### Diagnosis of TB: From conventional to modern molecular protocols

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### 1. ABSTRACT

Tuberculosis (TB) is an infectious disease caused by different strains of Mycobacterium tuberculosis complex. TB is a curable infection if diagnosed correctly and timely. Late diagnosis and improper treatment may lead to relapse or its escalation to MDR / XDR-TB. TB with HIV co-infection is difficult to diagnose by conventional set-up. Also, detection of child TB and extrapulmonary TB has its own set of problems and is not straightforward to diagnose.

The increasing complexity of TB due to the advent of new circulating strains and invasion to the regions where it was non-existent or thought to be eradicated is putting a severe strain on the health management services and making it an unmanageable pandemic. This increasing complexity has led to the spurt in the development of TB diagnostics platforms. This review focuses on the new emerging technologies that have changed the diagnostics landscape.

#### 2. INTRODUCTION

Tuberculosis (TB) is a transmissible disease caused by the bacterium Mycobacterium tuberculosis (MTB). The bacterium normally targets the lungs causing pulmonary TB (PTB). It can also infect and spread to the other parts of the body like kidney, spine and brain causing extra pulmonary TB (EPTB) there. It usually spreads through the air by means of respiratory droplets through the sneeze or cough of an infected person. For diagnosis of pulmonary TB (PTB) which is more common, samples from respiratory tract like sputum, induced sputum, broncho aleveolar lavage or lung biopsy is usually done. Diagnosis of extrapulmonary TB (EPTB) depends on the site of infection, e.g. stool sample for intestinal TB to detect Mycobacterium avium as in the case of AIDS patients: other samples may include biopsies, aspirates, pus, urine, and sterile body fluids like cerebrospinal, synovial, pleural, pericardial, and peritoneal fluids (1-3).

Tuberculosis continues to be the major cause of mortality despite all sorts of developments in medicare including diagnostics and medicines. It poses continued threat to the health programs and services in resource poor set-ups. Co-infection with HIV is the new factor that is responsible for the escalation of TB (9% of global TB incidence) (4). Majority of infection in human results in an asymptomatic latent infection and about one in ten latent infection progresses to active TB if left untreated, killing more than 50 percent of those infected. The situation becomes aggravated further due to its co-infection with HIV and incomplete treatment worsening it further to relapse cases and emergence of drug resistance in the form of multi drug resistant (MDR) and extensively drug resistant (XDR)-TB. MDR-TB develops due to bacterial resistant to first line TB drugs like rifampicin (RIF) and isoniazid (INH), while XDR-TB occurs due to bacterial resistant to fluoroquinone (FLQ), amikacin (AMK), kanamycin (KAN) and Capreomycin (CPM). Tuberculosis is difficult to diagnose due to the late onset of symptoms, so that a person acts as a reservoir of TB bacilli with potential to infect others. Similarly, co-infection with HIV makes TB difficult to detect via conventional sputum smear microscopy (SSM) / X-ray. Late diagnosis coupled with incomplete treatment escalates it to MDR and XDR-TB. Although MDR-TB is curable via second line anti-TB drugs, therapy is ineffective, marred with toxic effects and require almost 2 years of treatment which is certain to have its side effect (1-3).

20th century has seen spurt of development in TB diagnostics and medicare from conventional sputum microscopy to Nucleic acid amplification test (NAAT) based molecular and biosensing based protocols, culminating into genome sequencing. This has generated a ray of hope for the timely and accurate diagnosis and treatment, hence eradication of TB.

However, it is still the giant killer and major cause of mortality and morbidity. Main cause of concern is the emergence of TB bacilli in the regions where it is thought to be completely eradicated. Irrespective of the evolution of fancy and improvised technologies for TB diagnosis SSM and culture based (Lowenstein-Jensen (LJ) media) are still the most reliable and preferred diagnostics of choice for detecting AFB causing active TB, particularly for resource limiting set up of developing countries (2-5).

Timely detection and effective treatment are essential for managing tuberculosis. However, the rate of global case detection is considerably low. The diagnosis of pediatric TB and EPTB is particularly difficult and there are, as yet, no rapid screening tests that can help finding active cases in the target population. Tuberculosis diagnosis is primarily based on the culture method and non-culture methods. Recently, it is complemented by cutting edge molecular and genome sequencing techniques that have increased the sensitivity of TB diagnosis manifold and also provide epidemiological information about it prevalence, INTRODUCTION of new strains or cause of relapse etc (2-3).

This systematic review has incorporated a decadal growth in TB diagnostics from conventional SSM to NAAT based molecular and sequencing based protocols. The review also highlights the TB diagnostics in varied set-ups focusing on challenges of customized diagnostics. In addition, review elucidates on currently available and upcoming protocols and barriers to their successful uptake. The review further focuses on the advantage and disadvantages of each technique / protocol and intends to complement earlier reviews and reports, stimulating discussion and informing potential opportunities for market intervention to improve access to effective TB diagnostics.

### 2.1. Search strategy and selection criteria

The references for this review were identified through searches of PubMed for articles published from 2005 to 2016 by the use of terms "Tuberculosis", "diagnostics", "DST", "SSM", "NAAT". Relevant articles published during the aforementioned period were identified through searches from secondary sources (published reports, peer reviewed papers, WHO policies, UNITAID landscape report, and developer's website).

### 3. SPUTUM SMEAR MICROSCOPY (SSM)

Time tested diagnosis for active tuberculosis (TB) involves detection of *Mycobacterium tuberculosis* complex bacilli in specimens from the respiratory tract for PTB or specimens from other bodily sites for EPTB. Microscopic analysis of sputum smear detects AFB by

Table 1. Imaging techniques

1.	SSM	Developed by Robert Koch (1870)	Imaging	Sputum	Bacterial loadindicator, Rapid,economical	Fails to differentiate between tubercularand non- tuberculour, bias of technician, cannot detectEPTB	Lower, could beincreased byfluorescent staining or byincubating with CB-18 (20-30%)	Low
2.	TBDx	Signature mapping medical sciences, USA	Imaging	Sputum	High-thoughput (200slides), roboticloading of stained slides, output asdigitized images	High tier infrastructure and trained manpowerrequired, a limitationin low economical set-up	High	High
3.	CellScope	UC Berkeley / CITRIS /Banatao Instit.	Fluoresence Microscope	Blood sample	High resolution enlarged digitizedimage	Lack ofaccess in under developed countries,Insufficient studies tovalidate its utility	Sufficientstudy /data required	Sufficient study / data required
4.	CXRsalgorithm CAD4TB		Imaging	Chest X-ray	Rapid detection of active TB particularly in HIV co-infection, can be interfaced with other such platform, detects both active and passive TB	Trained expertise required	High	High
5.	Easyport able DCXR	(Delft ImagingSystems)	Imaging	Chest X- ray	Capacity to hold 50,000 DCXR images, can be operated from a laptop	Trained expertise required	High	High
6.	riView-TB	Advenio (Chandigarh, India)	Imaging,CAD4TB enabledautomated imageanalys issoftware similar toDCXR	Chest X-ray	Easy to interpret images, simultaneos diagnosis of pneumo nia and silicosis to complementPTB diagnosis	Prior training in clinical radiography required	High	High

Ziehl-Neelsen(Z-N) staining (6). Sputum test is good indicator of bacterial load hence the infectivity. It is the most common, rapid and economical diagnostic method for active PTB diagnosis but it has lower sensitivity, when bacterial load is less than 10,000 bacilli / ml of sputum (Table 1). However, sensitivity can be improved by fluorescent staining (auraminerhodamine) coupled with LED microscopes but it fails to distinguish between tuberculous and nontuberculous bacteria. Other limitations include prevalent epidemiological factors in the region of interest and skill of laboratory technician (7-8) (Table 1). Laserson et al. (2005) (9) in their study of Viatnamese immigrants have observed that incubation of sputum with detergent C18-Carboxypropylbetaine (CB-18) for 24 hours improves the sensitivity of AFB staining by 20-30% when compared to direct auramine and Z-N staining (9). Sputum analysis cannot detect EPTB. There might be a possibility that bacilli resides in cavities of lungs and may not be there in sputum. Decontamination process of sputum for mycobacterial culture may lead to the loss of bacilli from the sputum.

