WHO recommended tNGS, its application at the diagnostic algorithm and implementation consideration

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### **Pipeline of TB diagnostics/WHO guidelines**



2025

## **Comparison of different sequencing approaches**

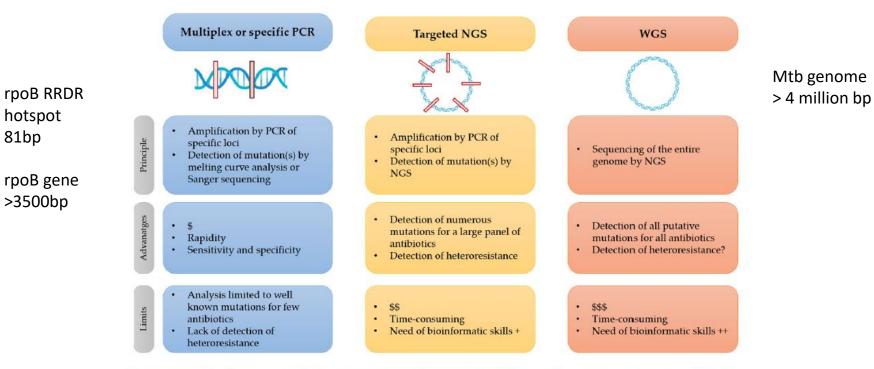


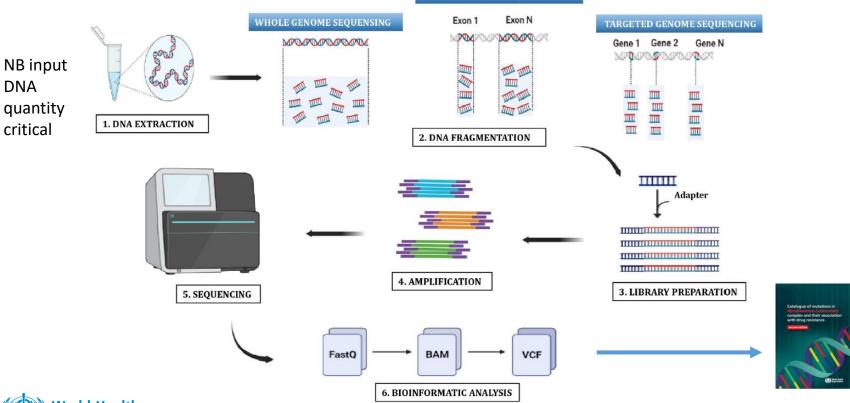
Figure 1. Advantages and limits of molecular techniques for detection of resistance mutations in MTB.



Beviere, et al. The Role of Next-Generation Sequencing (NGS) in the Management of Tuberculosis: Practical Review for Implementation -in Routine. Pathogens **2023**, 12, 978. <u>https://doi.org/10.3390/pathogens12080978</u>

## **Sequencing process**

- Depth of coverage
- Width of coverage





Satam, et al. Next-Generation Sequencing Technology: Current Trends and Advancements. Biology 2023, 12, 997. https://doi.org/10.3390/biology12070997

WHOLE EXOME SEQUENSING

### **Class of targeted NGS products**

- Uses massively parallel sequencing to detect resistance to TB drugs
- Starting from a processed clinical sample
- Ending with an end-user report that relates detected *M.tb* mutations to the presence (or absence) of drug resistance, based on the interpretation of a standard catalogue of mutations.

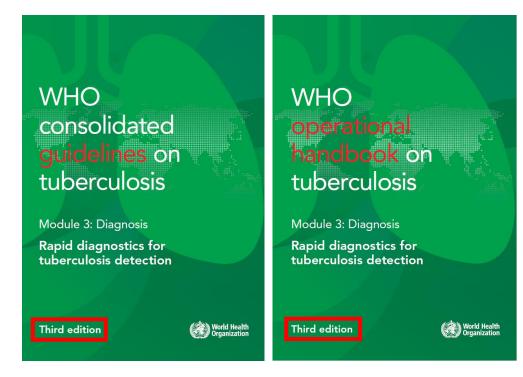


## WHO GDG process

- In 2022, WHO has commissioned a series of systematic reviews of published and unpublished data on the class of targeted NGS products that are commercially developed for TB drug resistance detection.
- The systematic reviews included data on diagnostic accuracy, economic information, and qualitative evidence on feasibility, acceptability, equity and end-user values and preferences.
- WHO convened a GDG on 2–5 May 2023 to discuss the findings of the systematic reviews and make recommendations on this technology.
- 25<sup>th</sup> July WHO has issued rapid communication including early results of the process
- Issue of full recommendations: March 2024



