

## Research Article

# Impact of GeneXpert MTB/RIF Assay for the Detection of Pulmonary and Extrapulmonary Tuberculosis from Clinical Samples

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## Abstract

**Background:** Tuberculosis (TB) is a remains major infectious disease, particularly in developing countries. Diagnosing TB remains challenging, and the rise of drug-resistant strains poses a serious risk in resource-limited settings. Therefore, improving TB diagnosis is a global priority for effective control. This study aimed to evaluate the diagnostic accuracy of the GeneXpert MTB/RIF assay for identifying both pulmonary TB and extrapulmonary TB. Additionally, it assessed the performance of the GeneXpert system in detecting rifampicin resistance among the patients involved.

**Methods:** Clinical samples were collected from patients undergoing clinical and radiological evaluations for either Pulmonary Tuberculosis or Extrapulmonary Tuberculosis. These samples were processed and tested for *Mycobacterium tuberculosis* presence using the GeneXpert assay.

**Results:** Its sensitivity is 99.8% for pulmonary tuberculosis, and specificity is 99.9% for extrapulmonary tuberculosis. Among the 15.01% of individuals suspected of having tuberculosis, the positivity rates vary by group: 7.54% in people living with HIV (PLHIV), 2.22% in paediatric patients, 20.53% in smear-negative cases, 14.45% in individuals with X-ray results suggestive of TB, and 16.97% in vulnerable groups according to Active Case Finding guidelines. Furthermore, 9.78% of contacts of TB and drug-resistant TB patients tested positive, along with 14.88% in EPTB cases. Contacts of TB and Drug-resistant TB patients have an increased risk, with 23.76% diagnosed with rifampicin-resistant tuberculosis. Patients who had contact with TB or DRTB patients have 7.87 times the odds of developing rifampicin-mono-resistant TB compared to those without such contacts.

**Conclusion:** The GeneXpert MTB/RIF assay is a rapid and highly sensitive method for diagnosing PTB and EPTB. Its simplicity and accuracy make it an impressive and valuable tool for detecting *M. tuberculosis* and rifampicin resistance.

**Keywords:** *Mycobacterium tuberculosis*; GeneXpert; Rifampicin resistant; Sensitivity; Specificity; Positive Predictive Value; Negative Predictive Value

## Abbreviations

RR-TB: Rifampicin-resistant Tuberculosis; PBS: Phosphate-Buffered Saline; DR-TB: Drug-Resistant Tuberculosis; DS-TB: Drug-Sensitive Tuberculosis; MDR TB: Multidrug-Resistant Tuberculosis; TB: Tuberculosis; MTB: *Mycobacterium Tuberculosis*; NTM: Non-Tuberculous Mycobacteria; EPTB: Extrapulmonary Tuberculosis; PTB: Pulmonary Tuberculosis; CI: Confidence Interval; HIV: Human Immunodeficiency Virus; WHO: World Health Organization; PMDT: Programmatic Management of Drug-Resistant Tuberculosis; NAAT: Nucleic Acid Amplification Test; CSF: Cerebrospinal Fluid; OR: Odds Ratio

## Introduction

Tuberculosis (TB), caused by the bacterium *Mycobacterium tuberculosis*, is one of the leading causes of mortality worldwide and ranks among the deadliest infectious diseases. Despite extensive global efforts, TB continues to be a significant public health threat, particularly in developing and underdeveloped countries. The World Health Organization (WHO) estimates that in 2023, approximately 10.8 million people were infected with TB, consisting of 55% men, 33% women and 12% children and young adolescents. That year, TB resulted in an estimated 1.25 million deaths, including 161,000 individuals living with HIV. Eight countries accounted for more than two-thirds

