

# Male Sex Is Associated With Worse Microbiological and Clinical Outcomes Following Tuberculosis Treatment: A Retrospective Cohort Study, a Systematic Review of the Literature, and Meta-analysis

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**Background.** Although the incidence of tuberculosis is higher in men than in women, the relationship of sex with tuberculosis treatment outcomes has not been adequately studied.

**Methods.** We performed a retrospective cohort study and a systematic review and meta-analysis of observational studies during the last 10 years to assess sex differences in clinical and microbiological outcomes in tuberculosis.

**Results.** In our cohort of 2894 Taiwanese patients with drug-susceptible pulmonary tuberculosis (1975 male and 919 female), male patients had higher adjusted hazards of 9-month mortality due to all causes (hazard ratio, 1.43 [95% confidence interval (CI), 1.03–1.98]) and infections (1.70 [1.09–2.64]) and higher adjusted odds of 2-month sputum culture positivity (odds ratio [OR], 1.56 [95% CI, 1.05–2.33]) compared with female patients. Smear positivity at 2 months did not differ significantly (OR, 1.27 [95% CI, .71–2.27]) between the sexes. Among 7896 articles retrieved, 398 were included in our systematic review describing a total of 3 957 216 patients. The odds of all-cause mortality were higher in men than in women in the pooled unadjusted (OR, 1.26 [95% CI, 1.19–1.34]) and adjusted (1.31 [1.18–1.45]) analyses. Men had higher pooled odds of sputum culture (OR, 1.44 [95% CI, 1.14–1.81]) and sputum smear (1.58 [1.41–1.77]) positivity, both at the end of the intensive phase and on completion of treatment.

**Conclusions.** Our retrospective cohort showed that male patients with tuberculosis have higher 9-month all-cause and infection-related mortality, with higher 2-month sputum culture positivity after adjustment for confounding factors. In our meta-analysis, male patients showed higher all-cause and tuberculosis-related mortality and higher sputum culture and smear positivity rates during and after tuberculosis treatment.

**Keywords.** culture; female; HIV; mortality; sputum.

The World Health Organization reported nearly 10 million new cases of active tuberculosis in 2019 [1]. Compared with women, men have a 1.8-fold higher incidence of active tuberculosis [1, 2]. The tuberculosis prevalence-to-notification ratio is also much higher in men [3, 4]. These sex disparities exist irrespective of geographic locale [5].

Many potential medical and cultural confounding factors, such as the increased prevalence of diabetes, alcohol use, and smoking among men, and decreased access to healthcare among women, may account for sex differences in tuberculosis

[1, 5]. Men have higher sputum bacterial loads and more severe tuberculosis-related lung disease seen at imaging than women [6, 7]. These clinical observations are supported by animal models showing more extensive lung disease, higher lung bacterial burdens, and accelerated mortality in males after *Mycobacterium tuberculosis* infection [8, 9]. Although prior population-based studies have highlighted sex differences in tuberculosis outcomes [10, 11], the findings are often unclear and inconsistent because of the inherent heterogeneity of study populations and the difference in the level of care between the sexes in low-income countries. In addition, many studies do not report sex-disaggregated data.

We analyzed a retrospective cohort of Taiwanese pulmonary patients with tuberculosis to better understand the role of sex on mortality and sputum microbiological status after tuberculosis treatment initiation, after adjusting for confounders. We also performed a systematic review and meta-analysis, given

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the lack of generalizability from individual cohort studies, and to assess the impact of other important parameters, including human immunodeficiency virus (HIV) coinfection status, site of tuberculosis involvement, drug resistance, duration of follow-up, and study country on the effect of sex differences.

## METHODS

### Retrospective Cohort Study

#### Study Design and Population

Our retrospective cohort consisted of adult patients (aged >18 years) with drug-susceptible pulmonary tuberculosis, treated according to the American Thoracic Society guidelines [12], enrolled at the National Taiwan University Hospital in Taipei from 2000 to 2016 [13]. All patients were sputum culture-positive at baseline by either MGIT-960 or Lowenstein-Jensen medium. There were no exclusion criteria. The institutional review boards at Johns Hopkins University and National Taiwan University Hospital approved the study. The methods are described in greater detail in the [Supplementary Materials](#) (Section Ia).

