

Occurrence of Dyslipidaemia in Patients with Tuberculosis

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Abstract

Background: Dyslipidaemia is an increasingly recognized comorbidity in individuals with tuberculosis (TB), potentially arising from infection-induced metabolic alterations and drug-related hepatotoxicity. In high-burden countries such as Nigeria, TB control strategies largely prioritize infectious disease management, with limited focus on coexisting metabolic disorders. This study evaluated the prevalence and pattern of dyslipidaemia among TB patients and explored its association with socio-demographic variables.

Methods: A hospital-based cross-sectional study was conducted among 171 newly diagnosed adult pulmonary TB patients attending the Directly Observed Treatment Short-course (DOTS) clinic at the University of Port Harcourt Teaching Hospital (UPTH), Rivers State, Nigeria. Data collection involved structured interviewer-administered questionnaires, and fasting venous blood samples were analysed for lipid profile parameters—total cholesterol, triglycerides, low-density lipoprotein (LDL), and high-density lipoprotein (HDL)—using enzymatic colorimetric assays. Dyslipidaemia was defined according to the National Cholesterol Education Program Adult Treatment Panel III (NCEP ATP III) guidelines. Associations between lipid abnormalities and patient characteristics were assessed using chi-square tests, with statistical significance set at $p < 0.05$.

Results: Dyslipidaemia was present in 75.4% of the study population. Abnormalities included elevated total cholesterol in 44%, raised triglycerides in 48%, low HDL in 15.8%, and elevated LDL in 75.4% of participants. Age was significantly associated with dyslipidaemia ($p = 0.001$), with the highest prevalence observed in those aged 40–49 years (88.1%) and 50–59 years (85.7%). Educational level was also significant ($p = 0.0001$), with tertiary education holders showing the highest prevalence (90%). Occupational status demonstrated a significant relationship ($p = 0.011$), with unskilled workers (80.2%) and skilled workers (77.8%) more affected than participants in the “Others” category (50%). No statistically significant association was found with sex or marital status.

Conclusion: Dyslipidaemia is common among TB patients in this setting, particularly among older adults, those with higher educational attainment, and individuals in formal employment. These findings underscore the importance of incorporating lipid profile assessment and cardiovascular risk evaluation into TB management protocols, especially within tertiary healthcare facilities. Early detection and appropriate intervention may reduce long-term cardiovascular complications in TB survivors.

Keywords: Tuberculosis, Dyslipidaemia, LDL, HDL, Cardiovascular risk, Nigeria, Metabolic disorders

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I. INTRODUCTION

Tuberculosis (TB) remains one of the most widespread and deadly infectious diseases globally, with a disproportionate impact on low- and middle-income countries. In 2022, the World Health Organization (WHO) estimated that 10.6 million individuals developed TB worldwide, resulting in approximately 1.3 million deaths among HIV-negative individuals and an additional 167,000 deaths among people living with HIV^{1,2}. The disease burden is particularly severe in regions marked by poverty, undernutrition, and weak healthcare systems.

Emerging evidence points to an increasing overlap between TB and non-communicable metabolic disorders such as dyslipidaemia^{3,4}. Dyslipidaemia—defined by abnormalities in lipid metabolism, including elevated total cholesterol, low-density lipoprotein cholesterol (LDL-C), triglycerides, or reduced high-density lipoprotein cholesterol (HDL-C)—is a well-established risk factor for cardiovascular disease^{5,6}. Its occurrence in TB patients is not fully understood but is believed to be influenced by factors such as chronic inflammation, nutritional deficits, drug-induced metabolic changes, and immune system activation^{7,8}. Classically, it presents as elevated plasma triglycerides and/or cholesterol, often accompanied by reduced HDL-C and increased LDL-C, typically associated with high dietary fat intake and increased hepatic lipid metabolism⁹.

The relationship between TB and lipid metabolism is complex. *Mycobacterium tuberculosis* has been shown to interact with host lipids, promoting the formation of lipid-rich “foam cells” that aid bacterial persistence and immune evasion^{10,11}. Systemic inflammation and metabolic stress during active TB can further disrupt lipid homeostasis^{5,8}, while anti-TB medications, particularly rifampicin and isoniazid, have been implicated in altering lipid profiles^{5,6}. Elevated cholesterol levels have been associated with increased TB risk, as the pathogen relies on host-derived lipids for survival, contributing to foam cell formation¹². In TB–diabetes comorbidity, impaired lipid metabolism may result in high very low-density lipoprotein (VLDL), low HDL-C, elevated sphingomyelin, and increased remnant total cholesterol lipoproteins, with LDL-C elevation linked to reduced LDL lipolysis. Lawson et al. reported elevated LDL-C in 29% of their cohort, high total cholesterol in 11.7%, high triglycerides in 6.6%, and low HDL-C in 50.5%¹³.