of the global total of TB cases: India, Indonesia, China, the Philippines, Pakistan, Nigeria, Bangladesh, and the Democratic Republic of the Congo. Notably, the top five countries together accounted for 56% of the global TB burden [1]. Diagnosing tuberculosis (TB) can be difficult due to its nonspecific symptoms and the paucibacillary nature of the disease. Accurately detecting *M.tuberculosis* is essential for diagnosing TB, which primarily affects the lungs and is transmitted through respiratory means. The GeneXpert MTB/RIF assay (Xpert) is a diagnostic tool that has significantly improved the accuracy of TB detection in clinical settings, offering enhanced sensitivity and specificity [2]. While early detection of TB can be challenging, Xpert has increased the effectiveness of the diagnostic process. However, the accuracy of Xpert can vary based on different diagnostic specimens and the sites of TB infection. Therefore, selecting the appropriate specimens is crucial when using Xpert to identify suspected TB cases. This study assesses the effectiveness of Xpert in diagnosing various types of TB using different specimens.

## Methods

### Study Settings

Early morning sputum samples (2-4 ml) were collected in pre-sterilized 50 ml Falcon tubes and processed at ten NAAT (Nucleic Acid Amplification Test) sites from January to December 2023 using GeneXpert for *M.tuberculosis* and Rifampicin-resistance detection. Extra pulmonary samples were processed at the Intermediate Reference Laboratory with GeneXpert instruments. In this study, we successfully enrolled a total of 37695 tuberculosis (TB) suspects from Puducherry, contributing 9,554 participants. Additionally, we collected data from nine neighboring districts in Tamil Nadu, which included Villupuram (2,837 participants), Kallakuruchi (3,243), Cuddalore (2,412), Tiruchirappalli (7,592), Perambalur (2,164), Thiruvarur (3,721), Nagapattinam (1,021), Thanjavur (2,077), and Tiruvannamalai (3,074). This comprehensive enrolment highlights our significant effort to address TB suspicion in the region.

### Sputum Sample Processing for GeneXpert MTB/RIF Assay

The GeneXpert sample reagent was added to each sputum specimen at a 2:1 (v/v) ratio using a sterile pipette. The sputum cup was shaken vigorously 10 to 20 times and then incubated at room temperature for 15 minutes, with at least one shake during incubation. Afterward, the sample should be liquefied without clumps. Each GeneXpert MTB/RIF cartridge was labelled with the lab accession number, either by writing on the cartridge or using the provided transfer pipette. The liquefied sample was drawn into the pipette and transferred into the cartridge. It's crucial to ensure the lab numbers match. After labelling, the pre-printed barcode on the cartridges was scanned, and the cartridge was loaded into the GeneXpert instrument. A green light indicates the test has started, and the results print automatically when complete. Wait for the system to release the door before removing the cartridge, and dispose of used cartridges in a biohazard waste container [3-4].

### Tissues Sample Processing for GeneXpert MTB/RIF Assay

Lymph nodes and tissue samples were cut into small pieces with sterile tools and placed in a sterile mortar. Approximately 2 mL of

sterile phosphate-buffered saline (PBS) was added, and the mixture was ground into a homogeneous suspension. Next, 0.7 mL of this homogenized sample was transferred to a sterile tube, followed by the addition of 1.4 mL of GeneXpert MTB/RIF Sample Reagent. The mixture was shaken vigorously for at least 10 seconds and incubated at room temperature for 10 minutes, with additional shaking. After incubation, 2 mL of the processed sample was transferred to a GeneXpert MTB/RIF cartridge, ensuring the correct laboratory number was recorded. The barcode was scanned, and the cartridge was loaded into the GeneXpert instrument. Testing initiated when the green light stopped blinking, and results were printed automatically. The cartridge was removed only after the system unlocked the door, and used cartridges were disposed of in a biohazard container [5].