#### Exposure and Outcomes

To ascertain biological sex differences, we considered male sex as the exposure group and female sex as the comparison group. The primary outcomes were 9-month all-cause and infection-related mortality after initiation of antituberculosis therapy (ATT). Infection-related mortality was a composite outcome of deaths due to pneumonia, sepsis, and tuberculosis. Secondary outcomes were positivity of sputum cultures and sputum smear acid-fast bacilli (AFB) by microscopy at 2 months after ATT initiation.

#### Statistical Analysis

Participant characteristics stratified by sex were compared using  $\chi^2$  tests for categorical variables, and 2-sided *t* and Mann-Whitney *U* tests for normally and nonnormally distributed continuous variables, respectively. Kaplan-Meier analysis was performed to evaluate the survival probability of all-cause and infection-related mortality between the sexes. Cox proportional hazards models were used to measure the association between sex and all-cause and infection-related mortality in separate univariable models. Person-time at risk of outcome stratified by sex was calculated from the time of tuberculosis treatment initiation up to 9 months or loss to follow-up or death, whichever occurred first. Separate bivariable models assessed the interaction between sex and major factors, such as age, body mass index, smoking status, and cavitary disease, and AFB smear at baseline was assessed using multiplicative interaction terms. Multivariable Cox regression was designed to adjust for confounding variables identified a priori through literature review, by simultaneously including them in the model. We used Charlson comorbidity index (CCI) as the composite index

to account for the comorbid conditions in the multivariable model. Variables that are components of the CCI were not separately adjusted for. The association of sex with sputum culture and smear positivity at 2 months was analyzed using univariable and multivariable logistic regression. Potential confounders for multivariable analyses were identified, as described above.

### Systematic Review

#### Search Strategy and Study Selection

The systematic review was conducted according to the PRISMA guidelines [14], using the Covidence platform [15]. We searched PubMed, Embase, and the Web of Science on 15 August 2020, for English-language research articles using the search strategy detailed in the [Supplementary Materials](#) (Section Ib) for studies published in the last 10 years.

Studies were required to report sex-disaggregated data on  $\geq 1$  of the following outcomes on adult patients with tuberculosis receiving multidrug ATT: all-cause mortality, mortality due to tuberculosis, sputum AFB smear, or culture positivity during or at the end of tuberculosis treatment, or 'treatment success' according to the World Health Organization definitions for reporting tuberculosis outcomes [16]. We included prospective and retrospective cohort studies and case-control studies. After removing the duplicates, the titles and abstracts screening followed by full-text screening of the retrieved articles were screened by at least 2 authors (V. C., N. L. T., M. G. M., A. K., or P. N.) independently, and the disagreements were resolved by V. C.

#### Data Extraction and Quality Assessment

At least 2 authors extracted data from the articles (V. C., N. L. T., S. K. A., R. K. S., E. P. W., E. J. A., S. W., or A. Z.) in the Qualtrics platform [17], and V. C. resolved discrepancies. Data on study country, funding source, patient comorbid conditions, site of tuberculosis involvement, pattern of resistance to ATT, HIV-tuberculosis coinfection, and time points for outcomes were extracted. Data on treatment outcomes were extracted from studies as either raw data or precalculated effect sizes, namely, odds ratio (OR), relative risk, or hazard ratio (HR), along with 95% confidence intervals (CIs). We performed quality assessment using the Newcastle-Ottawa scale (NOS) for observational studies [18].

#### Data Analysis

For each outcome, we performed a random-effects meta-analysis of the ORs. We pooled HRs separately for each of the outcomes. Differences were considered significant at  $P < .05$  (2 sided). We assessed heterogeneity using the  $I^2$  statistic. When the  $I^2$  was  $>60\%$ , we performed subgroup analyses and meta-regression (detailed in the [Supplementary Materials](#), Section II) with respect to HIV-tuberculosis coinfection status, resistance to ATT, extrapulmonary involvement, time of outcome

assessment, study country's income status classification according to the World Bank [19], and incidence of tuberculosis infection and HIV-tuberculosis coinfection [20]. Publication bias was assessed by means of funnel plot and Egger's test. We performed statistical analyses using Stata/IC 16.0 software (StataCorp) [21]. The study protocol is registered with PROSPERO (no. CRD42020219050).