Despite these findings, there is limited data on the prevalence and pattern of dyslipidaemia in TB patients, particularly within sub-Saharan Africa. Characterizing dyslipidaemia in this population is essential for developing integrated TB care strategies and reducing long-term cardiovascular complications in TB survivors. This study therefore seeks to determine the prevalence of dyslipidaemia among TB patients at the University of Port Harcourt Teaching Hospital and to examine its socio-demographic and clinical correlations.

II. MATERIALS AND METHODS

2.1 Study Design and Setting

This hospital-based descriptive cross-sectional study was carried out at the Directly Observed Treatment, Short-course (DOTS) Clinic of the University of Port Harcourt Teaching Hospital (UPTH) in Rivers State, Nigeria. UPTH serves as a tertiary referral center, offering diagnostic and treatment services to patients from both urban and rural areas across southern Nigeria. Data collection spanned a 12-month period.

2.2 Study Population

The study population comprised adult patients aged 18 years and above who had a new diagnosis of pulmonary tuberculosis and had initiated anti-TB treatment within the study period. Participants were recruited consecutively. Eligibility criteria included a confirmed diagnosis of pulmonary TB through GeneXpert testing or sputum smear microscopy and a willingness to provide informed consent. Individuals were excluded if they had a prior diagnosis or history of treatment for dyslipidaemia, were on lipid-lowering therapy, had HIV co-infection, chronic liver disease, or were pregnant.

2.3 Sample Size Determination

The minimum sample size for this study was calculated using Kish’s formula for sample size estimation in populations greater than 10,000:

$$n = (Z^2 \times p \times q) / d^2$$

Where:

- n = required sample size for large populations
- Z = standard normal deviate at 95% confidence level (1.96)
- p = estimated prevalence of diabetes mellitus among tuberculosis patients, set at 15.1% (0.151) based on findings from a similar study by Lawson et al. (Abuja, Nigeria)
- q = 1 – p = 0.849
- d = desired precision or margin of error = 0.05

Substituting the values:

$$n = (1.96^2 \times 0.151 \times 0.849) / (0.05)^2$$

$$n = (3.8416 \times 0.128199) / 0.0025 = 0.4926 / 0.0025 = 197$$

Since the total population of newly diagnosed TB patients at the University of Port Harcourt Teaching Hospital (UPTH) in the previous year was estimated at 707 (obtained from hospital records), a finite population correction was applied:

$$Nf = n / [1 + (n/N)] = 197 / [1 + (197/707)] = 197 / 1.2786 \approx 154$$

To account for an anticipated 10% non-response rate, the adjusted final sample size was:

$$154 + (10\% \text{ of } 154) = 154 + 15.4 \approx 169$$

The study therefore adopted a final sample size of 171 participants, allowing for adequate power and potential exclusions.

2.4 Data Collection

Data collection was conducted using a structured, interviewer-administered questionnaire that obtained information on socio-demographic characteristics (age, sex, education, marital status, occupation), lifestyle behaviours (smoking status, alcohol consumption), and clinical history (family history of cardiovascular disease, duration of TB illness). Anthropometric parameters, including weight and height, were measured using calibrated equipment, and body mass index (BMI) was subsequently calculated.

2.5 Laboratory Analysis

Following an overnight fast of 8–12 hours, venous blood samples were obtained from all participants. Lipid profile parameters—total cholesterol (TC), triglycerides (TG), high-density lipoprotein cholesterol (HDL-C), and low-density lipoprotein cholesterol (LDL-C)—were measured using enzymatic colorimetric assays on an automated biochemistry analyzer. Dyslipidaemia was classified based on the National Cholesterol Education Program Adult Treatment Panel III (NCEP-ATP III) criteria as: TC $\geq$ 200 mg/dL, LDL-C $\geq$ 130 mg/dL, HDL-C $<$ 40 mg/dL for men or $<$ 50 mg/dL for women, and TG $\geq$ 150 mg/dL.

2.6 Data Management and Analysis

Data were entered into Microsoft Excel and analyzed using IBM SPSS Statistics version 25. Descriptive statistics were applied to summarize participant characteristics and lipid profile parameters. Associations between dyslipidaemia and categorical variables were evaluated using the chi-square test or Fisher's exact test, with statistical significance defined as $p < 0.05$.