### CSF Samples Processing for GeneXpert MTB/RIF Assay

If the cerebrospinal fluid (CSF) sample is less than 2 mL, add an equal volume of GeneXpert MTB/RIF sample reagent and transfer about 2 mL of the mixture into the GeneXpert cartridge. Load it into the instrument as per the manufacturer's instructions. For samples over 2 mL, transfer the entire volume to a sterile conical tube and centrifuge for 15 minutes at 4000 rpm. Discard the supernatant in a suitable disinfectant. Add 2 mL of GeneXpert reagent to the deposit and transfer 2 mL to the cartridge. Ensure the laboratory number matches the cartridge and sputum cup. Switch on the GeneXpert, scan the pre-labelled barcode, and load the cartridge. Start the test by clicking the appropriate button—when the green light stops blinking, the test has begun. Wait for the green light to turn off at the end of the test, then remove the cartridge and dispose of it in the biohazard container [3].

### Ethical Consideration

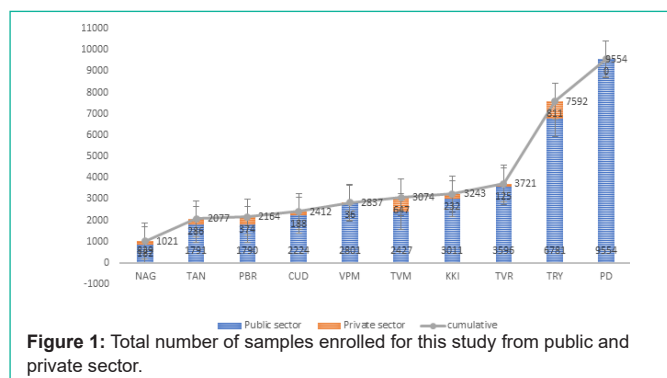
The study was approved by the Ethics and Scientific Review Committee at the General Hospital Institute, Puducherry, and conducted according to WHO guidelines and the National Tuberculosis Elimination Program.

### Statistical Analysis

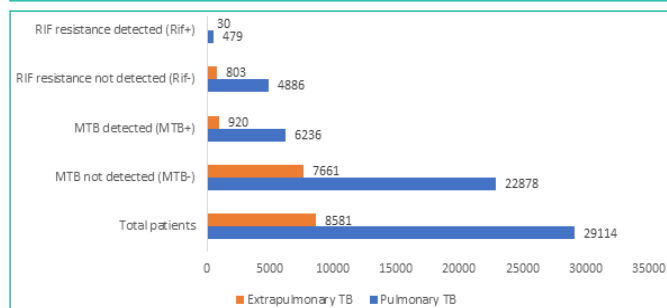
Logistic regression analysis was used to evaluate the odds ratio for multidrug-resistant/rifampicin-resistant tuberculosis, utilizing MedCalc software (version 22.026) [6].

## Results

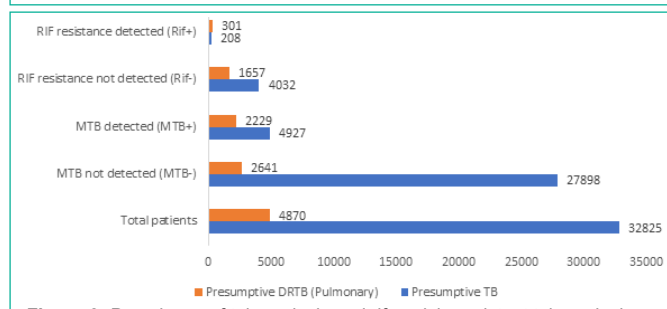
Of 37695 samples, 28141 were collected from nine adjoining districts of Tamil Nadu, while 9,554 samples came from Puducherry state. Among the 37695 samples, 74.45% (29114) were classified as pulmonary cases, and 21.94% (8,581) were categorized as extrapulmonary cases. Additionally, 87.08% of the samples were considered presumptive TB, whereas 12.92% were deemed presumptive DR-TB. Furthermore, of the 37695 samples, 92.36% (34814) were received from the public sector, and 7.64% (2,881) were obtained from the private sector, as illustrated in Figure 1. Of 29114 pulmonary samples, 21.42% (6,236) were positive for tuberculosis, and 1.65% (479) were rifampicin-resistant. Among 8,581 extrapulmonary samples, 10.72% (920) were positive for tuberculosis, and 0.35% were rifampicin-resistant, as shown in Figure 2. Out of 32825 presumptive patient samples, 15.01% (4927) tested positive for tuberculosis, with 0.63% (208) being rifampicin-resistant. Among 4870 presumptive