## WHO guidelines and handbook on diagnostics for TB





https://www.who.int/publications/i/item/9789240089488

### **Products included in the evaluation**

- Deeplex<sup>®</sup> Myc-TB (GenoScreen): for rifampicin, isoniazid, pyrazinamide, ethambutol, fluoroquinolones, bedaquiline, linezolid, clofazimine, amikacin and streptomycin.
- AmPORE-TB<sup>®</sup> (Oxford Nanopore Technologies): for rifampicin, isoniazid, fluoroquinolones, linezolid, amikacin and streptomycin.
- **TBseq**<sup>®</sup> (ShengTing Biotech): for ethambutol.



### The test performance among people with RR-TB

- Accurate for isoniazid, levofloxacin, moxifloxacin, pyrazinamide and ethambutol (pooled sensitivity ≥ 95%)
- Acceptable for bedaquiline (68%), linezolid (69%), clofazimine (70%), amikacin (87%) and pyrazinamide (90%)
- The specificity was ≥ 95% for all drugs except streptomycin (75%)



### **Recommendations (bacteriologically confirmed TB)**

In people with bacteriologically confirmed PTB, targeted NGS technologies may be used on respiratory samples to diagnose resistance to RIF, INH, Fq, Z and E rather than culture-based pDST. (Conditional recommendation, certainty of evidence moderate [INH and Z], low [RIF, Fq and E])

### Remarks

- Priority should be assigned to those at higher risk of resistance to first-line treatment medications, including individuals who:
  - continue to be smear or culture positive after 2 or more months of treatment, or experience treatment failure;
  - have previously had TB treatment,
  - are in contact with a person known to have resistance to TB drugs; or
  - reside in settings or belong to subgroups where there is a high probability of resistance to either RIF, INH or FQ (used in new shorter regimens), or where there is a high prevalence of *M*. *tb* strains harbouring mutations not detected by other rapid molecular tests.



### **Recommendations (rifampicin-resistant TB)**

In people with bacteriologically confirmed RR PTB, targeted NGS technologies may be used on respiratory samples to diagnose resistance to INH, Fq, Bdq, Lzl, Cfz, Z, E, Amk and S rather than culture-based pDST. (Conditional recommendation, certainty of evidence high [INH, Fq and Z], moderate [E], low [Bdq, Lzl, Cfz and S], very low [Amk])

### Remarks

- Priority should be given to those at a higher risk of resistance to medications used for the treatment of RR-TB, including individuals who:
  - continue to be smear or culture positive after 2 months or more of treatment or have experienced treatment failure;
  - have previously had TB treatment, including with the new and repurposed drugs;
  - are in contact with a person known to have resistance to TB drugs, including the new and repurposed drugs; or
  - have pre-XDR-TB with resistance to fluoroquinolones.



## Implementation considerations (general)

- **Regulatory approval** from national regulatory authorities or other relevant bodies is required before implementation of these diagnostic tests.
- In its current format, targeted NGS is a high-complexity test most suitable for centralized laboratories equipped with **specialised skills and infrastructure**.
- Targeted NGS tests do not replace existing rapid tests that are more accessible and easier to perform for detecting resistance to RIF, INH and FQ.
  - However, if targeted NGS can be performed rapidly, it can be considered as an alternative initial option for prioritized populations.
- Priority should be given to samples with a high bacillary load as determined by initial bacteriological tests (e.g. semi-quantitative high/medium or smear-positive grading).
  - In situations where the bacillary load is low (e.g. semi-quantitative low/very low/trace or smear-negative grading), the recommendations still hold, although rates of indeterminate results are likely to be higher;
  - therefore, pDST is likely still required for samples with a low bacillary load. orld Health

### Implementation considerations (BDQ, LZD, CFZ)

Since sensitivity for BDQ, LZD and CFZ resistance is **suboptimal**, consideration of the **pretest probability** is important in interpreting the targeted NGS results for these drugs. Further testing of samples with a susceptible result (using culture-based pDST) would be warranted, particularly when the risk of resistance is high. Since specificity is high, a result that indicates resistance may be used to guide the therapy, particularly among those at risk for resistance.