2.7 Ethical Considerations

Ethical clearance was granted by the Research and Ethics Committee of the University of Port Harcourt Teaching Hospital. Written informed consent was obtained from each participant before data collection and sample acquisition. Participant confidentiality was maintained through data anonymization and secure storage of all records. Individuals identified with dyslipidaemia were referred for appropriate clinical evaluation and management.

III. RESULTS

Table 1 summarizes the baseline characteristics of the 171 TB patients included in the study. Males constituted 53.8% of the cohort, while females accounted for 46.2%. The largest age category was 30–39 years (39.8%), followed by 18–29 years (27.5%), with a mean age of 30.03 ± 8.89 years. Most participants had attained tertiary education (64.3%), while 27.5% had secondary-level education and 8.2% had only primary education. More than half were married (59.1%). In terms of occupation, half of the participants (50.3%) were engaged in unskilled work, while 36.8% held skilled or professional positions.

Table 1: Socio-demographic Characteristics of patients

Characteristics	Frequency (n = 171)	Percent (%)
Sex		
Male	92	53.8
Female	79	46.2
Age Bracket (in years)		
18 – 29	47	27.5
30 – 39	68	39.8
40 – 49	42	24.6
50 – 59	14	8.2
Mean age $\pm$ SD	30.03 $\pm$ 8.89	
Education level		
Primary	14	8.2
Secondary	47	27.5
Tertiary	110	64.3
Marital status		
Single	66	38.6
Married	101	59.1
Widow/widower	4	2.3
Occupation		
Skilled/professional	63	36.8
Unskilled	86	50.3
Others	22	12.9

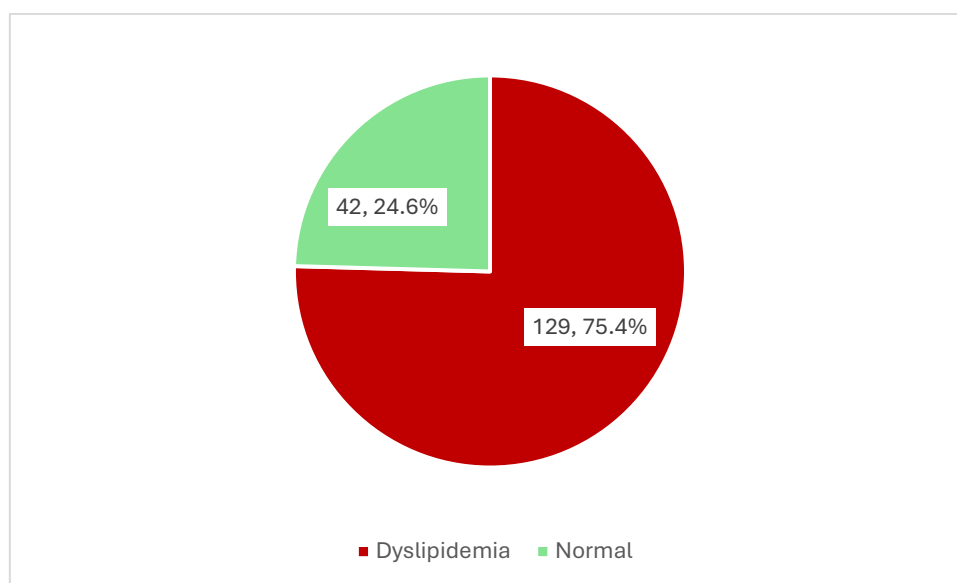


Figure 1: Prevalence of dyslipidaemia in TB Patients

Figure 1 shows that there was a 75.4% overall prevalence of dyslipidaemia among the study participants.

