



Early View

Original article

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The global prevalence of latent tuberculosis: a systematic review and meta-analysis

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Summary of messages:

Using a novel strategy, global estimates of latent tuberculosis were updated to 24.8% for IGRAs and 21.2% for TSTs using prevalence surveys of 351.811 individuals. Regional estimates varied between 11-27% and 12-33% for IGRAs and TSTs respectively.

Key words:

Latent Tuberculosis; Prevalence; Global Health; Interferon-Gamma Release Assay (IGRA); Tuberculin Skin Test (TST)

Abstract

In 1999, the WHO estimated that one-third of the world's population had latent tuberculosis infection (LTBI) which was recently updated to one-fourth. However, this is still based on controversial assumptions in combination with tuberculin skin test (TST) surveys. Interferon-gamma release assays (IGRAs) with a higher specificity than TST have since been widely implemented, but never used to estimate the global LTBI prevalence.

We conducted a systematic review and meta-analysis of LTBI estimates based on both IGRA and TST results published between 2005 and 2018. Regional and global estimates of LTBI prevalence were calculated. Stratification was performed for low, intermediate and high TB incidence countries and a pooled estimate for each area was calculated using a random effects model.

Among 3280 studies screened, we included 88 studies from 36 countries with 41 IGRA (n=67 167) and 67 TST estimates (n=284 644). The global prevalence of LTBI was 24.8% (95% CI: 19.7-30.0%) and 21.2% (95% CI: 17.9-24.4%) based on IGRA and a 10 mm TST cut-off respectively. The prevalence estimates correlated well to WHO incidence rates ($R_s=0.70$, $p<0.001$).

In the first study of the global prevalence of LTBI derived from both IGRA and TST surveys, we found that one-fourth of the world's population is infected. This is of relevance as both tests, although imperfect, are used to identify individuals eligible for preventive therapy. Enhanced efforts are needed targeting the large pool of latently infected as these individuals continuously constitutes an enormous source of potential active TB.

Introduction

The World Health Organization (WHO) estimated in 1999 that 1.8 billion people, or one-third of the world's population, were infected with *Mycobacterium tuberculosis* (Mtb) but without clinical symptoms of active tuberculosis (TB) which is the definition of latent TB infection (LTBI) [1]. Since, this estimate has been referred to frequently, but has not been updated until recently. In 2016, a WHO endorsed estimate updated the global prevalence of LTBI to 23% corresponding to 1.7 billion people infected worldwide [2,3].

The reactivation rate of LTBI into active disease is controversial, partly as reinfection may occur, but mainly because there are no methods to identify LTBI subjects at highest risk of developing active TB. Nevertheless, the current estimate of the LTBI burden clearly indicates a large reservoir of individuals at risk of developing active TB. Global incidence and mortality rates of active TB have declined since 1990, and the global incidence rate has been decreasing since the WHO goals were appointed in the beginning of the new millennium [3]. Improved attention to LTBI screening and preventive therapy has been pointed out as crucial for The End TB Strategy for 2050 to be achieved [4,5]. It is hardly possible to eliminate TB unless progression to active TB is prevented underlining the need to determine the actual prevalence of LTBI and define hot spot areas [6].

The previous WHO estimate was only based on tuberculin skin test (TST) to a small extent (13%) [1], but mainly on annual risk of infection (ARI) calculated from the incidence of smear positive cases using the Styblo rule [7], derived from empirical data and assumptions on duration of infectiousness and transmissions per year. The ratio assumes that each smear positive case transmits ten infections per year; whereas a newer estimate suggested that this number could be as low as 2-6 [8]. The Styblo rule also assumes that the duration of infectiousness is in general two years, which is now debated and likely to have decreased due to intensified case finding and treatment of active TB. Even in resource poor settings, treatment delay is reduced to an average of 3 months which enables a more rapid sputum conversion and reduction in infectiousness than when the rule was defined [9,10]. Additionally, transmission rates and infectiousness are highly dependent on age distributions, geographical location, drug availability, living conditions and population density [8]. Therefore, the assumption that the Styblo rule – even in its revised form used for the recent

update – still applies in the global TB settings of today could lead to an overestimation of the LTBI prevalence. Hence, basing LTBI prevalence on a rule of thumb, involving assumptions which may not be valid today, is likely to be more imprecise than using real data collected on populations in a large number of countries which does not involve assumptions on transmission rates or infectiousness but incorporates local conditions.

TST has traditionally been used as a screening tool for LTBI due to low direct costs and ease of use. In the last decades, commercial interferon-gamma release assays (IGRAs) have been introduced which solely contain antigens that are absent in Bacillie Calmette-Guérin (BCG)-vaccine strains and require no follow-up test [11]. Thus, IGRAs have superior specificity to TST in BCG-vaccinated populations and in regions with frequent non-tuberculous mycobacteria exposure [12].

So far, no global estimation of LTBI prevalence has been based on IGRAs, and it has been suggested that the prevalence of LTBI might be overestimated by TST compared with IGRA due to the improved specificity [13]. In this study, we aimed to investigate the global prevalence of LTBI based on TB incidence stratified estimates directly derived from both IGRA and TST as these are the tests currently applied to identify individuals for preventive therapy.

Methods

Search strategy and selection criteria

We performed a systematic review and meta-analysis using the Preferred Reporting Items for Systematic Reviews and Meta-analyses (PRISMA) guidelines (Supplementary F) [14]. The protocol was registered on PROSPERO (CRD42019125380).

Studies that reported on the prevalence of LTBI diagnosed with IGRA and/or TST were eligible. Two investigators (A.C. and V.D.M.) searched the databases MEDLINE, Embase, Scopus, and Web of Science for articles published between 1st of January 2005 and the 30th of July 2018 using the following terms in different combinations and constructions depending on the applied database: “Latent tuberculosis” and (“Tuberculosis” AND “Prevalence” AND “Latent”) combined with “Tuberculin test”, “Tuberculin tests”, “Tuberculin skin test”, “Tuberculin skin tests” ,”TST*”, “Mantoux*”, “Interferon-gamma”, “Interferon-gamma

release assay”, “Interferon-gamma release assays”, “Interferon-gamma release test”, “Interferon-gamma release tests”, “IGRA*”, “Quantiferon*”, “QFT*”, “T-SPOT.TB*”, “Enzyme-Linked Immunospot Assay” and “Enzyme-linked Immunospot Assays” (Detailed electronic search strategy available in supplementary A). All search terms were searched in both title, abstract and field keywords. Our search combined free text and subject identifiers as medical subject heading (MeSH) terms. Additionally, we examined several reference lists of relevant articles. We did not set any language restrictions. The search was initiated in 2005 reflecting the widespread availability of stable, quality controlled and commercial IGRAs on the market, and in order to provide an updated analysis with recent surveys. The QuantiFERON-TB (QFT) test introduced in 2001 measured response to the same antigen mixture (purified protein derivate) as TST, while the QFT Gold (QFT-G) introduced in 2005 omitted antigens present in the BCG vaccine and in the ubiquitous non-tuberculous mycobacteria.

Full-texts were obtained for all studies identified by either A.C. or V.D.M. as potentially relevant.

Study eligibility and quality assessment

Following deletion of duplicates, A.C. and V.D.M. screened titles, abstracts or entire articles for exclusion criteria and determined which studies met the eligibility criteria (Supplementary B). Only studies with a sample size of at least 200 were included to avoid selection bias from small studies. Meta-analyses, reviews, cost-effectiveness analyses and non-human studies were excluded. Studies on patients with presumed or active TB were excluded as well as if they did not report on prevalence of LTBI using IGRA-tests (any versions of QFT and/or T-SPOT.TB) or TST. Further, studies targeting risk-groups (i.e. not population-based) such as healthcare workers, drug-users, prison inmates, human immunodeficiency virus (HIV) positive, patients with inflammatory mediated disease among others were excluded to avoid overestimation of LTBI prevalence during targeted screening. Additionally, studies that applied interventions that could affect IGRA and TST results, and studies that selected their population based on specific test results, were excluded. In studies of risk groups along with control groups, control subgroups were included if no exclusion criteria were found (e.g. healthy controls). The most comprehensive paper, i.e. largest sample size or most detailed IGRA/TST results, was included when the same data were reported in more than one publication. Authors were contacted for clarification when the methodology was unclear.

We established criteria for assessment of the quality of the included studies, adopted from the Cochrane collaboration for analytical studies [15], on the STROBE guidelines for reporting observational studies [16] and according to a global prevalence review [17]. All included studies were assessed for quality on four criteria: Quality of sampling method, quality of selection method, response rate, and quality of prevalence assessment. These four criteria were evaluated on a three-point numerical scale (0, 1, or 2) and involved assessment of internal and external validity and attrition bias (Supplementary C).

Data extraction

The following information was extracted: First author, year of publication, study design, study date, study population description, age, exclusion criteria, eligible and invited study population, participants and finally included participants. Enrolment data was used for the full study enrolment when data for selected subgroups were missing. Further, we extracted measures of TB verification, number of individuals with TB at the time of screening, and whether they were excluded from the final results, sample size, type of test (i.e. version of QFT, T-SPOT.TB and TST), and IGRA and TST test-positive as well as TST cut-offs. We only used baseline results from studies employing two-step TST to avoid a boosting phenomenon [18]. Indeterminate QFT results were registered, and whether they were excluded or not, and similarly for indeterminate T-SPOT.TB. When IGRA results for more than one cut-off point were presented, we used the manufactures' instructions for interpretation [19,20]. Due to the availability of LTBI survey data, which varied between years and areas, the latest country-specific incidence rates of active TB (including people living with HIV) and country population sizes were gathered July 2018 from WHO's global TB database [21].