## RESULTS

### Retrospective Cohort

#### Baseline Characteristics

In our cohort of 2894 patients with culture-confirmed, drug-susceptible pulmonary tuberculosis, 1975 (68.2%) were male and 919 (31.8%) were female (Table 1). Male patients had a higher median age than female patients (68.9 vs 58.2 years;  $P < .001$ ). Compared with female patients, higher proportions of male patients had comorbid conditions, a history of smoking, and alcohol abuse. Men had a greater proportion of cavitory disease (15.7% vs 11.1%;  $P < .001$ ) but similar AFB smear positivity at diagnosis (Table 1). Baseline characteristics are described in detail in the Supplementary Materials, Section Ia.

#### All-Cause and Infection-Related Mortality

In the univariable Cox regression analysis, male sex was associated with higher hazards of 9-month all-cause mortality (HR, 1.75 [95% CI, 1.42–2.14]) and infection-related mortality (1.52 [1.17–1.99]). Infection-related causes accounted for 55.7% of all deaths (303 of 544) in the first 9 months (Table 2). Details of

the bivariable Cox models are detailed in the Supplementary Materials (Supplementary Table 1A–1E). In the multivariable Cox regression analysis obtained by simultaneously adjusting for the following parameters: body mass index, CCI, hypertension, transplantation status, alcoholism, smoking, cavitory disease, and baseline AFB (Table 2 and Supplementary Table 1F and 1G), male patients had an adjusted HR of 1.53 (95% CI, 1.08–2.17) for all-cause mortality (Figure 1A) and an adjusted HR of 1.81 (95% CI, 1.11–2.93) for infection-related mortality (Figure 1B). Although age is a component of CCI, we also adjusted for age in a separate model, which yielded similar results (Supplementary Materials, Section III, and Supplementary Table 1H).

#### Sputum Culture and Smear AFB Positivity

Male patients had significantly higher sputum culture (18.4% vs 11.6%;  $P = .003$ ) and AFB smear (8.3% vs 5.0%,  $P = .02$ ) positivity at 2 months compared with female patients, by  $\chi^2$  test, with a considerable difference primarily among individuals <50 years old (Supplementary Figure A). In the univariable logistic regression analysis, male sex was associated with higher odds of 2-month sputum culture (OR, 1.73 [95% CI, 1.27–2.35]) and sputum smear (1.70 [1.09–2.63]) positivity. Details of the bivariable logistic models are detailed in the Supplementary Materials (Supplementary Tables 1A–1E). In the multivariable logistic regression analysis obtained by adjusting for parameters described above (Table 2 and Supplementary Table 1F and 1G), male patients had adjusted ORs of 1.67 (95%

**Table 1. Characteristics of the Study Participants in the Retrospective Cohort From Taiwan, Stratified by Sex (N = 2894)**

Characteristic	Participants With Available Data, No.	Participants, No. (%) <sup>a</sup>			P Value
		Total (N = 2894)	Male (n = 1975)	Female (n = 919)	
Age, median (IQR), y	2894	66.6 (49.1–77.8)	68.9 (53.4–78.8)	58.2 (38.2–75.4)	<.001
BMI, mean (SD) <sup>b</sup>	2178	21.3 (4.2)	21.4 (4.4)	21.0 (3.6)	.049
Ever smoker	2296	930 (40.5)	884 (56.1)	46 (6.4)	<.001
Alcohol abuse	2894	81 (2.8)	79 (4.0)	2 (0.2)	<.001
Diabetes mellitus	2888	533 (18.46)	403 (20.4)	130 (14.2)	<.001
Hypertension	2894	1052 (36.4)	759 (38.4)	293 (31.9)	.001
CVD	2894	352 (12.2)	265 (13.4)	87 (9.5)	.002
COPD	2894	451 (15.6)	374 (18.9)	77 (8.4)	<.001
Cancer	2888	459 (15.9)	369 (18.7)	90 (9.8)	<.001
Cirrhosis	2894	5 (1.7)	43 (2.2)	7 (0.8)	.007
Transplant recipient	2894	26 (0.9)	18 (0.9)	8 (0.9)	.91
HIV	2894	65 (2.3)	64 (3.2)	1 (0.1)	<.001
CCI, median (IQR)	2894	4 (2–6)	4 (2–6)	3 (1–5)	<.001
Sputum smear AFB positivity at baseline	2842	1215 (42.75)	820 (42.40)	395 (43.50)	.58
Smear grade at baseline, mean (SD)	2842	0.99 (1.36)	1.00 (1.39)	0.96 (1.32)	.50
Prior tuberculosis	1598	98 (6.1)	75 (6.84)	23 (4.58)	.08
Cavitory disease	2894	413 (14.27)	311 (15.75)	102 (11.09)	<.001