#### Low pre-test prob

- Negative predictive value good
- High pre-test prob
- Positive predictive value is good



Table 65: Should TNGS be used to diagnose drug resistance to bedaquiline (BDQ) (pDST) in patients with bacteriologically confirmed rifampin-resistant pulmonary TB disease?

Sensitivity	0.68 (95% CI: 0.43 to 0.93)					Desuel	ences 1%	3% 5%				
Specificity	0.97 (95% CI: 0.94 to 1.00)					Prevalences 1% 3% 5%						
Outcome		Nº of studies (Nº of patients)	Study design	F	actors that may	y decrease certainty of evidence			Effect per 1,000 patients tested			Te
				Risk of bias	Indirectness	Inconsistency	Imprecision	Publication bias	pre-test probability of1%	pre-test probability of3%	pre-test probability	accur Co
True positives (patients with drug resistance to bedaquiline (BDQ) (pDST))		3 studies 31 patient <i>s</i> ª	cross- sectional (cohort type	not serious <sup>b</sup>	not serious	not serious <sup>e</sup>	very serious	none	7 (4 to 9)	20 (13 to 28)	34 (22 to 47)	⊕⊕( _Lov
False negatives (patients incorrectly classifi not having drug resistance bedaquiline (BDQ) (pDST)	to		accuracy study)						3 (1 to 6)	10 (2to 7)	16 (3 to 28)	
True negatives (patients without drug resis to bedaquiline (BDQ) (pDS	stance 5	4 studies 519 patient <i>s</i> °	cross- sectional (cohort type accuracy study)	not serious <sup>o</sup>	not serious	not serious	not serious	none	960 (931 to 990)	941 (912 to 970)	922 (893 to 950)	⊕⊕€ Hig
False positives (patients incorrectly classifi having drug resistance to bedaquiline (BDQ) (pDST)									30 (0 to 59)	29 (0 to 58)	28 (0 to 57)	

#### b. Explanations

a. This model is not controlled for CT value as that variable was collinear in the original model

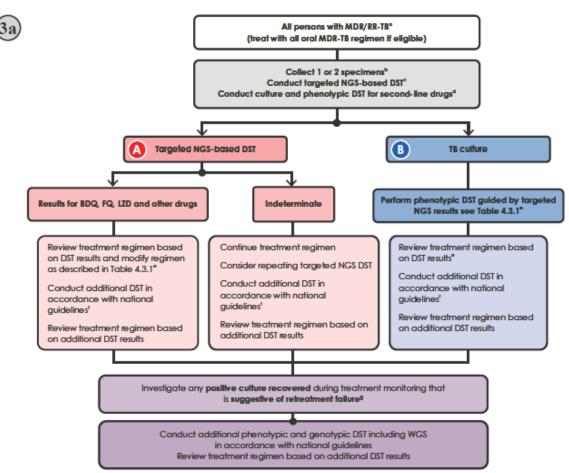
b. Prevalence of resistance to be daquiline across data used in the model was 6% (CI 4% to 8%)

c. One studyhad verylow sensitivity but it onlyhad 3 resistant samples. It identified 0/3

d. Very wide 95 % confidence intervals for sensitivity

e. This model is not controlled for rifampicin resistance as this variable was collinear in the original model. Instead, the data have been restricted to isolated that are resistant to rifampicin by Xpert, and then controlled for CT value.

### Updated diagnostic algorithm using tNGS



### Conclusion

- The new recommendations on targeted NGS marks a major shift in the diagnosis of drug-resistant TB
  - Addresses important gaps in the diagnosis of new and repurposed drugs despite the less optimal performance
- The cost and feasibility are potential barriers but offer important opportunities post-Covid
  - Many countries have sequencers available and have experience
- The best implementation approach is yet to be defined but will likely be context-dependent
- More evidence is needed on the new drugs and the impact of targeted NGS on patient important outcomes

### Acknowledgements

- WHO Global TB Programme
  - Nazir Ismail,
  - Alexei Korobitsyn,
  - Carl-Michael Nathanson
- Other WHO staff at HQ, Regions and Country offices
- Guideline Development Group & other experts
- Staff of national TB programmes in Member States
- USAID, Unitaid, The Global Fund





# Thank you

# It's time for action It's time to END TB

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