Definition of incidence intervals and evaluation of IGRA and TST results

In order to enable extrapolation of LTBI prevalence to countries with no data, we divided countries into three intervals of TB incidence: Low (0-10 cases / 100 000 person years⁻¹), intermediate (11-120 cases / 100 000 person years⁻¹) and high (>120 cases / 100 000 person years⁻¹). To our knowledge, there is no consensus on defining the upper limit of the intermediate TB incidence interval, and we based our definition on a combination of published data and visual inspection of the latest WHO estimates of TB incidence [3]. To accommodate this uncertainty, we sequentially performed statistical analysis on a upper

limit defined as all numbers between 20 and 150 and included the resulting range of final global IGRA and TST estimates. The upper limit of the low interval was based on WHO's Framework towards Tuberculosis Elimination in Low-Incidence Countries [22].

We calculated individual prevalence estimates for both IGRA and TST used in all studies and the primary estimate was based on a TST 10 mm cut-off and IGRA-tests without considering indeterminate results. However, other ways of calculating estimates were considered as described below. In studies including both QFT and T-SPOT.TB, we calculated a sample size weighted mean prevalence and used the QFT sample size as denominator. When both QFT-G, QFT Gold In-Tube (QFT-GIT) and/or QFT Gold Plus (QFT-Plus) results were presented, we prioritized data from the newest test version (i.e. QFT-GIT or QFT-Plus). The reported estimate was based on excluding indeterminate IGRA results, as it cannot be ascertained whether these are truly positive or negative. In order to compare different strategies for calculating IGRA results, we included the following strategies as a sensitivity analysis: (1) excluding indeterminate results when possible, (2) including indeterminate results in the denominator and regarding them as negative in the numerator, and in worst case scenario (3) including indeterminate results in the denominator and regarding them as positive in the numerator. We did not have access to quantitative IGRA data and thus, the proposed grey zone for QFT results (0.20-0.70 IU/ml) could not be evaluated [23]. We calculated TST results in three different ways: (1) cut-off at 5 mm or as close as possible, (2) cut-off at 10 mm or as close as possible and (3) cut-off at 15 mm or as close as possible. Exact 95% confidence intervals (CI) were calculated for all studies and estimates.

Meta-analysis and statistical analysis

Study prevalence proportions of LTBI based on IGRA and TST were divided into three groups according to the aforementioned TB incidence intervals using WHO TB incidence rates. Firstly, proportions were transformed using the Freeman-Tukey double arcsine method [24]. We assessed a great variation within the study populations, possibly affecting study estimates, and therefore chose to employ a random effects model as also used in similar studies [25]. In the random effects analysis, increased sample size increases the weight of a study, but the more the study result varies from the other studies in the analysis, the more the weight decreases. This prevents very large studies in one country from affecting the overall result, but provides very small studies with a relatively high weight considering their small

population sizes [26]. A pooled inverse variance weighted random effects analysis was performed on each TB incidence group using the DerSimonian and Laird method [27]. Clopper-Pearson 95% CI were calculated for each study and for the TB incidence interval pooled estimates. We calculated weights of each TB incidence interval estimate by dividing the pooled country population size of each TB incidence interval with the global population size. Followingly, we calculated a global prevalence of LTBI by weighting the TB incidence interval prevalence estimates according to the population size represented. Study estimate heterogeneity was evaluated using I^2 statistics for each incidence interval. Further, we assessed the impact on difference between IGRA and TST global estimates by excluding single test studies. Using the TB incidence interval estimates for LTBI, we calculated LTBI prevalence estimates for each WHO region by weighting the three incidence interval estimates according to pooled population sizes of the same TB incidence intervals of each WHO region and compared them with WHO estimates. Finally, a Spearman's rank correlation coefficient was calculated to evaluate the relationship between IGRA- and TST-based LTBI prevalences and WHO incidence rates. Statistical analyses were performed using the meta 4.9-2 package in R (version 3.5.1).

Results

In total, 8328 search results were identified through MEDLINE (n=2024), Embase (n=1936), Scopus (n=2394), and Web of Science (n=1974). After removal of duplicates, 3280 studies remained of which 770 full-text were assessed for eligibility, and 682 were excluded. Eighty-eight quantitative studies fulfilled the criteria for inclusion (Figure 1; studies listed in supplementary D) [13,28–113]. Among 36 countries represented, 41 IGRA (n=67 167) and 67 TST (n=284 644) estimates were available. Annual TB incidence rates ranged from 0.8/100 000 in the United Arab Emirates [28] to 781/100 000 in South Africa [106]. The mean age reported ranged from approximately 51 months [41] to 82.3 years [66].

Among the included studies, 36 used one or more variants of the QFT assay including individuals sampled with QFT Gold (n=8262) [29,36,48,58,59,70,78,83,94], QFT Gold In-Tube (n=56 327) [13,28,39,42,45,49,51,57,60,67,71,72,79–81,83,86–88,93,96,104–107,110,113] and a single study using QFT-Plus (n=829) [89]. Seven studies used TB.SPOT-TB (n=5547) [31,62,65,72,90,104,114] and two of these simultaneous QFT Gold In-Tube [72,104]. In total, 41 studies used one or more IGRA tools (n=67 167 individuals)

[13,28,29,31,36,39,42,45,48,49,51,57–60,62,65,67,70–72,78–81,83,86–90,93,94,96,104–107,110,113,114]. Sixty-seven studies had TST results (n=284 644 individuals) [13,30–38,40–47,49–56,61,63,65,66,68,69,72–77,80,82–87,91–93,95–109,112,113,115]. Twenty studies used both IGRA and TST [13,31,36,42,45,49,51,65,72,80,83,86,87,93,96,104–107,113]. The pooled sample sizes of studies using TST were larger than studies using IGRA in all intervals and largest in the high-incidence interval (Table 1).

The studies included country estimates (n=36) covering all incidence intervals and WHO regions (Supplementary E). A world map was compiled showing all countries with original LTBI prevalence data coloured in darker variants of blue, orange and red, depending on which of the three TB incidence intervals they were within (i.e. low, intermediate or high) (Figure 2). For the remaining countries without any current data (n=159), we used the weighted LTBI prevalence estimate of their respective TB incidence interval, and colored the countries in a lighter version of aforementioned colors.

The global prevalence of LTBI was 24.8% (95% CI: 19.7-29.9%) and 21.2% (95% CI: 17.9-24.4%) according to IGRA and TST (10 mm) results respectively. Prevalence of LTBI by TB incidence intervals for TSTs and IGRAs are shown in the forest plots in figure 3-4. There was a strong monotonic relationship between WHO TB incidence rates and LTBI prevalence based on both IGRAs ($r_s=0.706$, $p<0.0001$) and TSTs ($r_s=0.697$, $p<0.0001$). The between study estimate heterogeneity of the low, intermediate and high TB incidence interval, calculated with I^2 statistics, was 97%, 99% and 99% for IGRA respectively and 100% for all three incidence intervals with TST.

If including indeterminate results in the denominator, and regarding indeterminate results as negative, global prevalence based on IGRA was 24.2% (95% CI: 19.2-29.2%). In a worst-case scenario, regarding indeterminate as positive results, global prevalence was estimated to 26.3% (95% CI: 21.0-31.6%). Based on TST results, we calculated a global prevalence of 24.1% (95% CI: 20.2-28.0%), 21.2% (95% CI: 17.9-24.4%) and 17.4% (95% CI: 14.4-20.4%) using 5, 10 and 15-mm cut-offs. If only considering studies that used IGRA and TST tests concurrently, 20 studies remained with a pooled population of 43 861 (IGRA) and 44 238 (TST) [13,31,36,45,49,51,65,72,80,83,86,87,93,96,104–107,113]. The global IGRA estimate was then 25.2% (95% CI: 19.8-30.7%) (indeterminate results excluded) and the global TST estimate (10 mm) was 27.1% (95% CI: 18.9-35.3%). When calculations were performed sequentially with the upper limit of the intermediate TB prevalence interval

defined as all numbers between 20 and 150, the global estimate ranged between 22.6-25.0% (IGRA) and 20.6-22.3% (TST).

These new estimates based on both IGRA and TST (10 mm cut-off) were lower than WHO estimates in all WHO regions (Table 1). Most notably, new TST estimates of Southeast Asia and Western Pacific were more than one-third lower than the 1999 estimates. IGRA estimates were slightly higher in all WHO regions compared with new TST estimates.

The quality of the studies included varied ranging from 0-8 points out of eight possible (References available in supplementary C). Most studies employed convenience sampling (n=60/88). Information on response rate was presented in 42 out of 88 (47.7%) studies. Indeterminate results were reported in 30 of 41 (73.2%) studies on IGRA of which four had indeterminate results constituting more than 5% of all results [31,60,65,96]. Twenty-seven of 67 (40.3%) studies reported a 15 mm TST cut-off, fifty-six (83.6%) a 10 mm cut-off and 26 (38.8%) a 5 mm cut-off.

Discussion

In this study, we present an update of the global LTBI prevalence estimate, for the first time based directly on both IGRA and TST results, the tests currently being used to diagnose and select LTBI subjects eligible for preventive therapy. Data were gathered from more than 350 000 IGRA and TST results covering all WHO regions. New IGRA and TST estimates were comparable in most regions but systematically slightly higher for IGRAs. Our global estimate of the LTBI prevalence is an update of the WHO estimate from 1999 and very much in line with a similar ARI-based modelling study from 2016 suggesting a global prevalence of LTBI at 23% [2]. Our findings support that the global prevalence of LTBI is no longer a third of the world population, but closer to one-fourth, with large regional differences, which in this test-based study is similar to the old and new modelled estimates and the reductions are in many aspects, although not completely, aligned with the new modelled prevalences by Houben et al.