In addition, HIV patients and children might found it difficult to produce sputum. These constraints severely limit the utility of this very simple and economical diagnostics method<sup>10</sup>.

TBDx developed by signature mapping medical sciences, USA is an automated high-thoughput system with the capacity of 200 slides. It utilizes robotic loading of stained slides. Results are rapid in few minutes in the form of high resolution digital image (8, 11-12) (Table 1).

## 4. IMAGING BASED DIAGNOSIS: FUTURE OF AUTOMATED MICROSCOPY

Priority based treatment depending on the severity of symptoms have the potential to discriminate amongst those at risk from the people bearing symptoms of PTB but do not actually have TB. Chest X-ray (CXRs) serves as a potent algorithm in mathematical analysis of TB epidemiology. CXR integrated with user friendly, robust interface has evolved into an automated

algorithm to quantify abnormalities from digital CXR (DCXR) images. DCXR can be utilized for both active and passive TB diagnosis. It can be interfaced with other such platforms like as Xpert® MTB/RIF for rapid detection of active TB particularly in HIV co-infection setup. It is an economical tool with high sensitivity and specificity and could be a tool of need in low resource set-up (Table 1). Delft Imaging Systems (Netherlands) have come up with a software to score CXR image at the rate of one per minute. Fourth version of Computer aided detection for tuberculosis (CAD4TB) (Delft Imaging Systems) can also be integrated with DCXR platform (13-14). EasyPortable DCXR (Delft Imaging Systems) fits the requirement of limited resource set up. It includes an X-ray detector, CAD4TB software and Picture Archive and Communication System (PACS) with a capacity to hold 50,000 DCXR images and can operate from a laptop (2) (Table 1).

Advenio (Chandigarh, India) has launched riView-TB, a CAD4TB enabled automated image analysis software similar to DCXR. This platform provides easy to interpret images without prior training in clinical radiography. It is slated to be improvised for simultaneous diagnosis of pneumonia and silicosis to complement PTB diagnosis (2) (Table 1).

CellScope, a mobile cell microscope (CITRIS, University of California, Berkeley, San Francisco) is another innovation in the series of next generation microscopy. It is a fluorescence microscope that gives output in the form of high resolution enlarged digitized fluorescent images of blood samples to detect rod shaped bacterium (8, 15) (Table 1).

### 5. SAMPLE COLLECTION

A good quality sample in sufficient amount is a prime requirement for accurate diagnosis. This becomes a limitation with children and HIV patients who cannot cough up enough sputum. Lung flute from Medical Acoustics (USA) allows the patient to collect their own sample when they are not able to expectorate enough volume. It consists of a plastic device held via lips to exhale air into it. This generates vibrations in the lungs that soften and liquefy the sputum enabling patient to cough out more sputum (16).

Deton Corp. (USA) has come up with one liter bag to collect microdroplets presumed to contain MTB bacilli. These microdroplets are used for diagnosis (3). DNA Genotek Inc. (Ottawa, Canada) has developed OMNIgene® SPUTUM to liquefy and decontaminate the sputum. This facilitates its transport without cold chain for further diagnosis via microscopy, culture, Xpert® MTB/RIF and other molecular diagnostic protocols. prepIT® MAX technology from DNA Genotek Inc. (Ottawa, Canada) enables release of nucleic acid from MTBC cells via chemical lysis (3).

PrimeStore Molecular Transport Medium® (PS-MTM®) from Longhorn Vaccines & Diagnostics (USA) provides a NAAT enabled solution to inactivate, lyse and stabilize MTB DNA without cold chain. The genetic material produced can be utilized for range of other TB tests. PS-MTM® has further diversified into range of products like PrimeSwab™ and PS-MTM® for sample preparation, PrimeSwab™ and PS-MTM® for DNA extraction (PrimeXtract™) and PrimeMix® for real time PCR assay. Molzym GmbH & Co. KG (now Molzym) has launched the MTB-DNA Blood kit (25 to 50 reactions) that removes excess human genomic DNA and concentrates MTB DNA (3).

# 6. LATENT TUBERCULOSIS AND SKIN TESTING

Tuberculosis may remain latent in an infected patient before reactivation. This is one amongst the most challenging aspect in TB diagnostics. The advent of ELISPOT assay has provided promise to test for latency. This test involves the quantification of interferon production from whole blood or peripheral blood mononuclear cells (PBMC) stimulated with either purified protein derivative (PPD) or specific antigens (17-19). Currently available QuantiFERON®-TB Gold Test (Cellestis Ltd, WA, USA) for latent and active TB utilizes ESAT-6 or CFP-10 antigens (absent from BCG vaccine) (Table 4). Although the assay got the FDA approval it is yet to be commercialized. MPB skin patch test is another test with very high specificity and sensitivity (20-21). This test is free from reader bias, do not give false positive results due to booster phenomena, free from the effect of prior BCG vaccination and require only one patient visit to give sample and get result (22) (Table 4).

### 6.1. Culture based drug resistance testing (DST)

Increasing global burden of TB has diversified the diagnostic protocols for TB detection. Various phenotypic methods that include nitrate reductase assay (NRA / Griess method), thin-layer agar (TLA), colour test (Color Test), microscopic observation drug susceptibility assay (MODS), colorimetric redox indicator (CRI) method and phage-based assays can be directly set on the clinical samples and do not have to undergo sample preparation like decontamination process (2, 23-24). These methods can detect MDR-TB including resistance to the second line drugs like INH and RIF. Among different phenotypic methods available, MODS is widely validated and has perfect synergy with conventional drug susceptibility test for INH, RIF and MDR-TB (100%, 97% and 99%). MODS is rapid, economical and works well with all types of primary clinical samples and isolates. MODS depend on light microscopy. However, it requires a requisite expertise that involves extensive manpower training (25-27). The effectiveness of MODS for HIV

**Table 4.** Culture based phenotypic assays

S.No.	Assay	Manufacturer	Method	Material	Advantage	Disadvantage	Sensitivity % (95% CI)	Specif city % (95% CI)
1.	MODS	MODS (Hardy diagnostics)	DST	Sputum	Directly set on the clinical samples and do not have to undergo sample preparation like decontamination process, detect MDR-TB including resistance to the second line drugs like INH and RMP; Handling risk is entirely avoided, Safer as performed in closed system	Daily observation of culture, risk of laboratory transmission, requirement of inverted microscope, require extensive manpower training	High	High
2.	NRA	To be marketed	DST on LJ medium With KNO <sub>3</sub> (ability of MTB to reduce nitrate to nitrite)	Sputum	Simple to perform with good results, inexpensive reagents, advantage in resource limited settings	Culture time is 3-4 times of MODS, Require mature MTB culture to open to add specific reagent	High	High
3.	TLA	To be marketed	DST on solid medium	Sputum	Rapid, Accurate, Standard microscope Required which is Easily available, cost effective	Contamination rate is higher, data on DST is lacking	Low	Low
4.	CRI	To be marketed	Indirect, done on MTB isolates grown from conventional culture	Sputum	Less equipment and infrastructu, Consumables that are readily available	Biosafety level 3 required, additional staff skill	High	High
5.	TKME DIU M® SLC-L	Salubris Inc. USA	Colorimetry	Sputum	Rapid, does not require centrifugation step, no media preparation required, lower contamination	Lack of automation, Disadvantageous in terms of turn- around time, Future studies in differential settings are required	Better sensitivity over LJ for MTB detection	High
6.	Accuprobe	Gen-Probe Inc	Chemilumi nescent acridinium ester-labelled (AE)-DNA Probes (AE- MAC) probe	Sputum	Simple and rapid, Readily available In market, Validated	Labor intensive	High	High

co-infection is not yet established. Further, this test is only approved for INH and RIF leaving other first and second line TB drugs. TLA has also shown the similar promise for the detection of MDR-/XDR-TB colour test, MTB complex and resistance to INH, RIF and ciprofloxacin (CFP) in cultures (27), details given in Table 4.