**Figure 1:** Total number of samples enrolled for this study from public and private sector.



**Figure 2:** Prevalence of tuberculosis and rifampicin-resistant tuberculosis among pulmonary and extrapulmonary samples.



**Figure 3:** Prevalence of tuberculosis and rifampicin-resistant tuberculosis among presumptive DRTB and Presumptive TB

DR-TB patient samples, 45.77% (2229) tested positive for tuberculosis, and 6.18% were rifampicin-resistant, as shown in Figure 3.

Of the 15.01% of presumptive tuberculosis cases, the positivity rates were as follows: 7.54% in people living with HIV (PLHIV), 2.22% in paediatric cases, 20.53% in smear-negative cases, 14.45% in cases with X-ray suggestive of TB, 16.97% in vulnerable groups (according to Active Case Finding guidelines), 9.78% among contacts of TB and drug-resistant TB patients, 14.88% in extrapulmonary TB,

**Table 1:** Prevalence of TB and rifampicin resistant tuberculosis among various categories of presumptive TB and presumptive DRTB cases.

Stratification of patients		Total	MTB not detected (MTB-)	MTB detected (MTB+)	RIF resistance not detected (Rif-)	RIF resistance detected (Rif+)	Percentage of TB positivity	Percentage of R <sup>r</sup> TB
Presumptive TB	PLHIV out of presumptive TB	2374	2195	179	155	10	7.54	5.59
	Paediatric out of presumptive TB	2257	2207	50	49	0	2.22	0
	Smear Negative, X-ray suggestive of TB	11233	8927	2306	2088	113	20.53	4.9
	Other Vulnerable group (as per ACF guidelines)	1820	1557	263	197	10	14.45	3.8
	Contacts of TB & DRTB patients	595	494	101	77	24	16.97	23.76
	EP TB	7639	6892	747	669	26	9.78	3.48
Presumptive DRTB (Pulmonary)	Upfront Molecular test offered	4026	3427	599	191	6	14.88	1
	Notified TB patients (New)- UDST	3846	2273	1573	1210	268	40.9	17.04
	Notified TB patients (Previously treated) -UDST	462	275	187	165	19	40.48	10.16
	Non-responders (DS TB & H <sup>r</sup> TB)	562	93	469	282	14	83.45	2.99
Private sector	Pulmonary TB	1939	1430	509	472	15	26.25	2.95
	EPTB	942	769	173	134	4	18.37	2.31
		37695	30539	7156	5689	509	18.98	7.11

and among those offered upfront molecular tests as presented in Table 1. Contacts of TB and DRTB patients have a heightened risk, with 23.76% diagnosed with rifampicin-resistant tuberculosis. Among the 15.01% presumptive drug-resistant tuberculosis cases, positivity rates were 40.90% for newly notified TB patients (who underwent universal drug susceptibility testing), 40.48% for previously treated TB patients (also tested), and 83.45% for non-responders (including patients with drug-sensitive TB and high-risk TB). Both newly notified and previously treated TB patients demonstrate a higher likelihood of having rifampicin-resistant tuberculosis. From 2,881 samples collected from the private sector, 26.25% tested positive for pulmonary TB, while 18.37% tested positive for EPTB. The overall positivity rate for presumptive TB was 18.98%, and the prevalence of rifampicin-resistant cases among those with presumptive drug-resistant TB was 7.11%. Table 2 presents the sensitivity, specificity, positive predictive value, and negative predictive value accuracy of the GeneXpert assay for all presumptive tuberculosis and presumptive drug-resistant tuberculosis samples. The sensitivity and specificity of GeneXpert for pulmonary tuberculosis are 99.8% and 99.9%, respectively. For extrapulmonary tuberculosis, the sensitivity and specificity of GeneXpert are 98.7% and 99.8%, respectively.