In accordance with the previous WHO estimates from 1999 as well as with the new estimates by Houben et al, we found that Southeast Asia is the region with the highest LTBI prevalence [1,2]. In contrast to both previous estimates, we observed that Africa had the second highest prevalence with 26.6% (IGRA) and 33.6% (TST) whereas Houben et al interestingly reported

a regional prevalence of only 22% considerably lower than the previous WHO estimate of 35%. According to WHO, the TB incidence rates in Africa have been decreasing since 2005 [3]. However, before 2005, while end-targets for TB fell in other regions, Africa saw a rise since monitoring began in the early 1990's. This could partly explain the slower pace in reducing LTBI prevalence in Africa, as is indicated by our estimates. Especially sub-Saharan Africa is a high-endemic region, and active TB is prioritized due to the high burden of disease and limited resources while LTBI is mainly a concern for close contacts to smear positive TB patients and people living with HIV [116]. Of further interest, we found considerably lower estimates for the Western Pacific Region (WPR) with 20.7% (IGRA) and 20.3% (TST) whereas Houben et al report 27.9% closer to the previous WHO estimate of 36%. Our estimates are more in line with specific estimates for China of 19% [13] which makes the estimate plausible since China constitutes 73% of the WPR population.

Estimating the true rate of LTBI is highly challenging due to the absence of a gold standard for LTBI. As both IGRA and TST detect memory T-cell response to previous Mtb antigen exposure, a positive test is not necessarily associated with infection of viable bacteria [117]. Of note, the tests are insufficient in detecting progression into active TB with very low positive predictive values of 2.7% (95% CI: 2.3-3.2) for IGRAs and 1.5% (95% CI: 1.2-1.7) for TSTs in one systematic review [118]. Recently, a new version of IGRA, QFT-Plus, has been introduced containing additional TB antigens stimulating both CD4 and CD8 T-cells [119] which according to the manufacturer could result in an enhanced sensitivity; however, so far there is a high agreement (>95%) between the QFT-Plus and older versions of the QFT [120,121].

IGRA and TST are currently used to diagnose candidates for preventive LTBI therapy, and consequently, we applied these as surrogate markers for ongoing TB exposure [122]. We chose to present the estimate based on the TST 10 mm cut-offs and based on exclusion of indeterminate IGRA results from the numerator and denominator as a compromise between sensitivity and specificity. The variability for using other strategies was low. It was unexpected that despite a higher specificity of the blood test, almost all LTBI estimates using IGRA were higher than the senescent skin test. However, this may be dependent on the cut-off applied which is not clearly established for any of the tests with regards to LTBI. We speculate that one reason for the systematically higher IGRA estimates, compared to TST, may be that IGRAs are slightly more sensitive to detect LTBI than TST when using the 10

mm cut-off. When a 5 mm cut-off was applied, the tests were more comparable at 24.8% and 24.1% for IGRA and TST respectively. Further, the tests suffer from variability, and the most optimal cut-off levels are under discussion, in particular for the IGRAs where a grey zone for QFT in the range of 0.20-0.70 IU/ml has been suggested [23,123]. We chose to use the established cut-offs suggested by manufacturers and international guidelines as quantitative IGRA results were very rarely available. However, false positive QFT-results in the grey zone do exist and may have overestimated IGRA results marginally [23,124]. Another possible explanation could be a baseline difference in sampling, and we did find a slightly higher estimate from TST when limiting results to studies with both tests. This finding may indicate that IGRAs are not only more specific, but also more sensitive, although this is difficult to assess in the lack of a gold standard test for LTBI. But even though it has previously been perceived that TST is more frequently positive, as it also captures BCG vaccination and other mycobacteria, we may actually here see a display of the fact that in populations where BCG is given at infancy, it has limited impact on the test result [125]. On the other hand, impaired immunity such as HIV may have higher impact on TST results than IGRAs.

Our study has several limitations. Firstly, crude estimates were based on small study population sizes, especially for IGRAs, in studies of varying sampling technique and quality. We performed vast extrapolation with several assumptions; most notably that our pooled estimates of each incidence interval represented the mean prevalence among the large populations represented by each interval. Secondly, the exclusion of patients with inflammatory disease was based on the assumption that this group may show inferior sensitivity to IGRA and TST due to the underlying disease and/or concurrent immunosuppressive therapy. Thirdly, age was assumed representative of the global age prevalence in the study populations included, and was not accounted for in our extrapolation. Age could act as a surrogate marker for accumulated TB exposure and thus be a risk factor as several studies indicate [13,108]. As outlined in supplementary D, the LTBI prevalence data illustrates a consistent effect of age when comparing populations of younger and older surveyed study participants (i.e. South Africa 16% to 56%, Mexico 12% to 36.9%, Singapore 12.6% to 43.4%, Spain 0.9% to 9.3% and USA 1.5% to 8%) but that does not necessarily imply that the studies are not representative. In a meta-analysis based on published prevalence surveys, it is not possible to adjust reliably for age, and we acknowledge that this

may introduce bias. Yet, we have no indications that the surveyed populations are skewed towards being particularly young or old individuals, which would force the prevalence estimate up or down. Moreover, children and adults were represented in all incidence intervals. Our estimate has the strength of being based on individual measurements in populations across a large number of countries across the world instead of being based on mathematical models. Fourthly, the relatively wide confidence intervals around specific prevalence estimates, which makes monitoring of incremental statistical changes in LTBI prevalence difficult, is another limitation. This is a reflection of the data available with low sample sizes in some of the surveys and large total populations in the surveyed countries, including populations with extrapolated prevalences. This variability of study estimates due to large and diverse populations is also reflected in the high heterogeneity ($\geq 97\%$) calculated with I^2 statistics. We believe that although this represents uncertainty in our estimates, all studies not excluded represent important parts of the total background population which is undoubtedly diverse. Yet an indication that the prevalences should be interpreted with caution. Fifthly, the included studies spanned 15 years and we assumed no development in TB prevalence during this period of years. Although these parameters were accounted for in the modelling study by Houben et al, we found remarkably comparable estimates [2]. Finally, we have excluded data on high-risk populations, and their contribution to the global burden of LTBI may therefore not be sufficiently represented. Studies with a focus on migrant populations were excluded, and migrants may not have been well-represented in population survey, which may also have led to an underestimation of the LTBI prevalence; in low-incidence countries, they may constitute the majority of LTBI cases, e.g. in Australia where Australian-born are stipulated to contribute with only 6.8% of all with LTBI [126].

As always in meta-analyses, the selection of studies may lead to bias, and extrapolations to countries with no data will most certainly introduce bias. Yet, it is important to keep in mind that models and estimations are also not free of bias, in particular if based on data of active TB and assumptions on infectiousness, which we hold is more uncertain than published survey data on IGRA/TST results. Models using the Styblo rule will also face difficulties in taking the age factor into consideration in case of changing epidemiology and population structures; hence, modelling may not lead to a better estimate, if the estimate is based on assumptions that are difficult to adequately predict. Children for instance are less likely to transmit than adults but perhaps that is controlled for, as they are less likely to be smear positive. However, this depends on the age and may not apply to children >15 years. With no

golden standard for diagnosing LTBI, and no method available to measure viable Mtb and to distinguish between ‘true’ LTBI and cleared infection, we believe that assessment of global prevalences of LTBI must be based on the measurements currently available and those used in clinical practice. The decision to initiate preventive therapy will not be based on assumptions but on individual testing, and although age is important in the assessment of the probability of infection, in the end, it will be the test outcome that determines who is eligible for therapy and who is not.

In conclusion, we estimate one-fourth of the world’s population to be latently infected with TB, in the first study applying both IGRA and TST surveys. LTBI still represents an enormous reservoir of potential reactivated TB and this must be recognized as a considerable obstacle, and as a point of intervention, in reaching The End TB Strategy goals of 2050.

Support statement

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Contributors

CW and AC conceived and designed the study. All data were collected by AC and VDM. AC and VDM had access to all obtained data and conducted statistical analyses. AC drafted the first manuscript with contributions from VDM, TS and CW. All authors interpreted data as well as contributed with intellectual content to the final manuscript. CW and TS were study supervisors. All authors agree with the results and conclusions of this article.

Declaration of interests

We declare no competing interests.

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Tables

Table 1: A) Pooled sample sizes of included studies and B) new and old estimates of latent tuberculosis infection prevalence by WHO region

Panel A. Pooled sample sizes by interferon-gamma release assay (IGRA), tuberculin skin test (TST) (10 mm cut-off) and in total. Categorization of countries in TB incidence intervals are listed in the supplementary E. **Panel B.** Global and regional prevalences of latent tuberculosis infection displaying 1999 WHO estimates [1], new modelling estimates from 2016 by Houben et al [2], and our current estimates, including IGRA survey data.

A	Pooled sample sizes of included studies			
TB incidence intervals	IGRA sample size (%)	TST sample size (%)		Total sample size (%)
Low	16 628 (24.8)	104 379 (36.7)		121 007 (34.4)
Intermediate	37 392 (55.7)	63 432 (22.3)		100 824 (28.7)
High	13 147 (19.6)	116 833 (41.0)		129 980 (36.9)
Total sample size (%)	67 167 (19.1)	284 644 (80.9)		351 811 (100.0)
B	Prevalence of latent tuberculosis infection			
	1999 WHO estimates [1], %	2016 Houben et al estimates [2], % (CI)	New estimates including IGRA data, % (CI)	
WHO region	TST*	ARI based on TST surveys and WHO TB prevalence data	TST**	IGRA
Africa	35	22.4 (20.6-24.6)	26.6 (23.0-30.2)	33.6 (24.4-42.9)
The Americas	18	11.0 (7.0-20.0)	13.5 (9.7-17.2)	13.7 (11.0-16.3)
Eastern Mediterranean	29	16.3 (13.4-20.5)	21.1 (17.4-24.8)	24.0 (19.4-28.5)
Europe	15	13.7 (9.8-19.8)	11.8 (8.6-15.0)	12.2 (9.8-14.5)
Southeast Asia	44	30.8 (28.3-34.8)	27.7 (23.6-31.8)	36.0 (25.3-46.7)
Western Pacific	36	27.9 (19.3-40.1)	20.3 (15.0-25.7)	20.7 (16.8-24.5)
Total	32	23.0 (20.4-26.4)	21.2 (18.0-24.4)	24.0 (18.8-29.3)
TB, tuberculosis. IGRA, interferon-gamma release assay. TST, tuberculin skin test. WHO, The World Health Organization. CI, 95% confidence interval.				
*Partly based on annual risk of infection and Styblo’s rule. **10 mm cut-off for TST.				

