prevalence of any resistance to RMP and/or INH was 6.33% (25/395). There was no mono resistance to RMP. About 2.5% (10/395) of the total were resistant to INH alone. The total prevalence of MDR-TB was 3.8% (15/395) (95% CI (2.0–5.8%)). Besides, the prevalence of MDR-TB among new and re-treated patients was 1% (2/282) and 11.5% (13/113), respectively.

Factors Associated with MDR TB

In bivariate analysis; sex, residence, previous exposure to antituberculosis treatment, a contact history with MDR-TB/TB patients, HIV co-infection, treatment category, diabetes status, chewing khat, alcohol consumption, and smoking were found to be significant at p < 0.25 and were considered as a candidate for multivariable logistic regression analysis.

In multivariable analysis; sex, previous history of TB treatment, treatment category of failure and defaulter, HIV co-infection, and contact history with MDR-TB/TB patient had a significant association with MDR-TB.

Being males were 4.9 times (AOR = 4.9; 95% CI: 1.16, 20.51) more likely to be infected with MDR-TB than females. Patients who had a previous history of anti-TB treatment were 6.6 times (AOR = 6.6; 95% CI: 1.5, 29.6) more likely to have MDR-TB compared to their counterparts. The odds of having MDR-TB in patients with a failure of treatment were 8.5 times(AOR = 8.5; 95% CI: 1.61, 45.3) and default cases were 10.4 times (AOR = 10.4, 95% CI: 1.86, 58.0) higher compared to the new TB patients. Those participants with a history of contact with MDR-TB/TB patients were 14.4 times (AOR = 14.4; 95% CI: 3.09–67.6) more likely to have MDR-TB compared to those who had no contact with MDR-TB/TB patients, and HIV co-infected patients were 9.8 times (AOR = 9.8, 95% CI: 3.21, 30.1) more likely to develop MDR-TB compared to their counterparts (Table 3).

Discussion

In this study, the overall prevalence of MDR-TB was 3.8%. The prevalence of MDR-TB among new and re-treated patients was 1% and 6.5%, respectively. Males,

Characteristics	Category	M	MDR-TB		AOR 95% CI	P-value	
		Yes	No				
Age	<25 26-45 >46	4 (2.9%) 6 (2.9%) 5 (9.4%)	135 (97.1%) 197 (97.1%) 48 (90.6%)	0.356 (0.08,1.51) 0.79 (0.22,2.79) I	I.0 (0.12,8.64) I.8 (0.25,13.54) I	0.85 0.54	
Sex	Male Female	10 (10.0%) 5 (1.7%)	90 (90.0%) 290 (98.3%)	6.10 (2.012,18.52)* I	4.9 (1.16,20.51)** I	0	0.03
Residence	Rural Urban	9 (2.9%) 6 (7.1%)	302 (97.1%) 78 (92.9%)	I.59 (0.429,5.91) I	I.25 (0.174,8.98) I	0	0.83
Previous History TB Rx	Yes No	10 (5.2%) 5 (2.6%)	103 (91.2%) 277 (98.2%)	6.57 (1.46,29.51)* I	6.6 (1.5,29.6)** I	0	0.01
Treatment Category	Failure Relapse Default New	6 (9.5%) 2 (7.7%) 5 (8.8%) 2 (0.71%)	57 (90.5%) 24 (92.3%) 19 (79.2%) 280 (99.3%)	10.16 (1.99,51.71)* 1.48 (0.373,5.22) 2.702 (0.648,11.26) 1	8.5 (1.61,45.32)** 2.6 (0.35,19.67) 10.4 (1.86,58.01)** 1	0.03 0.29 0.01	
Contact history with MDR TB/TB Patients	Yes No	12 (11.3%) 3 (1.0%)	94 (88.7%) 286 (99.0%)	I5.17 (3.65,62.20)* I	I 4.4 (3.085,67.6)** I	0.001	
HIV Status	Positive Negative	9 (15.5%) 6 (1.8%)	49 (84.5%) 331 (98.2%)	9.7 (2.87,33.01)* I	9.83 (3.21,30.13)** I	0.01	
Diabetic status	Yes No	5 (16.6%) 10 (2.8%)	25 (83.4%) 355 (97.2%)	I.93 (0.42,9.0)	6.9 (0.81,57.96) I	0.77	
Smoking habit	Yes No	6 (6%) 9 (3.1%)	94 (96%) 286 (96.9%)	2.28 (0.703,5.85) I	0.9 (0.10,8.08) I	0.91	
Chew khat	Yes No	5 (3.2%) 10 (4.1%)	149 (96.8%) 231 (95.9%)	0.98(0.33,2.78) I	2.6 (0.39,17.17) I	0.34	
Drinking alcohol	Yes No	I (I.2%) I4 (4.5%)	83 (98.8%) 297 (95.5%)	3.19 (0.5,30.2)	0.26 (0033,1.97) I	0.19	

Table 3 Bivariate and Multivariable Analysis Among Smear-Positive Pulmonary Tuberculosis Patients Who Attended Health Facilitiesin Harari Regional State, Eastern Ethiopia, 2020

Notes: *COR at P-value <0.05. **AOR at P-value <0.05.

having previous history of TB, treatment category of TB (failure and defaulter), HIV co-infected, and history of contact with MDR TB/TB patients were predictors of MDR-TB.