TK MEDIUM® SLC-L (Salubris Inc. USA) has developed a liquid media that provides an advantage over conventional decontamination method using sodium hydroxide and N-acetyl cysteine. It does

not require centrifugation step thus completing the whole process in around 25 minutes. It shows better sensitivity over LJ for MTB detection (2) (Table 4). Gonzalo et al (2014) (27) have used Biphasic media assay (BMA) having two media L-J and Kirchner medium in tubes with 66mg/ml of pyrazinamide (PZA) to grow PZA susceptible and resistant strains of MTB (27). Reproducible results have shown the majority of starins to be susceptible to PZA while few were resistant. BMA might be a useful low cost, culture based alternative for DST for detecting PZA resistance particularly where MDR-TB is prevalent.

### 6.2. Phage amplification technique (PAT)

Phage Amplification Technique (PAT) is a bacteriophage based test that can detect MTB in sputum. Non-pathogenic mycobacteria (sensor cells) are used as a negative control. The phage replicate, infect and lyse the sensor cells leaving zones of clearing (holes) in the agar conforming that sputum contains MTB. This diagnostic protocol is rapid (2 days), sensitive, and requires less infrastructure. However, it can only be applied to sputum samples and requires expert manpower (28).

### 7. GENOTYPIC METHODS

Genotypic methods target the nucleic acid of the mycobacterium. Ribosomal rRNA, which has universal occurrence, abundance, and conserved specific sequences can be a very useful genetic marker. Accuprobe (Gen-Probe Inc) detection method targets rRNA. This method can detect M tuberculosis complex, M avium, M intracellulare, M avium complex, M gordonae, and M kansasii. Accuprobe is simple and rapid with accuracy around 90%. However, it covers a limited range of species (Table 4). Peptide nucleic acids (PNA) based assay provides another approach to differentiate M. tuberculosis complex from NTM cultures29. PCR based restriction enzyme analysis of the hsp65 also provides an economical alternative for mycobacterium detection from culture. However, technique is complicated and labor intensive (22,30). Similarly, sequencing 16sRNA can also provide rapid detection of mycobacterium but it is labor intensive (22, 31).

### 8. IMMUNOLOGICAL PROTOCOLS

Detection of antibodies like lipoarabinomannans (LAM, cell wall component of Mycobacteria), A60, 38Kd and 16 Kd against MTB is a simple and economical method for EPTB detection. However, inconsistency in results as covered by meta-analyses and systematic reviews hampers its commercialization (3,32-33).

Determine™ TB LAM Ag rapid assay (Alere Inc.) is an immune-chromatographic strip that detects LAM antigen in urine. Surprisingly, a 'negative result' does not rule out the possibility of TB. Test is also vulnerable to 'false positives' due to contamination with dust or faeces. For test to be accurate, the CD4 counts/ mm3 should not be less than 100 and best results are obtained with CD4 counts around 200 counts/mm (3, 32-35). It detects NTMs in addition to MTBC along with the site of mycobacterial infection (Table 5).

### 8.1. TB biomarker assays

The T-cell activation marker-tuberculosis (TAM-TB) assay is a flow cytometry based protocol

that detects fluorescently stained TB antigen-specific CD4 T-cells counterstained with an antibody to CD127. It measures the level of CD127 in the CD4 cells challenged with TB antigens (36) (Table 5).

### 8.2. Interferon-Gamma Release Assays (IGRA)

This test utilizes the ability of MTB antigens like Early Secretory Antigen Target 6 (ESAT-6) and Culture Filtrate Protein 10 (CFP-10) to stimulate the host for the production of interferon –gamma (IFN-γ). These antigens are absent in NTM or in BCG vaccine hence can differentiate between latent TB infection, BCG immunization and NTM infection. This procedure is simple and rapid. The commercially available assays for IFN-γ are; the Quantiferon-TB Gold (Cellestis) and T Spot-TB test (Oxford Immunotech). Intergam Rapid Immuno Suspension Array, IRISA-TB (Antrum Biotech Ltd, South Africa) detects IFN-γ in pleural, pericardial, ascitic, and cerebrospinal fluid. Testing IFN-γ has higher sensitivity for pleural TB detection as compared to the Xpert MTB / RIF (8,37) (Table 5).

### 9. VOLATILE ORGANIC CHEMICALS (VOCs)

Analysis of metabolic pathways for TB detection is another breakthrough in technology that is rapid, safe and useful for TB diagnosis in children and HIV patients. Maiga *et al* (2012) (6) have developed Rabbit urease breath test to analyze the sensitivity and specificity of urease based detection of *M. tuberculosis* (6). This test involves administration of urea labeled with isotope as substrate. Urea metabolized to 13 C-CO2 can be detected in exhaled breaths using IR spectrometers. Signal shows good relation with bacterial load that served well for diagnostic purposes and treatment monitoring (6). The test is under clinical trial to evaluate its efficacy.

BACTEC TB 460 radiometric system from Becton Dickinson, Sparks, MD, USA facilitates rapid detection of *M. tuberculosis* thus expediting evidence based therapy (6). Liquid crystal display (LCD) based TB Breathalyser from Rapid Biosensor Systems Ltd takes less than 4 minutes to detect active TB bacilli based on the MTB specific antigen in sputum and gives output as TB positive or negative (2).

# 10. NUCLEIC ACID AMPLIFICATION TESTS (NAAT)

PCR is commonly used for nucleic acid amplification tests (NAAT). NAAT can detect single copy of nucleic acid in a clinical sample. However, presence of PCR inhibitors in samples can significantly compromise the sensitivity. Also, processing of samples may lead to the loss of nucleic acids (38-42). This technique is further complemented by fluorescent probe based detection or melts curve analysis.