Figure legends

Figure 1: Study selection.

Flow chart of study inclusion. Studies are listed in supplementary D. TB, tuberculosis. HIV, human immunodeficiency virus. LTBI, latent tuberculosis infection.

Figure 2: World map of countries by TB incidence

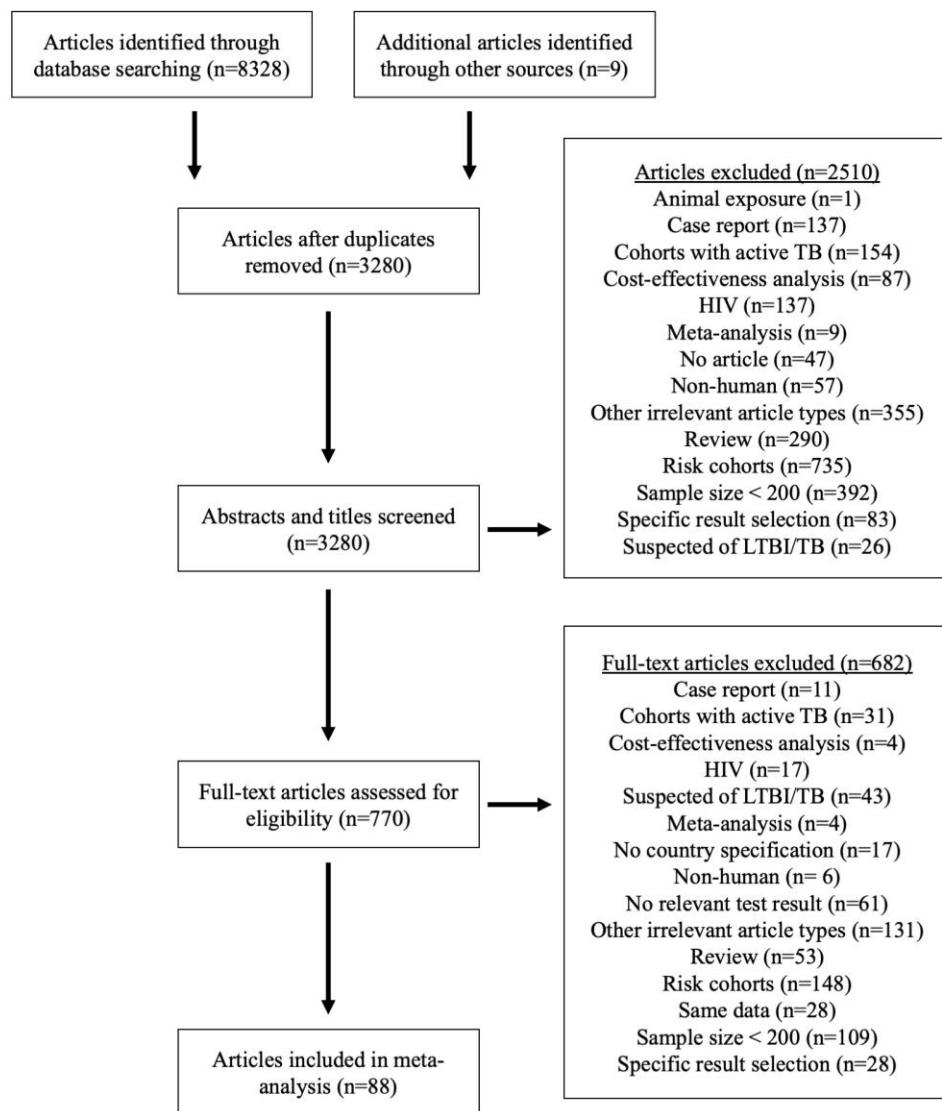
High, intermediate and low active tuberculosis (TB) incidence countries are coloured red, orange and blue respectively. The color red corresponds to an average LTBI prevalence of 28-36%, orange indicates 19-20% and blue 3-5% (from fig. 3 & 4). Darker shades of the colours indicate areas with original latent tuberculosis infection (LTBI) prevalence data, lighter shaded colours indicate countries where the weighted estimate of the countries TB incidence interval has been used.

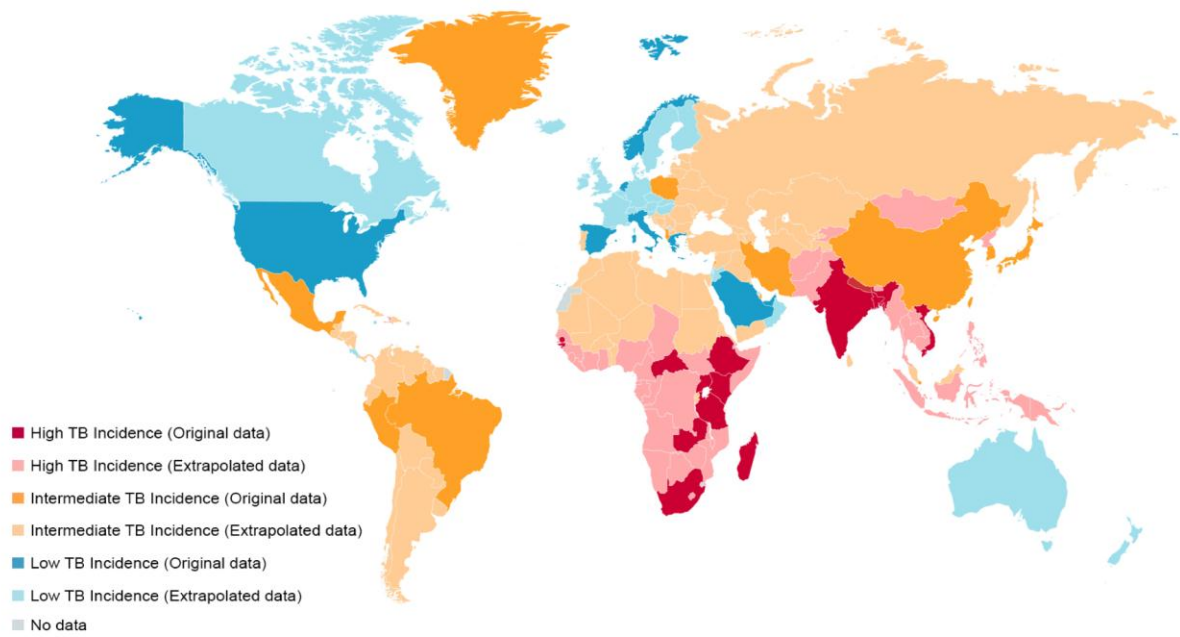
Figure 3: Forest plot of tuberculin skin test data

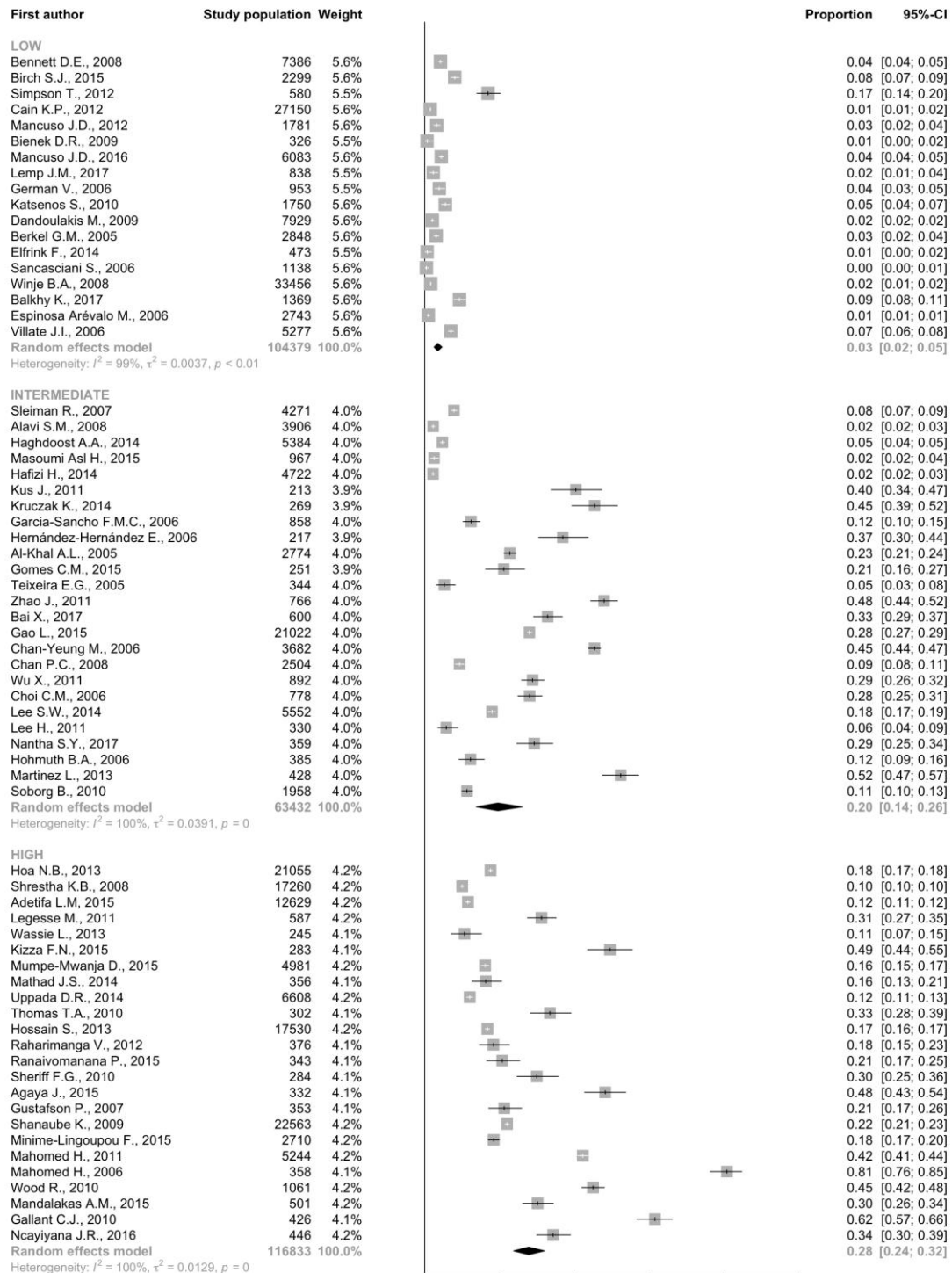
Prevalence of latent tuberculosis infection by tuberculosis incidence intervals using random effects model, weighted by standard error of the mean estimates. Latent infection based on tuberculin skin tests with a 10 mm cut-off. CI, confidence interval.