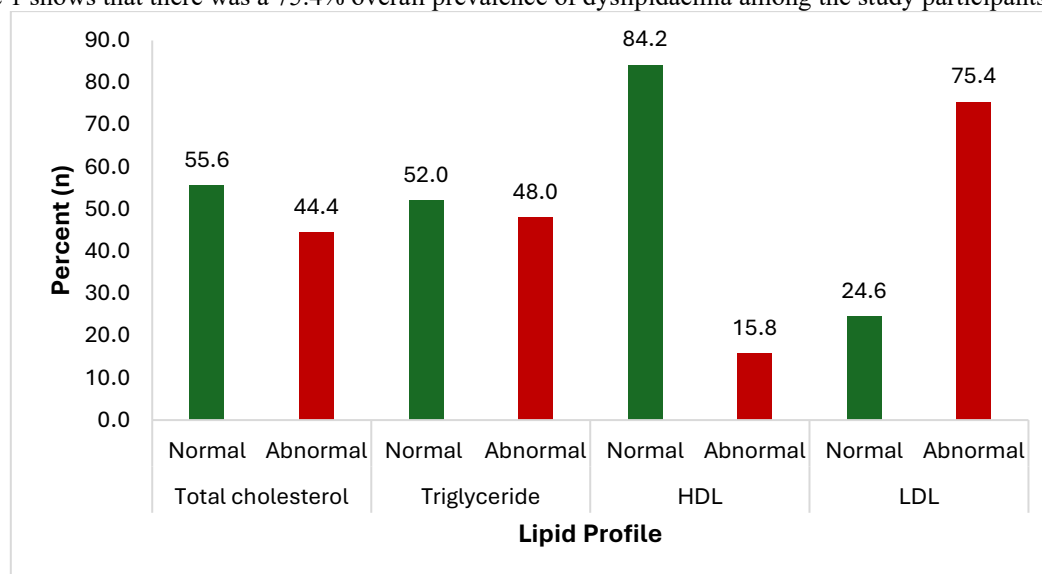


Figure 2: Distribution of Dyslipidaemia in TB Patients

Figure 2 shows the distribution of dyslipidaemia among the patients, 44% had abnormal cholesterol, 48% had abnormal triglyceride, 15.8% had abnormal HDL and 75.4% had abnormal LDL.

Table 2: Association of Demographic factors and dyslipidaemia in TB Patients

Characteristics	Dyslipidaemia n, (%)	No Dyslipidaemia n, (%)	Total n, (%)	Chi-square (p-value)
Sex				
Male	74(80.4)	18(19.6)	92(100.0)	2.68 (0.101)
Female	55(69.6)	24(30.4)	79(100.0)	
Age Bracket (in years)				
18 – 29	26(55.3)	21(44.7)	47(100.0)	15.28 (0.001)*
30 – 39	54(79.4)	14(20.6)	68(100.0)	
40 – 49	37(88.1)	5(11.9)	42(100.0)	
50 – 59	12(85.7)	2(14.3)	14(100.0)	
Education level				
Primary	7(50.0)	7(50.0)	14(100.0)	35.29 (0.0001)*
Secondary	23(48.9)	24(51.1)	47(100.0)	
Tertiary	99(90.0)	11(10.0)	110(100.0)	
Marital status				
Single	47(71.2)	19(28.8)	66(100.0)	2.80 (0.2456)
Married	80(79.2)	21(20.8)	101(100.0)	

Widow/widower	2(50.0)	2(50.0)	4(100.0)	
Occupation				
Skilled/professional	49(77.8)	14(22.2)	63(100.0)	8.93 (0.011)*
Unskilled	69(80.2)	17(19.8)	86(100.0)	
Others	11(50.0)	11(50.0)	22(100.0)	

*statistically significant ($p < 0.05$)

Table 2 shows the analysis of associations between socio-demographic factors and dyslipidaemia. The results revealed that although 80.4% of males and 69.6% of females had dyslipidaemia, the difference was not statistically significant ($p = 0.101$). A strong and statistically significant association was observed with age ($p = 0.001$), as the prevalence of dyslipidaemia was lowest among participants aged 18–29 years (55.3%) and highest among those aged 40–49 (88.1%) and 50–59 (85.7%). Educational attainment also showed a highly significant association with dyslipidaemia ($p = 0.0001$), with 90% prevalence among those with tertiary education compared to 50% and 48.9% among those with primary and secondary education, respectively. No significant association was found with marital status ($p = 0.2456$). In contrast, occupational status demonstrated a statistically significant relationship ($p = 0.011$), with the highest dyslipidaemia prevalence among unskilled workers (80.2%) and skilled/professional workers (77.8%), while participants classified under the others category had a considerably lower prevalence (50%).