The findings of this study indicate that patients with smear-negative tuberculosis and X-ray results suggestive of TB have significantly higher odds of developing additional diseases, with an odds ratio of 1.87 (95% CI: 1.76-1.99). Additionally, contacts of both TB and drug-resistant TB patients also face increased odds for the development of diseases, reflected by an odds ratio of 1.16 (95% CI: 0.94-1.44), as illustrated in Table 3. In contrast, non-responders to treatment (including those with drug-sensitive TB and those with isoniazid monoresistant TB) exhibit an even greater odds ratio for disease development, with an odds ratio of 7.3 (95% CI: 5.80-9.19), as shown in Table 4. Table 5 presents the results of a multivariable logistic regression analysis examining various factors associated with the development of rifampicin-monoresistant tuberculosis. Notably, people living with HIV (PLHIV) who were presumptive TB patients had 1.47 times the odds of becoming rifampicin-monoresistant. Among patients with smear-negative results, those with X-ray findings suggestive of TB had 2.19 times the odds of becoming rifampicin-monoresistant compared to those who were smear-negative without X-ray suggestions. Patients classified as part of vulnerable groups also faced a higher risk of developing rifampicin monoresistance, with an odds ratio of 1.15 compared to those not in vulnerable groups. Additionally, patients who had contacts with TB or drug-resistant TB

**Table 2:** Diagnostic Performance of GeneXpert for the detection of Tuberculosis and Rifampicin-resistance among pulmonary and extrapulmonary samples