 Table 5. Immunological biomarkers

S.N.	Assay	Manufacturer	Method	Material	Advantage	Disadvantage	Sensitivity % (95% CI)	Specif city % (95% CI)
1.	Determine™ TB LAM Ag Rapid assay	Alere Inc.	Immunochromatographic strip	Urine	Simple and economical method, detects TB in HIV positive patients, EPTB, NTMs, MTBC along with the site of mycobacterial infection	Inconsistency in results, negative result' does not rule out the possibility of TB, Test is also vulnerable to 'false positives' due to contamination with dust or faeces, test to be accurate, the CD4 counts/mm³ should not be less than 100	Increasd when combined with sputum microscopy	High
2.	TB Rapid Screen	Tashima Inc Global BioDiagnostics, USA, with FIND, Switzerland	One-step Chromatographic immunoassay, which specifically detect the antibodies to Mycobacterum tuberculosis in human serum or plasma	Blood	Rapid, simple and easy to use, Suitable to use in peripheral laboratories, use low-cost fluorescence reader	Limited to the qualitative detection Of antibodies to M. tuberculosis in human serum and plasma, Qualitative test, Negative result does not rule out TB, Need confirmation from other tests	High	High
3.	TAM-TB assay	Ludwig- Maximilians University of Munich (LMU), Germany	Flow cytometry, fluorescently stained TB antigen-specific CD4T-cells	Blood	Quick and efficient diagnosis, Paediatric TB	Need optimization for TB detection in HIV kids and reduced cost	High	High
4.	Intergam Rapid Immuno Suspension array, IRISA-TB	Antrum Biotech Ltd, South Africa	Measures IFN-® in pleural, pericardial, ascitic, cerebrospinal fluid	IFN- y In pleural, pericardial, ascitic, and cerebrospinal fluid	Rapid assay, simple and flexible format for simple or multiple assays, Test can be manual or automated	Antigen- specific T- cell IGRAs were limited by suboptimal accuracy and the inability to isolate sufficient mononuclear cells to perform the assay	High	High
5.	T-SPOT®TB	Oxford Immunotec, UK	ELISPOT / IGRA assay	Blood	Rapid, not influenced by previous BCG vaccination	Lower sensitivity for patients with prior BCG, diagnostic value of this test in detecting active TB approached 90% in sensitivity, specificity, limitation in detecting pediatric TB	High relative to TST	High relative to TST
6.	QuantiFERON®-TB Gold Test	Cellestis Ltd, WA, USA	ELISPOT assay	ESAT-6 or CFP-10	Results are not subject to reader bias, no booster phenomena (i.e., increased sensitivity on subsequent test, leading to false-positive results), prior BCG vaccination and require only one patient visit	Not tested for infants, Kids, Pregnant women	High	High

i.e. real time PCR (RT-PCR). PCR based protocol can differentiate between wild type and mutated sequence. Other methodologies include ligase chain reaction (LCR), strain displacement amplification (SDA), loop-mediated isothermal amplification (LAMP) and transcription mediated amplification (TMA) (43-45). These molecular techniques simultaneously detects resistance genes as well, e.g. DNA probe and DNA sequencing of MTB gene such as catalase (katG) or RNA polymerase (rpoB).

Mutations in katG and rpoB lead to the resistance to INH and RIF respectively. NAAT based methods are more suited for academic / research purposes than for commercial clinical purpose.

### 10.1. Commercial NAATs based protocols

COBAS® TaqMan® MTB Test (Roche Diagnostics) is a RT-PCR based high throughput detection protocol (100 samples/day) that targets the

gene responsible for 16S RNA. This protocol requires liquefied, decontaminated and concentrated human respiratory samples like sputum and bronchial alveolar lavages. This test is highly specific to the MTBC (2) (Table 6).

The RealTime MTB from Abbott Molecular is also an RT-PCR based assay. It can be interfaced with fully automated DNA extraction platform. It detects MTBC from sputum or bronchoalveolar lavage (46) (Table 6).

GenoXtract®96 with the FluoroCycler® 96 from Hain Lifescience is a 96 well format extraction and amplification platform provided with the the GenoLyse® and FluoroLyse® for nucleic acid extraction for LPAs or MTB assay (2). The Fluorotype® MTB (Hain Lifescience) assay provides PCR based amplification of MTBC DNA. The amplicons are detected via molecular beacons and melt curve analysis (47). Fluorotype® MDRTB is an improvised version of Fluorotype® MTB that detects RIF and INH resistance in addition to MTBC detection (2) (Table 6).

Hain Lifescience has also come up with a multiplexed assay that detects MTBC and also RIF and INH resistance like the previous case. This protocol utilizes a novel PCR technique known as linear-after-the-exponential (LATE) PCR coupled with 'lights on' or 'light off' probe technology for precise identification of mutations. This protocol creates specific fingerprints for each allelic variant of drug resistant mutations (48) (Table 6).

Tosoh Bioscience (Japan) has developed TRCRapid® MTB assay that targets MTBC 16S RNA. The protocol includes amplification and transcription-reverse transcription concerted (TRC) reaction (49). NanoBioSys Inc. (Republic of Korea) has reported a microfluidic chip (LabChip G2-3) based sample processing and amplification protocol (2, 50). Detection is via RT-PCR and takes approximately 30 minutes (Table 6).

Seegene (Republic of Korea) provides a highly multiplexed RT-PCR based NAAT assay. This provides a range of Anyplex™ series of assays that includes the Anyplex™ MTB/NTM for MTBC and NTM; the Anyplex™ plus MTB/NTM/MDR-TB for MTBC, NTM and genotyping RIF and INH resistance; and the Anyplex™ II MTB/MDR/XDR-TB that detects MTBC, genotypes MDR via typing for INH and RIF resistance alleles and also 13 alleles linked with XDR-TB (7 alleles for FLQ and 6 for injectable drugs). This platform detects MTBC and also XDR if the sample is MTBC and MDR positive (50-51) (Table 6).

Xiamen Zeesan Biotech Co. Ltd (China) has developed MeltPro® Drug-Resistant TB Testing Kits as

RT-PCR enabled detection of resistant alleles to RIF and INH. It is undergoing further refinement to detect alleles resistant to ethambutol (EMB), streptomycin (STR) and fluoroguinolones (FLQs) (52-53) (Table 6).

### 10.2. Xpert MTB/RIF assay for the diagnosis of PTB

Xpert® MTB/RIF (Cepheid, Sunnyvale, CA, USA) is modular / single cartridge based NAAT platform to diagnose MTB. It involves semi-nested RT-PCR based amplification of *rpoB* and genotypes RIF resistant alleles. Melt curve analysis of amplicons generate fluoremetric fingerprints for five molecular probes in real time, which discriminate allelic variant in each probe region. This platform is accurate and sensitive and suits to the requirement of low income, limited resource set-up. This platform can process 16 to 1000 clinical samples (INFINITI® Xpert®). WHO has approved Xpert® MTB/Rif for the diagnosis of PTB, EPTB, HIV associated TB and TB meningitis in adults and children (2) (Table 6).

Xpert MTB/RIF assay provides a rapid and automated molecular test for pulmonary TB with the sensitivity of 89%, and specificity of 99% (54). However, this test is yet to be approved for the detection of non-respiratory clinical samples (55). Truenat MTB (Molbio Diagnostics, India) is a chip enabled RT-PCR operated on a portable device. Results are out in an hour. Nucleic acid is extracted from sputum utilizing a nano-particle based procedure. It shows the specificity similar to the Xpert MTB/RIF (56) (Table 6). EasyNAT Diagnostic Kit detects MTB complex. It is an isothermal amplification kit and results are qualitative in terms of visual bands. For thinned and concentrated sputum the sensitivity is comparable to culture on LJ medium (57-58) (Table 6). The VereMTB assay (Veredus Laboratories, Singapore) assay is more for academic / research purpose. It is a chip enabled platform and combines PCR with microarray technology to detect M. tuberculosis, Mycobacterium avium, Mycobacterium intracellulare, Mycobacterium simiae / kansasii / scrofulaceum, Mycobacterium abscessus / chelonae, Mycobacterium xenopi, and M. fortuitum in addition to RIF and INH (8,59) (Table 6). GeneXpert® Omni Cepheid Inc. is a mobile operated device that uses the cartridges already in use with the Xpert® platforms (e.g. MTB/RIF). This platform offers miniaturized and durable assay. It can store upto 20,000 results (2) (Table 6).