IV. DISCUSSION

This study identified a markedly high prevalence of dyslipidaemia among tuberculosis (TB) patients, with 75.4% of participants affected, a figure substantially higher than the 25%–45% range reported in the general Nigerian population depending on geographic and demographic variations¹ and consistent with growing evidence that active TB perturbs lipid metabolism through inflammation-driven pathways and metabolic stress². Older age (≥ 40 years) showed a strong association with dyslipidaemia, mirroring observations by Sun et al.³ and Boillat-Blanco et al.⁴ that lipid abnormalities intensify with declining metabolic flexibility and cumulative inflammatory load, and dovetailing with population data that tie advancing age to atherogenic shifts in lipoprotein profiles²⁰. An unexpectedly high prevalence among participants with tertiary education likely reflects urban lifestyle transitions—reduced physical activity, energy-dense diets, and circadian disruption from knowledge-economy work—despite presumed health literacy, a pattern echoed among Nigerian urban professionals by Okafor et al.⁵ and by multi-country analyses that link rapid urbanization to cardio-metabolic risk accumulation^{21,22}. Occupational status also mattered: both skilled and unskilled workers showed elevated dyslipidaemia, consistent with Ekeke et al.⁶ and with literature connecting job strain, shift work, and precarious employment to adverse lipid phenotypes via neuroendocrine stress axes and sleep fragmentation^{23,24}. Mechanistically, chronic TB provokes a cytokine milieu (TNF- α , IL-6) that suppresses hepatic ApoA-I synthesis, impairs reverse cholesterol transport, and accelerates LDL oxidation⁷, while Mycobacterium tuberculosis co-opts host lipids and drives foam-cell formation to enable persistence and immune evasion⁸; first-line agents such as rifampicin and isoniazid further remodel lipid homeostasis through hepatic enzyme induction and mitochondrial stress, compounding the atherogenic shift^{8,9}. Beyond traditional lipids, functional changes matter: HDL becomes pro-inflammatory under oxidative stress, with altered paraoxonase-1 activity and compromised cholesterol efflux capacity that track worse outcomes in infectious and cardiometabolic settings^{25,26}, and TB-associated hypoalbuminaemia and micronutrient deficits can depress measured cholesterol while masking vascular risk²⁷. Additional evidence strengthens the metabolic-infection nexus: in TB–diabetes comorbidity, patients exhibit highly atherogenic signatures with elevated VLDL triglycerides, sphingomyelins, and remnant cholesterol together with reduced HDL, as shown by Vrieling et al.¹⁹; Mendelian randomization by Du et al. suggests differential, fraction-specific links between lipids and TB susceptibility, with HDL-C, LDL, and triglycerides positively associated with TB risk but total cholesterol inversely related⁸; cohort work by Chidambaram et al. ties higher baseline HDL and total cholesterol to lower mortality and dampened inflammation¹¹; and studies by Mani et al.² and Suresh et al.⁹ show that low lipid levels accompany greater TB severity yet tend to normalize on therapy, arguing for bidirectional causality and potential use of lipids as dynamic biomarkers. Host genetics may shape these trajectories—polymorphisms in APOE, CETP, and LDLR influence lipoprotein handling and infectious susceptibility in diverse populations^{28,29}—while gut dysbiosis during TB and antibiotic exposure likely perturbs bile-acid and short-chain-fatty-acid signaling that modulates hepatic lipid synthesis³⁰. Therapeutically, adjunct strategies deserve attention: statins exhibit immunomodulatory and antimycobacterial-adjunct effects in observational and preclinical work, with signals for improved TB outcomes and reduced mortality independent of lipid lowering^{13,31}, metformin has been associated with enhanced host-directed control and better clinical endpoints in TB–diabetes cohorts³², and omega-3 polyunsaturated fatty acids may temper TB-related inflammation and restore macrophage lipid handling in experimental models³³. Together, these strands underscore a syndemic interface between infection and metabolism that TB programs cannot ignore; integrating fasting lipids or non-fasting lipid panels, BMI and waist-related

metrics, and simple dietary/physical-activity screening into DOTS visits would enable early risk stratification, while targeted counseling, timely referral for lipid management, and careful pharmacovigilance for drug-induced dyslipidaemia could mitigate downstream cardiovascular risk, particularly in tertiary centers serving urbanizing catchments where cardiometabolic risk converges with infectious burden^{21,22,23}. Early identification and focused management of dyslipidaemia, coupled with prospective follow-up through intensive and continuation TB treatment phases, should clarify reversibility, disentangle causal pathways, and, critically, reduce the atherosclerotic complications that increasingly shape survivorship in high-burden settings.

V. CONCLUSION

This study not only confirms the high prevalence of dyslipidaemia among TB patients, but also identifies significant socio-demographic correlates, including age, educational status, and occupation. These findings reinforce the necessity of early lipid screening and comprehensive risk assessment in all TB patients, particularly those at heightened metabolic risk. Future studies should adopt a longitudinal design to determine whether dyslipidaemia resolves with TB treatment or persists as a chronic comorbidity. Moreover, prospective cohort data could help delineate whether metabolic disturbances influence TB outcomes such as treatment failure, relapse, or mortality.

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