	Number	Percentage (%)	Xpert Sensitivity (%) with 95% CI	Xpert Specificity (%) with 95% CI	Xpert Positive Predictive Value (%) with 95% CI	Xpert Negative Predictive Value (%) with 95% CI	Disease Prevalence (%) with 95% CI	Accuracy (%) with 95% CI
PTB	29114	74.45	99.87(99.75-99.94)	99.92(99.88-99.95)	99.71(99.54-99.82)	99.97(99.93-99.98)	21.38(20.92-21.86)	99.91(99.87-99.94)
EPTB	8581	21.94	99.45(98.73-99.82)	99.84(99.73-99.92)	98.70(97.73-99.25)	99.93(99.84-99.97)	10.64(9.99-11.31)	99.80(99.68-99.88)
Presumptive TB	32825	87.08	99.82(99.65-99.92)	99.91(99.87-99.94)	99.51(99.28-99.67)	99.97(99.94-99.98)	14.96(14.58-15.35)	99.93(99.89-99.95)
Presumptive DRTB (Pulmonary)	4870	12.92	99.82(99.54-99.95)	99.77(99.51-99.92)	99.73(99.40-99.88)	99.85(99.60-99.94)	45.73(44.32-47.14)	99.92(99.82-99.97)
<b>Presumptive TB:</b>								
PLHIV out of presumptive TB	2374	6.07	99.44(96.91-99.99)	99.91(99.67-99.99)	98.88(95.68-99.72)	99.95(99.68-99.99)	7.50(6.47-8.63)	99.87(99.63-99.97)
Paediatric out of presumptive TB	2257	5.77	97.96(89.15-99.95)	99.91(99.67-99.99)	96.00(85.72-98.97)	99.95(99.69-99.99)	2.17(1.61-2.86)	99.87(99.61-99.97)
Smear Negative, X-ray suggestive of TB	11233	28.72	100.00(99.84-100.00)	99.99(99.94-100.00)	99.96(99.69-99.99)	100.00(99.96-100.00)	20.52(19.78-21.28)	99.99(99.95-100.00)
Other Vulnerable group (as per ACF guidelines)	1820	4.65	99.62(97.89-99.99)	99.87(99.54-99.98)	99.24(97.03-99.81)	99.94(99.55-99.99)	14.40(12.81-16.09)	99.84(99.52-99.97)
Contacts of TB & DRTB patients	595	1.52	100.00(96.31-100.00)	99.40(98.25-99.88)	97.03(91.36-99.02)	100.00(99.26-100.00)	16.47(13.58-19.70)	99.50(98.53-99.90)
EP TB	7639	19.53	99.60(98.82-99.92)	99.90(99.79-99.96)	99.06(98.06-99.55)	99.96(99.87-99.99)	9.73(9.07-10.41)	99.87(99.76-99.94)
Upfront Molecular test offered	4026	10.29	99.83(99.07-100.00)	99.97(99.84-100.00)	99.83(98.83-99.98)	99.97(99.79-100.00)	14.88(13.79-16.02)	99.95(99.82-99.99)
<b>Presumptive DRTB (Pulmonary):</b>								
Notified TB patients (New)- UDST	3846	9.83	99.94(99.65-100.00)	99.87(99.62-99.97)	99.81(99.41-99.94)	99.96(99.69-99.99)	40.85(39.29-42.42)	99.90(99.73-99.97)
Notified TB patients (Previously treated)-UDST	462	1.18	99.47(97.06-99.99)	99.64(97.99-99.99)	99.47(96.34-99.92)	99.64(97.49-99.95)	40.48(35.97-45.11)	99.57(98.45-99.95)
Non-responders (DS TB & H <sup>r</sup> TB)	562	1.44	99.57(98.47-99.95)	97.85(92.45-99.74)	99.57(98.34-99.89)	97.85(91.94-99.45)	83.45(80.12-86.43)	99.29(98.19-99.81)
<b>Private sector:</b>								
Pulmonary TB	1939	4.96	100.00(99.28-100.00)	99.93(99.61-100.00)	99.80(98.62-99.97)	100.00(99.74-100.00)	26.20(24.25-28.22)	99.95(99.71-100.00)
EPTB	942	2.41	98.82(95.81-99.86)	99.35(98.50-99.79)	97.11(93.34-98.77)	99.74(98.98-99.93)	18.05(16.64-20.65)	99.26(98.47-99.70)

**Table 3:** Multivariable logistic regression analysis of presumptive TB diagnosis n (32825).

Study variables		MTB detected	MTB not detected	Total n(32825)%	Odds ratio	95%CI	p-value
Presumptive TB	PLHIV out of presumptive TB	No	4748	25703	30451		
		Yes	179	2195	2374	0.44	0.38-0.52
	Paediatric out of presumptive TB	No	4877	25691	30568		
		Yes	50	2207	2257	0.12	0.09-0.16
	Smear Negative, X-ray suggestive of TB	No	2621	18971	21592		
		Yes	2306	8927	11233	1.87	1.76-1.99
	Other Vulnerable group (as per ACF guidelines)	No	4664	26341	31005		
		Yes	263	1557	1820	0.95	0.83-1.09
	Contacts of TB & DRTB patients	No	4826	27404	32230		
		Yes	101	494	595	1.16	0.94-1.44
	Extra pulmonary	No	4180	21006	25186		
		Yes	747	6892	7639	0.54	0.50-0.59
	Upfront Molecular test offered	No	4328	24471	28799		
		Yes	599	3427	4026	0.99	0.90-1.08

**Table 4:** Multivariable logistic regression analysis of presumptive DRTB diagnosis n (4870).

Study variables		MTB detected	MTB not detected	Total n(4870)%	Odds ratio	95%CI	p-value
Presumptive DRTB (Pulmonary)	Notified TB patients (New)- UDST	No	656	368	1024		
		Yes	1573	2273	3846	0.39	0.34-0.45
	Notified TB patients (Previously treated)	No	2042	2366	4408		
		Yes	187	275	462	0.79	0.65-0.96
	Non-responders (DS TB & H <sup>r</sup> TB)	No	1760	2548	4308		
		Yes	469	93	562	7.3	5.80-9.19

**Table 5:** Multivariable logistic regression analysis for the detection of Rifampicin-resistant among presumptive TB cases n (4434).