# 10.3. Xpert MTB/RIF assay for the diagnosis of extrapulmonary tuberculosis (EPTB)

Globally, PTB is more common as compared to the EPTB, which is nearly 25% of all TB cases. However, EPTB has higher prevalence in children and in HIV-infected individuals (62-63). EPTB diagnosis is marred with the lack of accuracy, pace and requisite expertise,

Table 6. NAAT based molecular diagnostics

S.No.	Assay	Manufacturer	Method	Material	Advantage	Disadvantage	Sensitivity,% (95% CI)	Specif city % (95% CI)
1.	*B-SMA RT	(LabCorp),USA	PCR+ DST,MTB detection, Resistance to PZM	Sputm	Underdeve lopment	Under deve lopment	Underdevelopment	Underdevelopment
2.	iCubateSystem	iCubate, USA	Multiplex PCR	Sputum	Highly integratedSingle platform for multiple assays forresistance, TB, NTB bacilli	For research use only	NA	NA
<sub>හ</sub>	COBAS®TaqMan®MTBTest	RocheDiagnostics	RT-PCR	Sputum and bronchial alveolarLavages 16S RNA	Ease to use, Rapid, Reduced contaminat on, Enhanced quality, Automated, High throughput	Only sputumor BAL liquefied, decontaminated am concentrated using NALC-NAOH methods, false negative results dueto PCR inhibitors	High	Highly specific to the MTBC
4.	RealTime MTB	AbbottMolecular	RT-PCR	Sputum and bronchial alveolariavages	Interfaced with fullyAutomated DNA extraction platform, detection of 8-members of MTBC, No cross reactivity with NTB	Not suited for resource limited peripheral laboratories	High	High
.5	Geno Xract® 96	Hain Lifescience	LATE –PCR amplification, molecularbeaconsand meltcurve analysis	Pulmo nary and extrapulmonary samples, stool, blood, unne, serum, swabs or cellpellets	Efficient extraction, multifunctional, flexible processing, user friendly, RIF and INH resistance inaddition to MTBCdetection	Not suited for resource limitedPeripheral laboratories	High	High
9	TRC Rapid®MTBassay	Tosoh Bioscienc, Tokyo, Japan	Transcription-reversetranscription concerted (TRC) reaction	Respiratoryand non-respiratory	Rapid technique for the detection of MTBC in respiratory. Smear positive and Smear negative non-respiration specifies a peripherallaboratories	Gross-reactions are possible when a high number of M.avium and M. marinum isolates are present	High forrespiratory samples than non-respiratory	High forrespiratory samples than non-respiratory
7.	Lab ChipG2-3	NanoBioSys Inc. (Republic of Korea)	RT-PCR	Sputum	Ultra fast cost-Effective quantitativeanalysis, takes 30minutes only, Easy-to-use, User-friendly software	Trained expertise, infrastructural set- uprequired	High	High
œί	Anyplex™series	Seegene (Republic of Korea)	MultiplexRT-PCR with DPO™ and TOCE™technologies	Respiratory,non-respiratory samples	Automatedirquid handingworkstation from nucleic acid extraction to PCR seuto, Quick andacaurate RT-PCRoptimized to multiplepathogen detection, Specialized in multiple tests, Analyze upto 94 samples	Limitations regarding detection of INHresistance	High	High
о́	MeltPro® Drug-ResistantTB TestingKits	Xiamen ZeesanBiotech Co. Ltd (China)	RT-PCR	Sputum	High accuracy, short testing turn- aroundtime, low unit price, excellent tool forMDR /XDR-TB	Significantheterogeneity for thedetection of KANresistance	High	High
10.	MFqRT-PCR	Qiagen, Germeny	Multi-Fluorescence quantitatieRT- PCR	Sputum, cultureisolates	Efficient, accurate, reliable, easy- operated method for detection of RIF and INH resistance, distinction of MTC and NTM of clinicalisolates	False negative andfalse positive results, Further studies arerequired to optimize this method and evaluate its performance	High	High
Ė.	Xpert®MTB/RIF	Cepheid, Sunnyval, CA, USA	RT-PCRMelt curveanalysis	Respiratoryclinicalsamples	Suits to the requirement of lowincome, limitedresource setup, diagnosis of PTB, EPTB, HIV associated TB and TB meningitisin adults and children	Test yet to be approved for the detection of non-respiratory clinicalsamples	High	High
15.	TruenatMTB	Molbio Diagnostis, India	RI-PCR	sputum	Portable device,rapid, quick and accurate	Expensive, Sample care needed, Large studies are required to obtain better estimates of the performance	High	High

High	High	Low	High	High	High	High	NA	N N	Variable
Low, Acceptable forsmear positive Cases smearnegative cases	High for smearpositive, low for smearnegative specimens	Low	High	Low	High	High	NA	NA	Variable
Low detection rate for the smear- negative, culture-positive sputum samples, length offine needed to get results, technicalcomplexity, high costs and biosafetyrequirements	Manual preparationof samples, pretreatedsputum resulted inpoorer performance	Failed to meet the Performance standardsrecommended by WHO for SMreplacement, Epistem is working on modifications to improve the assay	Switching on theanalyzer beforeFinishing chip loading may renderchip unusable, High stringency for chip usage	Restirated due to the length of time neededto get results, technical complexity, high costs and biosafety requirements	Tediousmulti-stepprocess	Short cartridge  The Requirement of stable electric supply. Annual calibration equired.  The perstance celling is critical, unreliable in HIV positive condition	NA	٧٧	Require prospective validation with clinical outcomes
Useful in low-resources setting, simple and easy-to-use, equipment-free, compatible with otheramplification technologies, such as PCR, LAMP, SDA, HAD, Rapid, results are qualitativen terms of visual bands	High tier laboratoryto the peripherallaboratory, robustness ofequipment, reagent stability	High iter laboratoryto the peripherallaboratory, robustnessof equipment, reagent stability	Fully automated, integrated system, robust, Affordable, GPRS and bluetoothenabled	Promising diagnosticalgorithm for suspectedchildhood TBlymphadenitis, suits to the limited resource set up of peripheral laboratories requiringcentrifuges, water bath etc to perform the assay	Highly integrate, automate, Portable, Walk a way operation, No complex procedures, minimadualification and training requirement, Rapid case detection	Portable, Provencartridge technology Durable, Low powerconsumption, Integrated battery,cloud - based connectivity	Multiple tests can berun in parallel, single analyzer to generateresults that can be used with multipleprocessors	Battery operateddevice for rapidamplification (10 minutes), real timedetection of individual XCR***assays, no primerdimer, low cost reagents and portabe instruments tion, high multiplex and simplesamplepreparation, resistance to PCR inhibitors	TB rule out" test, minimum backgroundhoise and maximumspecificity in its output
Sputum	Sputum	Sputum	Sputum	Sputum /FNAB peripheraltuberculo uslymphadenitis	Sputum /Blood	Sputum	Sputum	Sputum	Sputum
Isothermalamplification	LAMP	PCR amplification, genotyping, sequenceanalysis	Nanotechnology, MicroElectro- mechanicalsystems (ME-MS) PCR	RT- PCRcoupledwithfluorescentprobe	NEAR-isothermalamplification of TB DNA	PCR amplification	GenotypingRIF and FLQ resistantalleles	POR amplification and isothermal testing	NAAT
Ustar Biotechnology	Eiken (Japan)	Epistem (UK)	Molbio Diagnostics(India)	Ustar Biotechnologies (China)	Alere Inc. USA	Cepheid Inc.	Wave 80 Biosciences(USA)	Fluorocentric Inc. (USA)	Epicentre/ Médecinssans FrontièresLaboratory
EasyNAT Diagnosic Kit	Eiken'sLoopamp™MTB	Genedrive® MTB assay	True lab™Real Time microPCR	EasyNAT™ TB	Alere™q	GeneXpert® Omni	EOS CAPE-TBSystem	PCR(XCR™)	TOP-PCR
13.	4-	15.	16.	17.	18.	19.	20.	21	22.

which usually requires invasive procedures. Absolute culture specificity can be achieved by complementing the routine diagnostic protocols with antigen tests or NAATs. In case of TB meningitis the low culture yield can be enhanced in terms of sensitivity by replicating the CSF examination or by complementing it with NAAT (62-63). High performance liquid chromatography (HPLC) based detection of tuberculostearic acid in CSF is highly sensitive but requires expert manpower and costly laboratory set up (22,64).