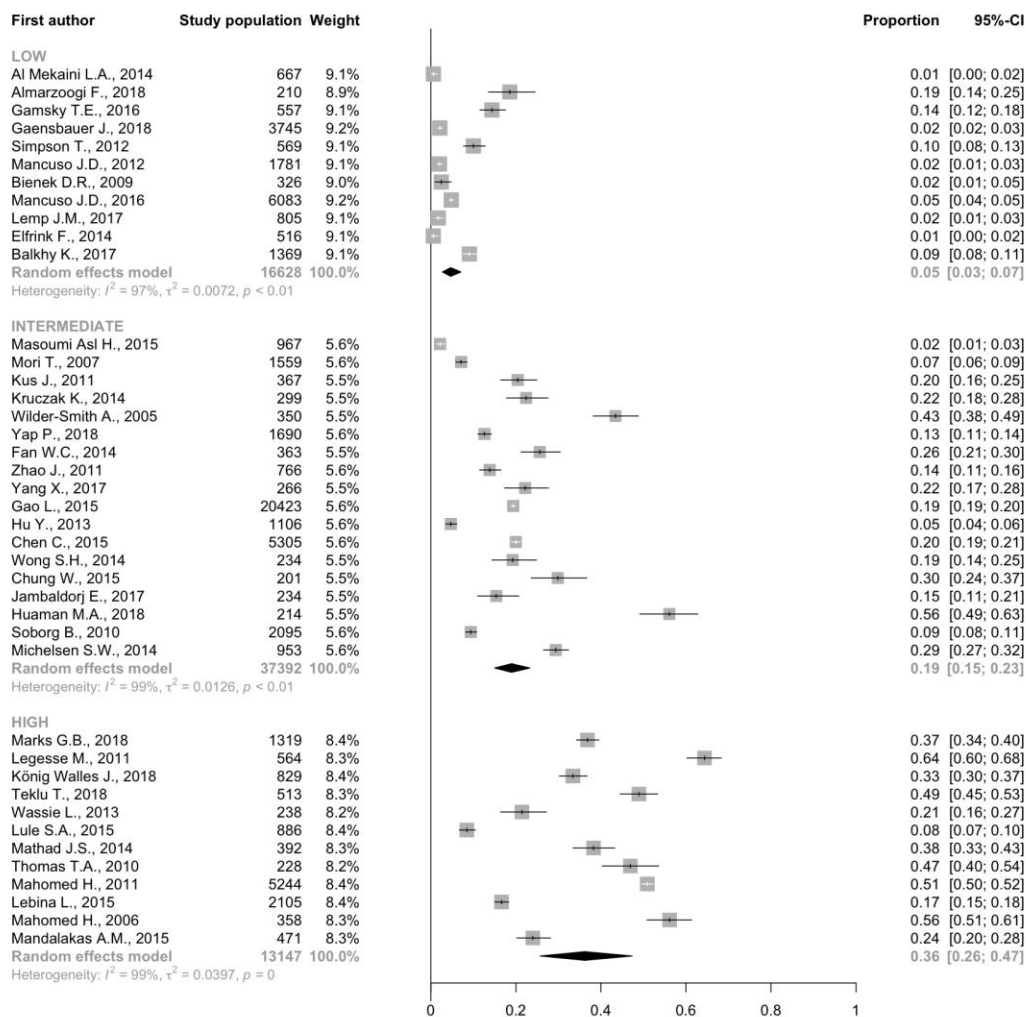
Figure 4: Forest plot of interferon-gamma release assay data

Prevalence of latent tuberculosis infection by tuberculosis incidence intervals using random effects model, weighted by standard error of the mean estimates. Latent infection based on interferon-gamma release assays with indeterminate excluded. CI, confidence interval.









Supplementary material

The global prevalence of latent tuberculosis: a systematic review and meta-analysis

Supplementary A. Search strategies

The following list shows the stepwise searches in the applied databases 30th of July 2018, resulting in the total number of abstracts screened.

MEDLINE

#	Searches	Results
1	"Latent Tuberculosis" OR "Latent Tuberculosis"[Mesh]	3891
2	"Tuberculosis" AND "Prevalence" AND "Latent"	943
3	"Tuberculin Test"[mesh] OR "Tuberculin test" OR "Tuberculin tests" OR "Tuberculin skin test" OR "Tuberculin skin tests" OR "TST*" OR "Mantoux*"	19 136
4	"Interferon-gamma Release Tests"[mesh] OR "Interferon-gamma" OR "Interferon-gamma release assay" OR "Interferon-gamma release assays" OR "Interferon-gamma release test" OR "Interferon-gamma release tests" OR "IGRA*" OR "Quantiferon*" OR "QFT*" OR "T-SPOT.TB*" OR "Enzyme-Linked Immunospot Assay" OR "Enzyme-linked Immunospot Assays" OR "Enzyme-Linked Immunospot Assay"[mesh]	87 491
5	(#1 AND #3) OR (#1 AND #4)	2128
6	(#2 OR #5) Filters activated: Humans. Publication date from 2005/01/01 to 2018/07/30.	2024

Embase

#	Searches	Results
1	exp latent tuberculosis/	4304
2	"latent tuberculosis".ab,ti	4204
3	1 or 2	5899
4	("Tuberculosis" and "Prevalence" and "Latent").ab,ti.	1200
5	exp tuberculin test/	20 006
6	("Tuberculin test" or "Tuberculin tests" or "Tuberculin skin test" or "Tuberculin skin tests" or TST* or Mantoux*).ab,ti.	15 607
7	5 or 6	28 249
8	exp interferon gamma release assay/	2793
9	exp enzyme linked immunospot assay/	9915
10	("interferon-gamma release assay" or "interferon-gamma release assays" or "interferon-gamma release test" or "interferon-gamma release tests" or "Interferon-gamma" or "enzyme linked immunospot assay" or "enzyme linked immunospot assays" or "IGRA" or QuantiFERON* or QFT* or "T-SPOT.TB").ab,ti.	54 582
11	8 or 9 or 10	63 586
12	3 and 7	2640
13	3 and 11	2407
14	4 or 12 or 13	3863
15	limit 14 to (human and yr="2005 -Current" and article)	1936

Scopus

#	Searches	Results
1	TITLE-ABS-KEY ("Latent tuberculosis")	5281
2	TITLE-ABS-KEY ("Tuberculosis" AND "Prevalence" AND "Latent")	1293
3	TITLE-ABS-KEY ("Tuberculin test" OR "Tuberculin tests" OR "Tuberculin skin test" OR "Tuberculin skin tests" OR "TST*" OR "mantoux")	31 039
4	TITLE-ABS-KEY ("Interferon-gamma" OR "Interferon-gamma release assay" OR "Interferon-gamma release assays" OR "Interferon-gamma release test" OR "Interferon-gamma release tests" OR "Enzyme-Linked Immunospot Assay" OR "Enzyme-linked Immunospot Assays")	71 287
5	TITLE-ABS-KEY ("IGRA*" OR "Quantiferon*" OR "QFT" OR "T-SPOT.TB*")	6170
6	(#4 OR #5)	75 236
7	(#1 AND #3)	2430
8	(#1 AND #6)	2132
9	(#2 OR #7 OR #8) Filters activated: Articles. Publication date from 2005 to 2018/07/30.	2394

Web of Science

#	Searches	Results
1	TOPIC: ("Latent tuberculosis")	4316
2	TOPIC: ("Tuberculosis" AND "Prevalence" AND "Latent")	1027
3	TOPIC: ("Tuberculin test" OR "Tuberculin tests" OR "Tuberculin skin test" OR "Tuberculin skin tests" OR "TST*" OR "Mantoux*")	10 713
4	TOPIC: ("Interferon-gamma" OR "Interferon-gamma release assay" OR "Interferon-gamma release assays" OR "Interferon-gamma release test" OR "Interferon-gamma release tests" OR "IGRA*" OR "Quantiferon*" OR "QFT*" OR "T-SPOT.TB*" OR "Enzyme-Linked Immunospot Assay" OR "Enzyme-linked Immunospot Assays")	77 515
5	#1 AND #3	1388
6	#1 AND #4	1708
7	#5 OR #6	2138
8	#2 OR #7 Filters activated: Articles Publication date from 2005 to 2018/07/30. (Indexes=SCI-EXPANDED, SSCI, A&HCI, CPCI-S, CPCI-SSH, ESCI)	1974

Supplementary B. Exclusion criteria

1. The article is a review or meta-analysis.
2. The article is a case report or case series.
3. The study is a cost-effectiveness analysis.
4. The study population is non-human.
5. The study population is tested with neither the skin tuberculin test nor a variant of the interferon-gamma release assay (QFT-G, QFT-GIT, QFT-Plus, T-SPOT.TB).
6. The sample size of neither test is ≥ 200 .
7. The study was published before 2005.
8. The study includes an intervention before testing that could affect results.
9. Test results are not stratified according to country
10. The study population is selected with culture or radiographic findings indicating infection with any mycobacteria as a mandatory criterion.
11. The study population is selected with specific results of TST or IGRA as a mandatory criterion.
12. The study population is a high-risk group: Drug users, prison inmates, psychiatric patients, indigenous, healthcare workers, homeless, HIV positive, part of a contact investigation or refugee/migrant screening, pre- or post-organ transplanted, sarcoidosis, silicosis, miners, patients with end-stage renal disease and/or in dialysis, patients with inflammatory mediated disease (IMID) or patients with diabetes mellitus.