Study variables		RR detected n(189)%	RR not detected n(4245)%	Total n(4434)%	Odds ratio	95%CI	p-value
Presumptive TB	PLHIV out of presumptive TB	No	179	4090	4269		
		Yes	10	155	165	1.47	0.76-2.84
	Paediatric out of presumptive TB	No	189	4196	4385		
		Yes	0	49	49	0.22	0.01-3.64
	Smear Negative, X-ray suggestive of TB	No	76	1939	2015		
		Yes	113	2306	2419	1.25	0.93-1.68
	Other Vulnerable group (as per ACF guidelines)	No	179	4048	4227		
		Yes	10	197	207	1.15	0.60-2.21
	Contacts of TB & DRTB patients	No	165	4168	4333		
		Yes	24	77	101	7.87	4.85-12.77
	Extra pulmonary	No	163	3576	3739		
		Yes	26	669	695	0.85	0.56-1.30
	Upfront Molecular test offered	No	183	4054	4237		
		Yes	6	191	197	0.7	0.30-1.59

**Table 6:** Multivariable logistic regression analysis for the detection of Rifampicin-resistant among presumptive DRTB cases n (1958).

Study variables			RR detected n(301)%	RR not detected n(1657)%	Total n(1958)%	Odds ratio	95%CI	p-value
Presumptive DRTB (Pulmonary)	Notified TB patients (New)- UDST	No	33	447	480			
		Yes	268	1210	1478	3.01	2.06-4.38	0.0001
	Notified TB patients (Previously treated)	No	282	1492	1774			
		Yes	19	165	184	0.61	0.37-01.00	0.0482
	Non-responders (DS TB & H <sup>+</sup> TB)	No	287	1375	1662			
		Yes	14	282	296	0.24	0.14-0.41	0.0001

(DRTB) patients had 7.87 times higher odds of becoming rifampicin-mono-resistant than those who did not have such contacts. Table 6 further indicates that notified TB patients had a higher odds ratio of 3.01 for developing rifampicin mono-resistance.

## Discussion

Tuberculosis remains a significant public health threat, with an increasing death rate, particularly in low-resource settings. Early detection and initiation of proper treatment are crucial to reducing mortality rates. Acid-fast bacillus (AFB) smear microscopy and culture are fundamental for diagnosing tuberculosis. While culture is considered the gold standard for TB diagnosis, it is time-consuming and requires appropriate infrastructure and technical expertise. In contrast, AFB smear microscopy is a rapid and inexpensive option, but its sensitivity can be variable, ranging from 20% to 80%. Additionally, due to its limited specificity, it cannot distinguish between *M. tuberculosis* and non-tuberculous mycobacteria (NTM). Given these limitations, the fully automated Xpert MTB/RIF assay has been endorsed by the World Health Organization (WHO) as the most rapid test for diagnosing pulmonary tuberculosis. The Xpert MTB/RIF method is prioritized for diagnosing *M. tuberculosis* because it is quick, reliable, easy to use, and cost-effective. The GeneXpert system employs DNA PCR technology to detect *M. tuberculosis* and mutations related to rifampicin resistance simultaneously [7].