Abbreviations: AFB, acid-fast bacilli; BMI, body mass index; CCI, Charlson comorbidity index; COPD, chronic obstructive pulmonary disease; CVD, cardiovascular disease; HIV, human immunodeficiency virus; IQR, interquartile range; SD, standard deviation.

<sup>a</sup>Data represent no. (%) of participants unless otherwise specified.

<sup>b</sup>BMI was calculated as weight in kilograms divided by height in meters squared.

**Table 2. Association of Male Sex With Tuberculosis Treatment Outcomes in the Retrospective Cohort Using Cox and Logistic Regression Models (N = 2894)**

Variable	Mortality, No. of Deaths/Total No. in Population (%)			Estimate	Unadjusted Effect Size <sup>a</sup> (95% CI)	P Value	Adjusted Effect Size <sup>b</sup> (95% CI)	P Value
	Total	Men	Women					
All-cause mortality (n = 2667)	544/2667 (20.4)	427/1837 (23.2)	117/830 (14.1)	HR	1.75 (1.42–2.14)	<b>&lt;.001</b>	1.53 (1.08–2.17)	<b>.03</b>
Infection related mortality (n = 2667)	303/2667 (11.4)	231/1837 (12.6)	72/830 (8.7)	HR	1.52 (1.17–1.99)	<b>.002</b>	1.81 (1.11–2.93)	<b>.009</b>
Sputum culture positivity at 2 mo (n = 1640)	265 (16.2)	203 (18.4)	62 (11.6)	OR	1.72 (1.27–2.34)	<b>.003</b>	1.67 (1.06–2.63)	<b>.03</b>
Sputum smear AFB at 2 mo (n = 1640)	118 (7.2)	91 (8.3)	27 (5.0)	OR	1.70 (1.09–2.63)	<b>.02</b>	1.30 (.67–2.55)	<b>.42</b>

Abbreviations: AFB, acid-fast bacilli; CI, confidence interval; HR, hazard ratio; OR, odds ratio.

<sup>a</sup>Unadjusted effect size obtained from the univariable model.

<sup>b</sup>Adjusted for body mass index, Charlson comorbidity index (CCI), chronic obstructive pulmonary disease, alcoholism, smoking, and cavitory disease at baseline. The variables were selected because they differed significantly between male and female participants (see Table 1). The components of the CCI were not adjusted for separately.

CI, 1.06–2.63) for sputum culture positivity and 1.30 (.67–2.55) for sputum smear positivity. Sensitivity analysis also adjusting for age yielded similar results (Supplementary Table 1H).