### 10.4. Semi-automated NAATs for use in peripheral laboratories

The biggest challenge in molecular diagnostics of TB is to shift the set up from the higher tier laboratory to the peripheral laboratories. This may represents a diversity of issues that may range from the availability of sufficient infrastructure, robustness of equipment, reagent stability, user expertise in addition to improved performance over SSM (65-66). Four different platforms like the Loopamp™ MTBC assay from Eiken (Japan); Genedrive® MTB assay from Epistem (UK); Truelab™ RealTime micro PCR System from Molbio Diagnostics (India); and the EasyNAT™ TB assay from Ustar Biotechnologies (China) are sighted with potential for delivery (2).

Eiken's Loopamp™ MTB provides a (LAMP) based amplification assay with better sensitivity over SSM. This assay involves manual preparation of samples which are subjected to RT-PCR. Genedrive® MTB assay provides for MTBC diagnostics only with future plan of including genotyping MDR-TB through RIF resistance. This assay involves manual preparation of samples followed by RT-PCR. Truelab™ RealTime micro PCR involves semi automated DNA extraction and amplification for MTBC detection. Truelab™ QUATTRO is an improvisation over Truelab™ RealTime micro PCR in being fully automated and integrated system. EasyNAT™ TB suits to the limited resource set up of peripheral laboratories requiring, centrifuges, water bath etc to perform the assay (2) (Table 6).

Alere™ q platform is highly integrated and automated NAAT for diagnosis and assessing the viral load of HIV-1 and 2 (67). It is a cartridge based system, where cartridge is equipped with the reagents to liquefy and inactivate the sample. This is followed by extraction of MTBC-DNA and amplification. This assay involves nicking enzyme amplification reaction (NEAR), an isothermal amplification of TB-DNA (2) (Table 6).

GenePOC Diagnostics (Canada) offers an automated platform that uses RT-PCR and fluorescence detection for TB diagnostics (2). TBDx system from Keck Graduate Institute (KGI; USA) provides a low cost instrumentation where all process starting from

sample processing occurs on single cartridge platform. Nucleic acid preparation is simplified as it does not require chaotropic salts and organic solvents (2). The Northwestern Global Health Foundation (USA) in partnership with Quidel Inc. (USA) is coming up with an integrated MTBC assay. The assay involves lysing MTBC cells via heat block followed by purifying DNA using immiscible oil interface (68) (Table 6).

Point-of-Need (PON) Qiagen offers isothermal nucleic acid amplification assay for the detection of PTB and MDR-TB. Q-POC™ platform, under development from QuantuMDx (UK) offers diagnosis for PTB. The assay involves coughing the sputum in a cup that contains reagents for its liquification and decontamination. This is followed by concentration of MTBC cells via paramagnetic bead concentration step and quick nucleic acid amplification (4 minutes) via isothermal asymmetric PCR (2).

EOSCAPE-TB System from Wave 80 Biosciences (USA) offers detection of MTBC via genotyping RIF and FLQ resistant alleles. Using this platform multiple tests can be run in parallel and single analyzer generates results that can be used with multiple processors (2) (Table 6).

PCR (XCR™) assay from Fluorocentric Inc. (USA) offers a RT-PCR based assay that works via proprietary primer design algorithms. It is powered by a battery operated device for rapid amplification (10 minutes) and real time detection of individual XCR™ assays (2) (Table 6). Co-diagnostics (USA) and Thisis (a spinoff company from Boston University School of Medicine, USA) offers a RT-PCR based platform for detection of MTBC via 16S RNA gene. This platform can be integrated with variety of other RT-PCR platforms. It is claimed to detect MTBC at LOD of 4 cfu and can genotype MTBC into five geno-groups using totally optimized PCR (TOP) (Table 6). Scanogen, though not strictly a NAAT assay detects MTBC specific DNA from body fluids using single molecule scanning technique without PCR amplification. This platform claims to have minimum background noise and maximum specificity in its output (2).

### 10.5. Spoligotyping

Spoligotyping is very useful technique in providing epidemiological information on the causative strain. It utilizes amplification of polymorphic direct repeat sequences in *M. tuberculosis* genome. Higher cost of laboratory infrastructure and requirement of trained expertise are biggest limitations (5,69).

### 10.6. Line Probe Assay (LPA)

LPAs or reverse blot hybridization assays are a relatively low-cost tool and have been developed

for a variety of areas around MTB diagnostics. including MTBC-specific assays, speciation of NTMs and assays that genotype first or second-line drug resistance alleles. DNA extraction can be manual or automated and DNA is amplified by asymmetric PCR to create a population of biotin-labelled single-(ssDNA) amplicons. Oligonucleotide capture probes with sequences specific to the target amplicons are printed as an array of stripes on a nitrocellulose strip. The amplification mixture is then exposed to a strip under conditions to optimize the specific hybridization of amplicons to their sequence match probe printed on the nitrocellulose. Detection of hybridization of amplicons to the probe stripe is achieved via a streptavidin-linked enzyme that creates a colorimetric reaction visualized as defined stripes that corresponds to the presence of TB and a sensitive or resistant genotype (22). The developed LPAs are read manually with comparison to a printed score chart or can be scanned and automatically analysed by attendant software. The methodology requires open tubes containing amplified DNA, which has potential to contaminate the workspace thus compromising all future test data. The automated workstations alleviate this to some degree by providing controlled conditions that contain reaction mixtures (2).

Commercially available LPAs for the detection of TB, MDR/XDR-TB are: INNO-LiPA Rif. TB (Innogenetics, Belgium) for RIF, GenoType®MTBDR/ MTBDRplus for RIF and INH and Geno-Type® MTBDRs/ (both Hain Lifescience, Germany) for FLQ, injectables, second line drugs and ETM (22). INNO-LiPA Rif. TB is highly sensitive when isolates from culture are used. However, the accuracy goes down when it is applied directly on clinical samples. GenoType®MTBDR/MTBDRplus can be applied for both culture isolates and clinical samples. GeneXpert MTB/ RIF utilizes heminested RT-PCR to amplify specific sequence of the *rpoB* gene of *M. tuberculosis*. The amplicons are then probed with molecular beacons for mutations within RIF resistance determining region (6, 70-72) (Table 2).

MTBDRsI v1.0. and MTBDRsI v2.0. can detect resistance to FLQ, KAN, STR and XDR-TB. MTBDRsI v2.0. is equipped with 27 probes to detect resistance to second line drugs and also targets new gene regions for improved detection of FLQ and KAN (73). LPAs show high sensitivity and give better results when TB culture or SSM positive samples are used as a substrate (Table 2). These are economical when compared to culture based DST (3). However, it is practically impossible to determine accurately resistant genotypes for many drugs like pyrazinamide (PZA). Silent mutations in drug sensitive strains can be misinterpreted as drug resistance leading to avoidable treatment regimens (2).