In the current study, the magnitude of MDR-TB was 3.8% (95% CI: 2.0%, 5.8%). This finding is consistent with previous reports from Armachiho and Metema, Ethiopia (5.6%).¹⁴ However, higher rates of MDR-TB have been reported elsewhere in Ethiopia: Jigjiga (10.2%),²⁰ Amhara regional state (6.5%),²¹ and Amhara regional state (18.2%).²² The variation between the studies might be explained by the difference in sample size, variation in the study population, access to health care facilities, and discrepancy in effectiveness of TB control programs.

In the present study, the magnitude of MDR TB among new cases was 1% (95% at CI: 0.85–2%), which is consistent with other studies in eastern Ethiopia (1.1%),²³ Amhara regional state, Ethiopia (1.8%),²¹ and Zimbabwe (2%).²⁴ While, it is lower when compared to the previous study conducted in Jigjiga, Ethiopia $(4.5\%)^{20}$ and Armanchiho and Metema district (2.3%).¹⁴ This variation between the studies might be explained by the difference in sample size, study population, access to health care facilities, and effectiveness of TB control programs.

The magnitude of MDR-TB observed in this study among formerly treated cases was 11.5% (95% CI: 8.8– 16.8). This result is in line with the study conducted in Zimbabwe (6.4%).²⁴ However, this finding is lower than the study conducted in other parts of Ethiopia: Jigjiga (22.6%),²⁰ eastern Amhara region (18.5%),²¹ and Armanchiho and Metema district (13.9%).¹⁴ This variation might be explained by the difference in sample size, study population, access to health care facilities, and effective-ness of TB control programs.

In the current finding, 100% of RIF resistance was also found to be resistant to INH, which suggests RIF's resistance might be considered as a surrogate marker for checking MDR-TB.

Males have a higher rate of MDR-TB than females. As a result, being male was significantly associated with MDR-TB. A similar conclusion was reached in Europe²⁵ and Addis Ababa.¹⁵ The possible reason might be due to higher occupational exposure of males to the infected environment, smoking, and alcoholism.

In this current study, a previous history of TB infection was an important determinant to the development of MDR-TB. A similar pattern of results is obtained in the study conducted in other parts of Ethiopia; the eastern Amhara region,²¹ West Armanchiho and Metema district.¹⁴ This high risk of acquisition of MDR among previously-treated cases might be due to naturally occurred mutated resistance in the mycobacterium and inadequate or interrupted treatment with anti-tuberculosis agents.

In this study, patients with a history of treatment failure and default were associated with the occurrence of MDR-TB. Similar findings are reported from Jimma, Ethiopia,¹² and Amhara regional state.²⁶ The association between TB treatment failure and MDR-TB might be related to lack of supervision of treatment, improper drug regimens, and inadequate or irregular drug supply that may have a potentiate role in drug resistance.^{13,27}

Previous contact history with MDR-TB/TB patients was also identified as an aggravated factor for MDR-TB. This finding is comparable with the previous study reported from Jigjiga, Ethiopia.²⁰ Several other studies have also shown this finding, that contact with a known TB patient is linked with MDR-TB due to exposure to resistant TB.^{13,27,28} In addition, either the transmission of MDR strains or the selection of single-drug-resistant strains may also contribute to the occurrence of MDR-TB.²⁷

In the current study, HIV co-infected TB patients were more at risk of developing MDR-TB. Other previous studies conducted in eastern Ethiopia,²³ East Shoa, Ethiopia,¹⁶ and Nigeria²⁹ were broadly in line with this finding. In this study, khat chewing was not associated with MDR-TB. This finding is supported by the study conducted in Oromia Region, Ethiopia which reported that chewing khat was not a forecaster of MDR-TB.³⁰

This finding highlights that drinking alcohol and a smoking habit were not associated with MDR-TB. Whereas, the association of drinking alcohol with MDR-TB was identified in studies conducted elsewhere in Ethiopia; Bishoftu Hospital¹⁶ and Amhara National Regional State.²⁶ Similarly, the association of a smoking habit with MDR-TB was established in studies conducted in Dessie and Debre Birhan towns, Ethiopia³¹ and Varanasi, India.³² This difference might be due to variation in sample size, study population, access to health care facilities, and effectiveness of TB control programs.

Limitation of the Study

The limitation of the study was related to the study design nature, which is the cross-sectional nature of the study does not confirm a definitive cause and effect relationship. Drug resistance based on routine DST is important to determine the magnitude and trends of anti-TB drug resistance. Nevertheless, DST was not performed in this study.

Conclusions

The overall magnitude of MDR-TB was 3.8% and the prevalence of MDR-TB among new and re-treated patients was 1% and 11.5%, respectively. Being male, HIV co-infected, patients in treatment category failure and default, history of TB infection, and a contact history with MDR TB/TB patients were aggravating factors in the development of MDR-TB. Strengthening the TB control program by giving special focus to the HIV co-infection patients, treatment failure and default, had a previous history of TB infection, and a history of contact with MDR-TB/TB patients may help to alleviate the transmission of MDR-TB.

Ethical Approval

Ethical approval for this study was obtained from the Institutional Health Research Ethics Review Committee of the College of Health and Medical Sciences, Haramaya University. Besides, a permission letter was obtained from health institution officers. Moreover, respondents were fully informed about the purpose of the study and signed their consent. Information obtained during the study was kept confidential. This study was conducted in accordance with the Declaration of Helsinki.

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Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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Disclosure

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