Supplementary C. Quality Assessment

The table summarizes the quality assessment of all included reports using the following quality assessment framework. Studies are listed in the same order as Supplementary D (by WHO countries and incidence rates per 100,000).

Scoring matrix for 4 quality assessment elements:

C.1. Quality of sampling method

- 0 Convenience
- 1 Randomization
- 2 Multisite randomization / National survey

C.2. Quality of selection method

- 0 No exclusion criterion stated / risk factor is an exclusion criterion
- 1 Exclusion criterion stated (risk factor is not a criterion)
- 2 Means of identification of TB is stated

C.3. Response rate

- 0 Not recorded/reported
- 1 Reported and under 65%
- 2 Reported and 65% or above

C.4. Quality of prevalence assessment

- 0 TST cut-off at 10 mm was not present / Indeterminate IGRA results were not stated.
- 1 TST cut-off at 10 mm was present / Indeterminate IGRA results were stated.
- 2 TST cut-off at 5 or 15 mm was present as well / Indeterminate IGRA results constituted < 10%.

First author and publication year	C.1. Quality of sampling method	C.2. Quality of selection method	C.3. Response rate	C.4. Quality of prevalence assessment	Total
Al Mekaini, L.A. 2014	2	0	0	2	4
Almarzoogi, F. 2018	0	1	1	0	2
Bennett D.E. 2008	2	0	2	1	5
Cain, K.P. 2012	0	0	0	0	0
Mancuso, J.D. 2012	0	1	2	2	5
Lempp, J.M. 2017	0	1	2	2 / 2	5 / 5
Gamsky, T.E. 2016	0	1	0	2	3
Mancuso, J.D. 2016	2	2	2	1 / 2	7 / 8
Simpson, T. 2012	0	0	0	2	2
Gaensbauer, J. 2018	0	1	0	0	1
Birch, S.J. 2015	0	0	0	0	0
Bienek, D.R. 2009	0	1	0	1 / 1	2 / 2
Katsenos, S. 2010	0	2	0	2	4
Dandoulakis, M. 2009	0	0	0	1	1
German, V. 2006	0	0	0	0	0
Berkel, G.M. 2005	0	1	0	2	3
Elfrink, F. 2014	0	1	2	1 / 0	3 / 4
Winje, B.A. 2008	2	2	2	0	6
Sancasciani, D. 2006	0	0	1	1	2
Villate, J.I. 2006	0	1	2	2	5
Espinosa Arévalo, M. 2006	0	0	0	0	0
Balkhy, H.H. 2017	2	1	2	1 / 2	7 / 8
Sleiman, R. 2007	1	2	2	2	7
Alavi, S.M. 2008	2	2	2	2	8
Asl, H.M. 2015	2	2	0	2 / 0	4 / 6
Haghdoost, A.A. 2014	2	0	2	1	5
Hafizi, H. 2014	2	1	0	2	5
Mori, T. 2007	0	0	0	0	0
Kruczak, K. 2014	0	0	0	1 / 2	1 / 2
Kuś, J. 2011	0	1	0	2 / 2	3 / 3
Garcia-Sancho, F.M.C. 2006	2	2	1	2	7

Hernández-Hernández, E. 2006	0	2	2	1	5
Al-Khal, A.L. 2005	0	0	0	2	2
Teixeira, E.G. 2005	0	1	2	1	4
Gomes, C.M. 2015	0	0	0	2	2
Yap, P. 2015	2	0	1	2	5
Wilder-Smith, A. 2005	0	1	2	2	5
Gao, L. 2015	2	2	2	2	8
Wong, S.H. 2014	0	0	0	2	2
Fan, W.C. 2014	0	2	0	2	4
Hu, Y. 2014	2	2	0	0	4
Chan, P.C. 2008	0	0	0	1	1
Wu, X. 2011	0	0	0	2	2
Zhao, J. 2011	0	2	0	2 / 2	4 / 4
Chan-Yeung, M. 2006	0	1	2	1	4
Chen, C. 2015	0	1	2	2	5
Bai, X. 2017	0	0	0	0	0
Yang, X. 2017	0	0	1	0	1
Choi, C.M. 2006	0	2	0	2	4
Jambaldorj, E. 2017	0	2	0	0	2
Chung, W. 2015	0	2	0	2	4
Lee, H. 2011	0	0	0	0	0
Lee, S.W. 2014	0	2	0	2	4
Nantha, S.Y. 2017	0	1	1	1	3
Martinez, L. 2013	2	2	0	2	6
Hohmuth, B.A. 2006	0	1	1	1	3
Huaman, M.A. 2018	0	2	0	2	4
Michelsen, S.W. 2014	0	0	2	2	4
Soborg, B. 2010	0	0	2	2	4
Marks, G.B. 2018	2	2	2	2	8
Hoa, N.B. 2013	2	1	2	1	6
Shrestha, K.B. 2008	2	1	2	1	6
Adetifa, I.M. 2011	2	1	1	1	5
Wassie, L. 2013	2	1	0	1 / 2	4 / 5

Legesse, M. 2011	2	1	0	2 / 2	5 / 5
Teklu, T. 2018	2	1	0	2	5
König Walles, J. 2018	0	0	0	2	2
Lule, S.A. 2015	0	1	1	0	2
Kizza, F.N. 2015	2	0	0	1	3
Mumpe-Mwanja, D. 2015	0	0	2	0	2
Mathad, J.S. 2014	0	1	0	1	2
Uppada, D.R. 2014	0	1	0	1	2
Thomas, T.A. 2010	0	1	2	1 / 1	4 / 4
Hossain, S. 2013	2	0	2	0	4
Raharimanga, V. 2012	0	1	2	2	5
Ranaivomanana, P. 2015	0	2	2	0	4
Sheriff, F.G. 2010	0	2	2	1	5
Agaya, J. 2015	1	0	2	1	4
Gustafson, P. 2007	1	2	0	1	4
Minime-Lingoupou, F. 2015	0	1	2	2	5
Mandalakas, A.M. 2015	0	2	0	1 / 1	3 / 3
Mahomed, H. 2011	0	2	1	2 / 2	5 / 5
Mahomed, H. 2006	0	2	0	2 / 0	4 / 2
Wood, R. 2010	0	1	0	2	3
Gallant, C.J. 2010	0	1	0	0	1
Lebina, L. 2015	0	1	0	2	3
Ncayiyana, J.R. 2016	1	0	1	0	2
Shanaube, K. 2009	2	0	1	2	5

Supplementary D. Supplementary table of LTBI prevalence studies included in the review (by WHO countries and incidence rates per 100,000)

*IGRA prevalence is presented with indeterminate excluded in both the denominator and the numerator. TST prevalence is presented with 10 mm cut-offs or the closest possible. Twenty-eight of 88 studies stated randomization strategies.

In total, 58 studies stated exclusion criteria; 25 studies stated diagnostic strategies for identifying active TB and 31 directly stated current TB disease as an exclusion criterion. 42 studies reported a response rate.

WHO TB incidence per 100,000	Country	Study date	Study area	Sampling method	Age	Sample size (IGRA/TST)	IGRA prevalence*	TST prevalence*	First author and year of publication
0.8	United Arab Emirates	2013/04-2013/09	7 ambulatories in 1 state	Randomly selected children	1-19 yrs	669	0.6%	-	Al Mekaini, L.A. 2014 [28]
	United Arab Emirates	2016/08-2017/05	5 departments at 1 hospital	Convenience sampling	42 yrs (mean)	210	18.6%	-	Almarzoogi, F. 2018 [29]
3.1	United States of America	1999-2000	National multicentre	Random, probability cluster sampling	≥ 1 yrs	7386	-	4.2%	Bennett, D.E. 2008 [30]
	United States of America	2002/03-2006/12	95 health departments, 1 state	Convenience sampling	-	27150	-	1.5%	Cain, K.P. 2012 [31]
	United States of America	2009/04-2009/05	1 military base	Convenience sampling	18-29 yrs	1781	2.0%	3.2%	Mancuso, J.D. 2012 [32]
	United States of America	2004/01-2004/02	1 military base	Convenience sampling	-	810/838	1.7%	2.3%	Lempp, J.M. 2017 [33]
	United States of America	2007/01-2013/10	Emergency responders in one county	Convenience sampling	-	565	14.4%	-	Gamsky, T.E. 2016 [34]
	United States of America	2011-2012	National Health and Nutrition Examination Survey	Probability cluster sampling	> 6 yrs	6083	4.8%	4.4%	Mancuso, J.D. 2016 [35]
	United States of America	2008/01-2009/06	Public health screening	Convenience sampling	-	580	10.0%	16.9%	Simpson, T. 2012 [36]
	United States of America	2011/01-2014/08	14 school-based health centers, 3 paediatric primary care clinics, and 6 family medicine clinics	Convenience sampling	8 yrs (mean)	3745	2.1%	-	Gaensbauer, J. 2018 [37]
	United States of America	2012-2013	1 university	Convenience sampling	-	2299	-	8.0%	Birch, S.J. 2015 [38]
	United States of America	-	Recruit training boot camp	Convenience sampling	18-14 yrs	326	2.5%	0.6%	Bienek, D.R. 2009 [39]
4.4	Greece	2007/11-2008/11	1 military base	Convenience sampling	24.3 yrs (mean)	1750	-	5.5%	Katsenos, S. 2010 [40]
	Greece	1990-2005	5 primary and 3 secondary schools	Convenience sampling	6-14 yrs	7929	-	2.0%	Dandoulakis, M 2009 [41]
	Greece	2005/11-2006/02	Military training center	Convenience sampling	23.5 yrs (mean)	953	-	3.9%	German, V. 2006 [42]