### 10.7. Microarray-based platforms

Microarray based platforms acts as portable LPA. Microarrays have high potential for MTB diagnosis, speciation to analyze variation in drug resistant alleles. The probes utilized in microarray can be printed as defined spots and due to their small size more spots can be printed to analyze amplified DNA. The INFINITI® PLUS Analyzer platform from Autogenomics is one such platform. INFINITI® High Throughput System (HTS) and MDR-TB BioFilmChip® for the detection of MTBC and first-line drug (RIF, INH and PZA) are high-throughput array based platform (Table 3). CapitalBio has launched twin microrray based platforms, i.e. Mycobacterium Identification Array Kit for MTB diagnostics and NTM speciation and Drug Resistance Detection Array Kit for MTBC diagnosis and for genotyping RIF and INH resistance (2, 74) (Table 3).

The TruArray® MDR-TB microarray is fluorescent probe based single tube multiplexed asymmetric PCR reaction. It can detect MTB and *M. avium* along with resistant alleles to RIF and INH (2). VereMTB™ Detection Kit from Veredus Laboratory is solely for research purpose. This array based platform can detect MTBC through *IS6110*. In addition the array can detect MTBC, speciate eight NTMSs (*M. avium, M. simiae, M. intracellulare, M. Kansasii, M. abscessus, M. scrofulaceum, M. chelonae, M. xenopi*) and probe alleles associated with RIF and INH resistance (74) (Table 3).

HYDRA 1K developed by Stanford University (USA) together with Insilixa Inc. (USA) is an array based platform to detect PTB MDR / XDR-TB through the genotyping of mutations. This uses complementary metal-oxide semiconductor (CMOS) technology for digital imaging on the array. PCR amplification happens in the array area so that binding with amplicon with probes is measured in real time that saves lot of time and energy. CMOS enables this platform to have melt curve analysis of amplicon binding at each probe spot. It is developed to detect PTB, MDR / XDR-TB. However, higher cost of instrument and other peripherals makes this technique less likely to commercialize (2, 74) (Table 3).

### 11. WHOLE GENOME SEQUENCING

Whole genome sequencing (WGS) is recently applied to analyze molecular evolution of *M. tuberculosis* Beijing strain and appearance of an Iberico-American strain in Tibet. WGS provides greater resolution when compared to mycobacterial interspersed repetitive unit-variable-number tandem repeats (MIRU-VNTR, the existing protocol for molecular genotyping of circulating strains) (75-77).

**Table 2.** Line probe assay (LPA)

S.No.	Assay	Manufacturer	Method	Material	Advantage	Disadvantage	Sensitivityy % (95% CI)	Specificity % (95% CI)
1.	INNO-LiPA Rif. TB	Innogenetics, Belgium	LPA	Sputum	Economical, reliable, widely applicable assay, valuable tool for routine diagnostics, simple, rapid	Accuracy goes down when it is applied directly on the clinical sample	Highly sensitive when isolates from the culture are used	High
2.	GenoType®MTBDR/ MTBDR +	Hain Life Sciences, Germany	LPA	Sputum, broncho- alveolar lavage, bronchial aspirate, pleural effusion, lymph node sample	Can be applied for both culture isolates and clinical samples	Contaminate the work space, may have the potential to compromise all future test data, cannot be used for direct testing of smear —negative paucibacillary sputum samples	High	Modest and variable
3.	GenoType®MTBDR sI	Hain Life Sciences, Germany	LPA	Sputum	Can be applied for both culture isolates and clinical samples, sensitive detection, efficient diagnosis, flexible processing, efficient results	Contaminate the work space, may have the potential to compromise all future test data, further research is required on test accuracy in different settings, cannot be used to rule out XDR-TB	Highly specific tool for XDR detection in direct specimens for HIV co- infected TB patients	High
4.	LiPA pyrazinamide	Nipro	LPA	Sputum/ culture based	High throughput, identify the major clinical isolates of Mycobacterium species, detect mutations associated with resistance to INH, RIF, PZA and FQ, rapid, reliable detection of DR-TB NTM	Automated sequencing is required for which there is no LiPA probes, modern molecular instrumental set-up is required	High	High
5.	REBA MTB-MDR	YD- Diagnostics	LPA revere hybridization assay	Sputum	Rapid detection of RIF, INH, FQ, +SLID, simple, economical, no costly equipment is required	Underestimate the true number of resistant strains because of limitations of knowledge regarding the limited knowledge regarding the genotype markers of drug resistance	High	High

Table 3. Microarray based diagnostic platforms

S.No.	Assay	Manufacturer	Method	Material	Advantage	Disadvantage	Sensitivity, % (95% CI)	Specif city, % (95% CI)
1.	INFINITI® PLUS	Autogenomics	Microarray platform	Clinical samples	High- throughput array, high performance genetic marker detection with varying options in automation and throughput	Can only be used with human nasopharyn geal aspirates, negative results do not preclude infection	NA	NA
2.	VereMTB assay	Veredus Laboratories, Singapore	PCR-Micro array	Sputum / culture based	Rapid detection, differentiation, identific ation of MTBC, Resistance to RIF, INH non-MTBC species, diagnosing patients after treatment failure and relapse	More for academic / Research purpose	High	High
3.	VereMTB™ Detection Kit	Veredus Laboratory	Array based platform	Sputum / culture based	Detect MTBC, speciate eight NTMSs	Research purpose only	High	High
4.	HYDRA1K	Stanford University (USA) together with Insilixa Inc. (USA)	Array based platform	Sputum / culture based	Miniaturization Saves time and energy	Higher cost of instrument	NA	NA
5.	TruArray® MDR-TB microarray	Akonni Biosystems	Fluorescent probe based Single tube multipled asymmetric PCR reaction	Bacterial culture / sputum	Minitaurize, Bench top test, single integrated work- flow approach, accurate, rapid, broad coverage and field ready	Research purpose only	High	High
6.	Twin Microrray	CapitalBio	Mycobacterium identification array kit for MTB, NTM speciation, drug resistance, MTBC diagnosis, genotyping RIF, INH resistance	Clinical specimens	Both MTC and NTM rapid system, sensitive, reliable, fast, spinal tuberculosis	Infrastructure and trained personnel required reduces its clinical utility in poverty- stricken zones, Additional large- scale studies are needed to evaluate its diagnostic performanc in spinal tuberculosis	High, variable for different clinical specimes	High,variable For different clinical specimens

QIAxel automated genotyping system from Qiagen provides high-throughput genotyping of MTBC hence providing epidemiological informations like prevalence or introduction of a particular genotype. This protocol targets variable sized regions of MTB genomic DNA using

MIRU-VNTR approach. This platform is further improvised to detect RIF and INH resistant alleles (80-81).

Unlike other bacterial pathogens that require drug resistance through plasmid acquisition, transposon or phage mediated elements, *M. tuberculosis* acquire drug resistance through chromosomal mutations particularly, single-nucleotide polymorphisms (SNPs) (80-81). This makes identification of drug resistant alleles difficult particularly for those lying outside of drug resistant hotspots, e.g. RIF resistance-determining region for RIF or the alleles that lie outside of *rpoB* and associated with resistance (77). Newer approaches

to detect drug resistance using Next-generation-sequencing (NGS) involves PZAresistance via deletions, STR, INH and screening allelic variation associated with resistance to novel drugs (80,82-85). NGS is very useful in determining whether the case of relapse is because of infection with new strain or incomplete treatment (86). WGS is already applied to the patients with MDR-TB (87-88). McKinsey has even listed NGS as one of the 12 technologies that will transform our lives.