Almost one in seven individuals seeking evaluation at a public health facility for presumed Tuberculosis had a history of treatment for active TB. Additionally, nearly half of those evaluated had previously undergone treatment for active TB, a rate that is higher than what was reported by Mateyo et al [8]. in Zambia. Our study found that the co-infection rate of HIV and Tuberculosis was 7.54%, while the rate of rifampicin mono-resistance was 5.59%. These figures are higher than those reported by Qi et al [9]. In terms of paediatric tuberculosis prevalence, our study showed a rate of 2.22%. According to Surve et al [10] nearly five percent of new TB cases in India occur among children, which is higher than our findings. Additionally, Mane et al [11] recently reported that the prevalence of paediatric tuberculosis in India is between 6% and 7%. Our study found a prevalence of 20.53% among 11233 smear-negative X-rays suggestive of tuberculosis cases enrolled in this research. This prevalence is lower than the 23.61% reported by Khadka et al [12] in India. Additionally, Moyo et al [13] reported a prevalence of 21.34% of tuberculosis among smear-negative X-rays suggestive of TB cases in South Africa. Reta et al [14] reported a prevalence of pulmonary tuberculosis of 11.70% among key and vulnerable populations living in hotspot settings in Ethiopia, which is lower than the 14.45% reported in our study. In addition, Balakrishnan et al [15] found a TB prevalence of 14.16% among vulnerable populations, a finding that aligns closely with our results. Our study found that the prevalence of tuberculosis

and rifampicin-resistant tuberculosis among contacts of TB and DRTB patients was 16.97% and 23.76%, respectively. These rates are lower than the 21.04% reported by Paryani et al [16] in India. In our study, the prevalence of tuberculosis among extrapulmonary patients was 9.78%, while rifampicin-resistant tuberculosis was found in 3.48% of cases. These figures are lower than the 29.7% reported by Singhal et al [17] in India. Extrapulmonary tuberculosis accounts for approximately 20–30% of all active TB cases and primarily affects children and adults with compromised immune systems.

Our study indicates that individuals who are smear-negative and have X-ray results suggestive of tuberculosis are over 1.87 times more likely to develop TB compared to those who are smear-negative but have non-X-ray suggestive results. This finding aligns with research conducted by Kebede et al [18] and a survey from southwest Ethiopia, which shows that smear-negative, non-X-ray suggestive TB patients are 2.7 times more likely to have diseases. Our study indicates that contacts of TB and DRTB patients are at an increased risk of developing tuberculosis, consistent with the findings of Warriia et al [19] from Kenya. Our study suggests that people living with HIV who are presumptively diagnosed with tuberculosis, have smear-negative results, show X-ray signs suggestive of TB, belong to vulnerable groups, or are contacts of TB and drug-resistant TB patients are at an increased risk of developing rifampicin-resistant tuberculosis. This finding aligns with the results reported by Qi et al [9], Carter et al [20], and Velayutham et al [21]. This study had two significant strengths. First, it collaborated closely with the National Tuberculosis Elimination Programme, resulting in very few individuals declining participation. The extensive study population allowed for precise estimates of the prevalence and incidence of pulmonary tuberculosis. Second, involving existing health systems and community structures boosted community engagement, contributing to the project's sustainability. This study emphasizes the need for ongoing operational research on the effects of GeneXpert MTB/RIF and other factors related to the diagnosis of tuberculosis and rifampicin-resistant tuberculosis (RR-TB), which could significantly impact TB control programs overall.

## Ethical Approval and Consent to Participate

This retrospective study received approval from the Ethics and Scientific Review Committee of the General Hospital Institute in Pondicherry (Approval No. GHIEC/2023/244, dated 08-09-2023), which waived informed consent. The research followed the ethical principles of the Helsinki Declaration, and all data were kept confidential.

## Availability of Data and Material

All primary and secondary data are available with the corresponding author and in the Nikshay portal, Government of India, but can be requested from the corresponding author.

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## Competing Interests

The authors declare that they have no conflicts of interest.

## Contributors

All authors contributed to the conception and design of the study. SS, MJV, VP, UB, VR, SP, AM, RMB, SSR, GP, and GS all participated in data analysis and interpretation. MM drafted the manuscript, and all authors contributed to revisions and approved the final version.

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