5.9	Netherlands	2000-2001	8 clinics, 1 state	Convenience sampling	--	2848	-	3.1%	Berkel, G.M. 2005 [43]
	Netherlands	2008/12-2011/11	Travel clinic	Convenience sampling	25 yrs (median)	516/473	0.6%	0.6%	Elfrink, F. 2014 [44]
6.1	Norway	2005-2006	School children	Convenience sampling	14-15 yrs	33456	-	1.6%	Winje, B.A. 2008 [45]
6.1	Italy	2002-2003	21 high schools	Convenience sampling	18 yrs	1138	-	0.4%	Sancasciani, S. 2006 [46]
10	Spain	1998	All children turning 7, 1 city	Convenience sampling	7 yrs	5277	-	7.0%	Villate, J.I. 2006 [47]
	Spain	-	22 primary pediatric care clinics	Convenience sampling	51.3 months (mean)	2743	-	0.9%	Espinosa Arévalo, M. 2006 [48]
10	Saudi Arabia	2010/07-2013/03	11 primary health centres	Stratified random sampling technique	26.3 yrs (mean)	1369	9.1%	9.3%	Balkhy, H.H. 2017 [49]
12	Lebanon	2004/02-2004/05	Several schools	Convenience sampling	3-19 yrs	4271	-	7.8%	Sleiman, R. 2007 [50]
14	Iran	2006-2007	Children, southwest Iran	Randomized multi-cluster sampling	10 yrs (mean)	3906	-	2.2%	Alavi, S.M. 2008 [51]
	Iran	2009/10-2010/03	24 schools, 11 day care centers	Multistage random sampling	1-15 yrs	967	2.2%	2.5%	Asl, H.M. 2015 [52]
	Iran	2012-2013	25 urban and 53 rural primary schools	Randomized multi-cluster sampling	7 yrs (mean)	5384	-	4.7%	Haghdoust, A.A. 2014 [53]
16	Albania	2010	National screening	Students	10-13 yrs	4722	-	2.2%	Hafizi, H. 2014 [54]
16	Japan	2003/09-2013/10	1 rural community	Community screening	40-69 yrs	1559	7.1%	-	Mori, T. 2007 [55]
18	Poland	2007/07-2009/09	3 long term care facilities and community, 1 city	Convenience sampling	49 yrs (mean)	300/269	22.4%	45.5%	Kruczak, K. 2014 [56]
	Poland	-	Blood donors and controls, 1 province	Convenience sampling	43.8 yrs (mean)	367/213	20.4%	40.4%	Kuś, J. 2011 [57]
22	Mexico	2000/09-2001/02	Selected schools	Convenience sampling	6 yrs (mean)	858	-	12.4%	Garcia-Sancho, F.M.C. 2006 [58]
	Mexico	1989/01-2000/12	Renal donors	Convenience sampling	31 yrs (median)	217	-	36.9%	Hernández-Hernández, E. 2006 [59]
23	Qatar	2000/01-2003/16	Garment workers	Convenience sampling	28.6 yrs (mean)	2774	-	22.7%	Al-Khal, A.L. 2005 [60]

42	Brazil	2002/03-2003/09	Preclinical medical students, 5 hospitals	Convenience sampling	22 yrs (mean)	344	-	5.2%	Teixeira, E.G. 2005 [61]
	Brazil	-	Primary care unit database	Convenience sampling	50.2 yrs (mean)	251	-	21.1%	Gomes, C.M. 2015 [62]
51	Singapore	2014/04-2015/03	Singapore	National household randomization	18-19 yrs	1690	12.6%	-	Yap, P. 2018 [63]
	Singapore	2002	Vaccination sites	Convenience sampling	49.2 yrs (median)	357	43.4%	-	Wilder-Smith, A. 2005 [64]
64	China	2013/07-2013/09	4 areas	Multisite, randomisation	All	21022/20979	19.4%	28.0%	Gao, L. 2015 [13]
	China	-	Hong Kong	Convenience sampling	48.3 yrs (mean)	234	19.2%	-	Wong, S.H. 2014 [65]
	China	2011/01-2012/08	4 hospitals, Taiwan	Convenience sampling	17.8 yrs (mean)	391	25.6%	-	Fan, W.C. 2014 [66]
	China	2010/01-2010/08	8 schools, 7 districts	Convenience sampling	11-18 yrs	1106	4.7%	-	Hu, Y. 2013 [67]
	China	2002-2004	1 city	City screening	6-14 yrs	2504	-	9.3%	Chan, P.C. 2008 [68]
	China	2009/12-2010/03	Military base	Convenience sampling	17-24 yrs	892	-	28.8%	Wu, X. 2011 [69]
	China	2008	1 university	Convenience sampling	17-24 yrs	766	13.8%	47.9%	Zhao, J. 2011 [70]
	China	2006	35 old age homes, Hong Kong	Convenience sampling	82.3 yrs (mean)	3682	-	45.3%	Chan-Yeung, M. 2006 [71]
	China	2013/07	2 villages, Eastern China	Convenience sampling	≥ 5 yrs	5305	20.0%	-	Chen, C. 2015 [72]
	China	2014	Army recruits Beijing	Convenience sampling	19 yrs (mean)	600	-	32.8%	Bai, X. 2017 [73]
	China	-	Vaccinated control group	Convenience sampling	18-76 yrs	266	22.2%	-	Yang, X. 2017 [114]
77	South Korea	2005/03-2006/01	1 military base	Convenience sampling	20 yrs (mean)	788	-	28%	Choi, C.M. 2006 [74]
	South Korea	2009/01-2015/12	1 hospital	Convenience sampling	45.9 (mean)	234	15.4%	-	Jambaldorj, E. 2017 [75]
	South Korea	2012/08-2014/07	Controls undergoing routine health examinations	Convenience sampling	≥ 18 yrs	201	29.9%	-	Chung, W. 2015 [76]

	South Korea	2006/05-2006/12	Paediatric departments at 4 hospitals	Convenience sampling	3-7 yrs	330	-	5.8%	Lee, H. 2011 [77]
	South Korea	2005 + 2008-2011	Korean Reserve Force Battalion	Convenience sampling	20 yrs (median)	552	-	18.0%	Lee, S.W. 2014 [78]
92	Malaysia	2014/10-2015/12	Primary care clinic	Convenience sampling	63 yrs (mean)	359	-	29.2%	Nantha, S.Y. 2017 [79]
117	Peru	2011/07-2012/01	Cluster of shantytowns, 1 city	Convenience sampling	32.7 yrs (median)	428	-	52.3%	Martinez, L. 2013 [80]
	Peru	2002/03-2003/08	1 university	Convenience sampling	21.1 yrs (mean)	385	-	12.2%	Hohmuth, B.A. 2006 [81]
	Peru	2015/07-2017/03	Large national public hospital networks	Convenience sampling	62 yrs (median)	215	56.1%	-	Huaman, M.A. 2018 [82]
118	Greenland	2012/09-2013/04	Birth cohort, 1 district	Convenience sampling	5-30 yrs	953	29.4%	-	Michelsen, S.W. 2014 [83]
	Greenland	2006-2007	Schoolchildren, 5 towns	Convenience sampling	≥ 5 yrs	2117/1958	9.4%	11.3%	Soborg, B. 2010 [84]
133	Vietnam	2015/06-2016/03	60 subcommunes	Randomized multi-cluster sampling	41 (median)	1319	36.8%	-	Marks, G.B. 2018 [85]
	Vietnam	2006/09-2007/07	Nationwide survey of school children	Stratified cluster sample survey	6-14 yrs	21055	17.6%	-	Hoa, N.B. 2013 [86]
154	Nepal	2006/02-2006/09	Nationwide survey of school children	Randomized multi-cluster sampling	5-7 yrs	17260	10.0%	-	Shrestha, K.B. 2008 [87]
174	Gambia	2011	Schools, national screening	Randomized multi-cluster sampling	6-11 yrs	12629	-	11.5%	Adetifa, I.M. 2011 [88]
177	Ethiopia	2009/03-2009/09	7 schools, 1 city	Randomly picked schools	14.8 yrs (median)	245	21.4%	10.6%	Wassie, L. 2013 [89]
	Ethiopia	2010/04-2010/06	Several administrative units, 1 community	Randomly picked administrative units	30 yrs (median)	570/587	64.4%	31.2%	Legesse, M. 2011 [90]
	Ethiopia	2015/05-2016/02	Pastoral communities in southern Ethiopia	Randomly picked sub-districts	37.2 yrs (mean)	497	48.9%	-	Teklu, T. 2018 [91]
	Ethiopia	2015/11-2016/08	3 public health clinics	Convenience sampling	-	829	33.4%	-	König Walles, J. 2018 [92]
201	Uganda	-	1 city	Convenience sampling	5 yrs	886	8.5%	-	Lule, S.A. 2015 [93]
	Uganda	-	1 city	Randomized sampling	≥ 15 yrs	283	-	49.5%	Kizza, F.N. 2015 [94]