### 11.1. New sequencing protocols

from Illumina. USA facilitates MiSea focused applications like targeted gene sequencing, metagenomics, small genome sequencing, targeted gene expression, amplicon sequencing and HLA typing. Only 10ng of DNA is required as starting material. MiSeq gives output of 15 Gb with 25 M sequencing reads of 2x300 bp read lengths). Personal Genome Machine® (PGM™) System (Life Technologies Incorporated; USA) is the conjugation of semiconductor sequencing technology with natural biochemistry to provide realtime chemical information as digital data in less than 3 hours. This technique is accessible to all kind of set-ups. Use of simple natural sequencing chemistry eliminates the requirement of costly optics and complicated chemistries to measure DNA extension. 454 FLX Junior (Roche Diagnostics; Switzerland) provides rapid and robust platform to sequence and assembly genome of any size without the need for supplementary Sanger data. It provides the output having double (1000 bp) the read length of the Titanium kits, hence an improvisation over current GS FLX. The applications like transcriptome and amplicon sequencing enable an all inclusive picture of gene space of organisms. GeneReader from Qiagen can complement available protocols for resistance determination. It can detect allelic variants at 1% allelic frequency. Data from liquid biopsy pipeline show 100% conformity with PCR-based variant detection methods. GnuBIO from Bio-Rad Laboratories is a novel low cost desktop sequencer. It comes up with a highly integrated platform that incorporates all the functions of DNA sequencing like target selection. DNA amplification. DNA sequencing and analysis into a single, integrated system. Single user interface with single step process ensures rapid genomic results in hours. GridION™ and the Min- ION™ from Oxford Nanopore Technologies. UK facilitates rapid assembly of genome data for a MDR isolate, which elucidates the distribution and underlying mechanism of drug resistance. It enables the selection of specific DNA molecules for sequencing by reversing the driving voltage across individual nanopores (89).

# 12. ASSAYS BEING DEVELOPED / EVALUATED UPCOMING/PIPELINE TECHNOLOGIES

There is a co-evolution of sort between TB bacilli invading to newer regions and newer

techniques / protocols as a counter offensive. Many such existing protocols are improvised and upgraded in response to the challenges and threat posed by rapidly emerging and evolving drug resistant TB bacilli. Following protocols / platforms are under development to spearhead the fight of humanity against TB:

Acombine of nano- and biosensing technology is being evaluated for its use as a portable, rapid on-the-spot biosensor for MTB detection. For instance, detection of trans-renal DNA opens up new avenues for TB detection at molecular level. This technique is not yet commercialized due to the problems in the development of TB detection / readout assays.

High-resolution melting (HRM) curve analysis coupled with closed-tube RT-PCR provides a good screening method with a positive predictive value (PPV) of 100% and negative predictive value (NPV) of 99.9.%, for high throughput screening of large number of specimens in any TB laboratory. Improvized amplification techniques in conjunction with 'molecular beacon' approach (LATE-PCR) offer future advancements, particularly in drug resistance analysis (90-91).

IR spectrophotometer based detection of isotopic urea in exhaled breath is another future technique that holds promise. It utilizes MTB urease as a bacterial virulence factor (6).

A biophotonic detection platform is being developed that utilizes reporter enzyme fluorescence to detect  $\beta$ -lactamase produced by MTB. This innovative technology is now being adapted for point of care (POC) use.

Becton Dickinson is in the process of designing an automated platform that will do the staining of AFB in sputum in addition to imaging. This platform will provide the sensitivity similar to the fluorescent microscopy at the rate of processing around 40 slides per day with slide scanning algorithm for scoring (2).

Xiamen Zeesan Biotech Co. Ltd (China) has developed MeltPro® Drug-Resistant TB Testing Kits as RT-PCR enabled detection of resistant alleles to RIF and INH. It is undergoing further improvisation to detect alleles resistant to ethambutol (EMB), streptomycin (STR) and fluoroquinolones (FLQs) (52-53, 92).

Genedrive® MTB assay is upgraded to include genotyping MDR-TB through RIF resistance in addition to MTBC diagnosis. This assay involves manual preparation of samples followed by RT-PCR115 (93-94).

The Northwestern Global Health Foundation (USA) in partnership with Quidel Inc. (USA) is coming

up with an integrated MTBC assay. The assay involves lysing MTBC cells via heat block followed by purifying DNA using immiscible oil interface (68).

Q-POC™ platform, under development from QuantuMDx (UK) offers diagnosis for PTB.

The assay involves coughing the sputum in a cup that contains reagents for its liquification and decontamination. This is followed by concentration of MTBC cells via paramagnetic bead concentration step and quick nucleic acid amplification (4 minutes) via isothermal asymmetric PCR (2).

Tangen Biosciences Inc. (USA) is going to launch a fully integrated TB detection platform. This platform will have provision for sputum collection in a cup to capture MTB cells followed by lysis and isothermal amplification via LAMP. It comes up with the storage capacity of 10,000 results (2).

### 13. CONCLUSIONS

Accurate and timely diagnosis of TB is the only requirement for its control and management. Currently, TB diagnosis and treatment is plaqued by factors, like inability to diagnose correctly latent and active TB; lack of precise diagnostic protocols for EPTB, pediatric TB and TB associated with HIV; prolonged diagnostic protocol aggravating the otherwise curable infection and escalating it to MDR/XDR-TB; absence of readily available testable biomarkers; Inaccessibility to the POC diagnostic facilities in the peripheral laboratories. These are further challenged by the rapidly emerging resistant strains of TB bacilli in the regions where it is thought to be eradicated or where it was non -existent historically. Need of the hour is the tools and techniques that are accessible. accurate, rapid, economical and most importantly rapid as time lag in diagnosis aggravates the infection. Although TB diagnostics have seen tremendous upsurge in the newer technologies and protocols from imaging, molecular NAAT based, immunological, culture based phenotypic methods, array based to genome sequencing protocols. Unfortunately, they offer promise in some aspect but lack or have severe limitation in other aspects, which limits their applicability on 'one -shoe-fit-all' basis. Another, problem with newer protocols is the reluctance of user groups to embrace it either due to lengthy approval mechanism. lack of WHO approval in Toto or simply the cost and expertise involved. There is urgent need to evolve the mechanism to accept the newer protocols on the emergency basis. As an extension to the problem, many small companies with promising diagnostic kits (in terms of good performance on heterogeneous set-ups) do not have sufficient funds to market their products or scale up their production. Also, most of the newer technologies are accepted in the developed

countries but there is a need to evolve the technologies compatible to the requirement of resource limited developing countries. There is a dire need of all inclusive diagnostic protocols with applicability in wide spectrum of patients and set-ups with high sensitivity and specificity. A dynamic understanding of existing and forthcoming technologies is key in facilitating access to appropriate TB diagnostic tools through market-based interventions. As such, this review is intended to be a living document, updated as the TB diagnostics market evolves, to highlight potential.

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Abbreviations: CXRs, Chest X-ray; CAD4TB, Computer aided detection for tuberculosis; DST, Drug susceptibility test; PAT, Phage Amplification Technique; PNA, Peptide nucleic acids; LAM, lipoarabinomannans; TAM-TB, T-cell activation marker-tuberculosis; IGRA, Interferon-gamma release assays; ESAT-6, Early secretory antigen target 6; LAMP, loop-mediated isothermal amplification; LCR, ligase chain reaction; LPA, line probe assay; WGS, Whole genome sequencing; MIRU-VNTR, mycobacterial interspersed repetitive unit-variable-number tandem repeats; SNP, single-nucleotide polymorphism.

**Key Words:** *Mycobacterium tuberculosis* complex, MDR, XDR-TB, extrapulmonary TB, EPTB, pulmonary TB, PTB, sputum smear microscopy (SSM), Nucleic acid amplification tests, NAAT, Review

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