**Systematic Review**

**Characteristics of Included Studies and Quality Assessment**

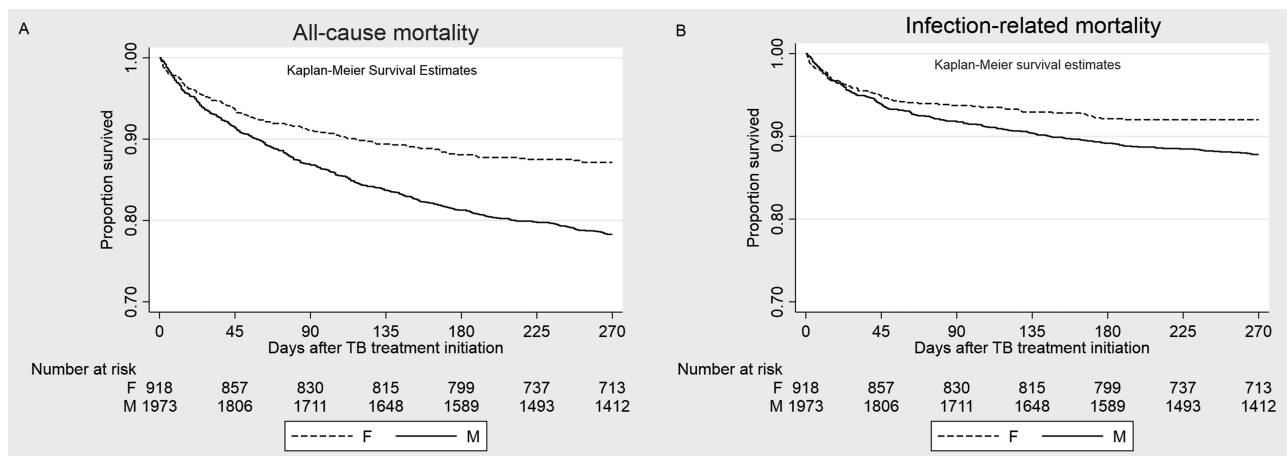
Among the 7896 studies screened, data from 398 studies, reporting on a total of 3 957 216 individuals, were analyzed in our review (Figure 2 and Supplementary Table 2). Characteristics of included studies are described in greater detail in the Supplementary Materials, Section II. Quality assessment using the NOS revealed that 2 studies (0.5%) scored the maximum of 9 points, 190 (47.7%) scored 8, 182 (45.7%) scored 7, and 11 (2.7%) scored 6. A total of 13 studies were of low quality (NOS, ≤5) (Supplementary Table 2).

**All-Cause Mortality and Subgroup Analysis**

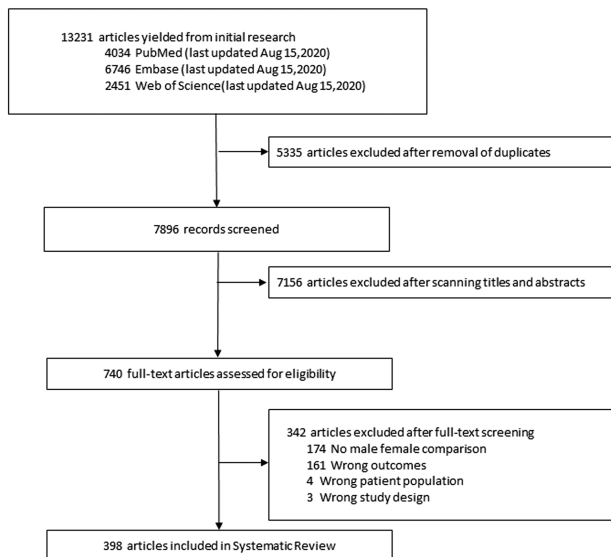
Male patients had higher unadjusted pooled effect sizes for mortality compared with female patients, with a pooled OR of

1.26 (95% CI, 1.19–1.34) from 135 studies ( $I^2 = 77.9\%$ ), and a pooled HR of 1.17 (1.07–1.27) from 44 studies ( $I^2 = 64.5\%$ ) (Table 3). Eighty-two of the 197 studies reporting all-cause mortality made adjustments for ≥1 confounding variable (Supplementary Table 3). The pooled adjusted OR was 1.26 (95% CI, 1.16–1.37) among 52 studies ( $I^2 = 90.9\%$ ), and the pooled adjusted HR was 1.20 (95% CI, 1.05–1.37) among 30 studies ( $I^2 = 82.5\%$ ) (Table 3).

Considerable heterogeneity was present in the meta-analyses of ORs and HRs for all-cause mortality. Studies reporting mortality rates for patients with tuberculosis in intensive care units showed no significant differences in mortality when stratified by sex (Table 3). Further subgroup analyses were performed after excluding studies reporting outcomes in patients receiving intensive care (Table 4). Meta-regression analysis showed that the time point used for assessing mortality did not change the association between male sex and all-cause mortality (Figure 3). Subgroup and meta-regression analyses for all-cause mortality are detailed in the Supplementary Materials (Sections II and III and Supplementary Table 4).



**Figure 1.** Kaplan-Meier survival graphs for all-cause (A) mortality and infection-related (B) mortality among patients treated for tuberculosis in the retrospective cohort. M = Males, F = Females.