	Uganda	-	1 university	Convenience sampling	12-18 yrs	4981	-	16.1%	Mumpe-Mwanja, D. 2015 [95]
211	India	2011/01-2012/07	1 antenatal clinic	Convenience sampling	-	401	38.3%	16.3%	Mathad, J.S. 2014 [96]
	India	2007/02-2008/05	1 school, 1 junior college	Convenience sampling	11-18 yrs	6608	-	12.0%	Uppada, D.R., 2014 [97]
221	Bangladesh	2009/04-2009/06	Slum area, 1 city	Convenience sampling	12.5 yrs	302	46.9%	33.4%	Thomas, T.A. 2010 [98]
	Bangladesh	2007-2009	School children from 20 urban and 20 rural clusters	Randomized multi-cluster sampling	5-14 yrs	17530	-	16.7%	Hossain, S. 2013 [99]
237	Madagascar	2008/04-2008/05	2 primary schools	Convenience sampling	6-7 yrs	376	-	18.4%	Raharimanga, V. 2012 [100]
	Madagascar	2010/07-2010/10	2 primary and 2 secondary schools	Convenience sampling	6-14 yrs	343	-	20.7%	Ranaivomanana, P. 2015 [101]
287	Tanzania	2008/06-2008/08	1 clinic	Convenience sampling	25.3 yrs (mean)	284	-	29.9%	Sheriff, F.G. 2010 [102]
348	Kenya	2013/06-2013/10	School workers from 34 primary and secondary schools	Randomized sampling	36 yrs (median)	332	-	48.2%	Agaya, J. 2015 [103]
374	Guinea Bissau	1999/05-2000/11	1 neighbourhood	Randomized sampling	All	353	-	21.2%	Gustafson, P. 2007 [104]
407	Central African Republic	2011/02-2011/02	57 primary schools, 2 cities	Convenience sampling	6-12 yrs	2710	-	18.4%	Minime-Lingoupou, F. 2015 [105]
781	South Africa	2008/01-2012/07	Neighbourhood controls	Convenience sampling	0.25-15 yrs	471/501	24.0%	30.1%	Mandalakas, A.M. 2015 [106]
	South Africa	-	1 high school	Convenience sampling	12-18 yrs	5422	50.8	42.2%	Mahomed, H. 2011 [107]
	South Africa	-	Workplaces, 1 rural town	Convenience sampling	18-40 yrs	358	56.1%	80.7%	Mahomed, H. 2006 [108]
	South Africa	2006	Townships, 1 city	Convenience sampling	5-40 yrs	1061	-	45.0%	Wood, R. 2010 [109]
	South Africa	-	2 suburbs, 1 city	Convenience sampling	13.9 yrs (mean)	426	-	61.5%	Gallant, C.J. 2010 [110]
	South Africa	2011/02-10	Schools and creeches, 1 city	Randomly selected schools and creeches	5-7 yrs	2105	16.6%	-	Lebina, L. 2015 [111]
	South Africa	2013/05-2014/03	Urban township	Randomized sampling	32 (median)	446	-	34.3%	Ncayiyana, J.R. 2016 [112]

781/376	South Africa & Zambia	2005	98 schools, 2 countries	Randomized multi-cluster sampling	6-11 yrs	22563	-	22.0%	Shanaube, K. 2009 [113]
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Supplementary E. WHO TB incidence intervals and categorization of countries

TB incidence interval	Countries with original data	Countries prevalence was extrapolated to
Low (0-10 cases / 100,000 person years ⁻¹)	Greece, Italy, Netherlands, Norway, Saudi Arabia, Spain, United Arab Emirates, United States of America	Andorra, Antigua and Barbuda, Australia, Austria, Barbados, Belgium, Canada, Costa Rica, Cuba, Cyprus, Czech Republic, Denmark, Dominica, Finland, France, Germany, Grenada, Hungary, Iceland, Ireland, Israel, Jamaica, Jordan, Luxembourg, Monaco, New Zealand, Oman, Samoa, San Marino, Slovakia, Slovenia, St. Kitts and Nevis, St. Lucia, St. Vincent and the Grenadines, Sweden, Switzerland, Tonga, United Kingdom
Intermediate (11-120 cases / 100,000 person years ⁻¹)	Albania, Brazil, China, Greenland, Iran, Japan, Lebanon, Malaysia, Mexico, Peru, Poland, Qatar, Singapore, South Korea	Algeria, Argentina, Armenia, Azerbaijan, Bahamas, Bahrain, Belarus, Belize, Benin, Bolivia, Bosnia, Brunei, Bulgaria, Burkina Faso, Burundi, Chile, Colombia, Comoros, Cook Islands, Croatia, Dominican Republic, Ecuador, Egypt, El Salvador, Eritrea, Estonia, Fiji, Georgia, Guatemala, Guyana, Honduras, Iraq, Kazakhstan, Kuwait, Latvia, Libya, Lithuania, Maldives, Mali, Malta, Mauritania, Mauritius, Moldova, Montenegro, Morocco, Nauru, Nicaragua, Niger, Niue, Panama, Paraguay, Portugal, Romania, Russia, Rwanda, Sao Tome and Principe, Serbia, Seychelles, Solomon Islands, Sri Lanka, Sudan, Suriname, Syria, Tajikistan, the Former Yugoslav Republic of Macedonia, Togo, Trinidad and Tobago, Tunisia, Turkey, Turkmenistan, Ukraine, Uruguay, Uzbekistan, Vanuatu, Venezuela, Yemen
High (>120 cases / 100,000 person years ⁻¹)	Bangladesh, Central African Republic, Ethiopia, Gambia, Guinea Bissau, India, Kenya, Madagascar, Nepal, South Africa, Tanzania, Uganda, Vietnam, Zambia	Afghanistan, Angola, Bhutan, Botswana, Cambodia, Cameroon, Cape Verde, Chad, Congo, Cote d'Ivoire, Democratic Republic of the Congo, Djibouti, Equatorial Guinea, Gabon, Ghana, Guinea, Haiti, Indonesia, Kiribati, Kyrgyzstan, Laos, Lesotho, Liberia, Malawi, Marshall Islands, Micronesia (Federated States), Mongolia, Mozambique, Myanmar, Namibia, Nigeria, North Korea / Democratic People's Republic of Korea, Pakistan, Palau, Papua New Guinea, Philippines, Senegal, Sierra Leone, Somalia, South Sudan, Swaziland, Thailand, Timor-Leste, Tuvalu, Zimbabwe
Total countries	36	159

Supplementary F. PRISMA checklist

Section/topic	#	Checklist item	Reported on page #
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TITLE			
Title	1	Identify the report as a systematic review, meta-analysis, or both.	1
ABSTRACT			
Structured summary	2	Provide a structured summary including, as applicable: background; objectives; data sources; study eligibility criteria, participants, and interventions; study appraisal and synthesis methods; results; limitations; conclusions and implications of key findings; systematic review registration number.	2
INTRODUCTION			
Rationale	3	Describe the rationale for the review in the context of what is already known.	3-4
Objectives	4	Provide an explicit statement of questions being addressed with reference to participants, interventions, comparisons, outcomes, and study design (PICOS).	3-4
METHODS			
Protocol and registration	5	Indicate if a review protocol exists, if and where it can be accessed (e.g., Web address), and, if available, provide registration information including registration number.	4
Eligibility criteria	6	Specify study characteristics (e.g., PICOS, length of follow-up) and report characteristics (e.g., years considered, language, publication status) used as criteria for eligibility, giving rationale.	4
Information sources	7	Describe all information sources (e.g., databases with dates of coverage, contact with study authors to identify additional studies) in the search and date last searched.	4
Search	8	Present full electronic search strategy for at least one database, including any limits used, such that it could be repeated.	4
Study selection	9	State the process for selecting studies (i.e., screening, eligibility, included in systematic review, and, if applicable, included in the meta-analysis).	4
Data collection process	10	Describe method of data extraction from reports (e.g., piloted forms, independently, in duplicate) and any processes for obtaining and confirming data from investigators.	5
Data items	11	List and define all variables for which data were sought (e.g., PICOS, funding sources) and any assumptions and simplifications made.	5
Risk of bias in individual studies	12	Describe methods used for assessing risk of bias of individual studies (including specification of whether this was done at the study or outcome level), and how this information is to be used in any data synthesis.	5
Summary measures	13	State the principal summary measures (e.g., risk ratio, difference in means).	-
Synthesis of results	14	Describe the methods of handling data and combining results of studies, if done, including measures of consistency (e.g., I^2) for each meta-analysis.	6-7

Section/topic	#	Checklist item	Reported
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			on page #
Risk of bias across studies	15	Specify any assessment of risk of bias that may affect the cumulative evidence (e.g., publication bias, selective reporting within studies).	-
Additional analyses	16	Describe methods of additional analyses (e.g., sensitivity or subgroup analyses, meta-regression), if done, indicating which were pre-specified.	6
RESULTS			
Study selection	17	Give numbers of studies screened, assessed for eligibility, and included in the review, with reasons for exclusions at each stage, ideally with a flow diagram.	7
Study characteristics	18	For each study, present characteristics for which data were extracted (e.g., study size, PICOS, follow-up period) and provide the citations.	7
Risk of bias within studies	19	Present data on risk of bias of each study and, if available, any outcome level assessment (see item 12).	8
Results of individual studies	20	For all outcomes considered (benefits or harms), present, for each study: (a) simple summary data for each intervention group (b) effect estimates and confidence intervals, ideally with a forest plot.	Fig 3-4
Synthesis of results	21	Present results of each meta-analysis done, including confidence intervals and measures of consistency.	7-8, fig 3-4
Risk of bias across studies	22	Present results of any assessment of risk of bias across studies (see Item 15).	-
Additional analysis	23	Give results of additional analyses, if done (e.g., sensitivity or subgroup analyses, meta-regression [see Item 16]).	-
DISCUSSION			
Summary of evidence	24	Summarize the main findings including the strength of evidence for each main outcome; consider their relevance to key groups (e.g., healthcare providers, users, and policy makers).	8
Limitations	25	Discuss limitations at study and outcome level (e.g., risk of bias), and at review-level (e.g., incomplete retrieval of identified research, reporting bias).	8-11
Conclusions	26	Provide a general interpretation of the results in the context of other evidence, and implications for future research.	11
FUNDING			
Funding	27	Describe sources of funding for the systematic review and other support (e.g., supply of data); role of funders for the systematic review.	12

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