**Figure 2.** Study selection for systematic review.

### Mortality Due to Tuberculosis

We included 12 studies that reported death due to tuberculosis (Supplementary Table 7). The pooled OR for death due to tuberculosis for male compared with female patients was 1.28 (95% CI, 1.1–1.6) in 10 studies, and the pooled HR was 1.34 (1.09–1.55) in 2 studies.

### Microbiological Outcomes

Compared with female patients, male patients had higher pooled unadjusted and adjusted ORs for sputum culture positivity after 2 or 3 months of ATT (Table 5). Among studies reporting HR for sputum culture conversion to negativity, men had a pooled unadjusted HR of 0.83 (95% CI, .70–.97) and a pooled adjusted HR of 0.76 (.59–.97), compared with women

**Table 3. Association of Male Sex With All-Cause Mortality Among Patients With Tuberculosis, Using Random Effects Meta-analysis**

Characteristic	All-Cause Mortality					
	No. of Studies	OR (95% CI)	$I^2$ Statistic	No. of Studies	HR (95% CI)	$I^2$ Statistic
<b>Unadjusted</b>						
All studies	135	1.26 (1.19–1.34)	77.90	44	1.17 (1.07–1.27)	63.59
During treatment	122	1.26 (1.19–1.34)	71.30	26	1.16 (1.06–1.27)	59.72
General setting	111	1.26 (1.19–1.35)	77.39	25	1.18 (1.05–1.30)	66.87
ICU setting	11	0.92 (.71–1.18)	0	1	2.57 (.73–9.08)	...
During follow-up	13	1.39 (1.09–1.77)	78.32	18	1.14 (1.00–1.30)	60.27
<b>Adjusted</b>						
All studies	52	1.31 (1.18–1.45)	74.80	30	1.19 (1.05–1.67)	82.5

Abbreviations: CI, confidence interval; HR, hazard ratio; ICU, intensive care unit; OR, odds ratio.

<sup>a</sup>During follow-up after completion of tuberculosis treatment.

(Supplementary Materials, Section IV). With respect to sputum AFB smear positivity, men had higher pooled unadjusted and adjusted ORs compared with women, both at the end of the intensive phase and at treatment completion (Table 5). The odds of treatment success were lower in male patients in the meta-analysis of both adjusted and unadjusted effect sizes (Supplementary Table 5). Data on sputum culture or AFB smear positivity had very low heterogeneity and thus did not warrant further subgroup analysis.

### Meta-Regression Analysis

In the meta-regression analysis, the mean age of the population and the proportions of patients with diabetes mellitus, hypertension, alcohol abuse, smoking, cardiovascular events, chronic obstructive pulmonary disease, and HIV did not significantly modify the association between male sex and any of the outcomes assessed (Supplementary Table 6). Publication bias measured by Egger’s test was not present for any of the outcomes (Supplementary Materials, Section IV). The forest, funnel, and bubble plots from the analysis are available in Supplementary Figures 1–19.

## DISCUSSION

Findings from our retrospective cohort and systematic review, comprising a total of 398 peer-reviewed publications from more than 80 high-, middle-, and low-income countries, suggest that male sex is associated with higher all-cause mortality and death due to tuberculosis, both during treatment and during follow-up after ATT. This association held true even after adjusting for confounders in the cohort study and during meta-regression analysis in the systematic review. Subgroup analysis revealed no impact by the drug susceptibility or the site of tuberculosis disease on the higher mortality rates in male patients. Male patients also had higher positivity for *M. tuberculosis* by both AFB smear microscopy and culture at the end of the intensive phase, as well as at ATT completion.

Women in low- and middle-income countries tend to have greater patient-related and healthcare system delays in tuberculosis diagnosis and treatment initiation [22–24], owing due to financial dependence, fear of social isolation, and initial access to less qualified providers [25–27]. Although our cohort did not include information on patients’ economic status and healthcare delays the patients, we expect these factors to have minimal impact on our study, because Taiwan is a high-income country with universal health insurance. This was supported by the subgroup analysis in our systematic review, which showed higher ORs for mortality in men compared with women in high-income than in low- or middle-income countries (Table 4). This suggests that the observed sex differences are clearer in high-income countries owing to minimal differences in healthcare access between the sexes.

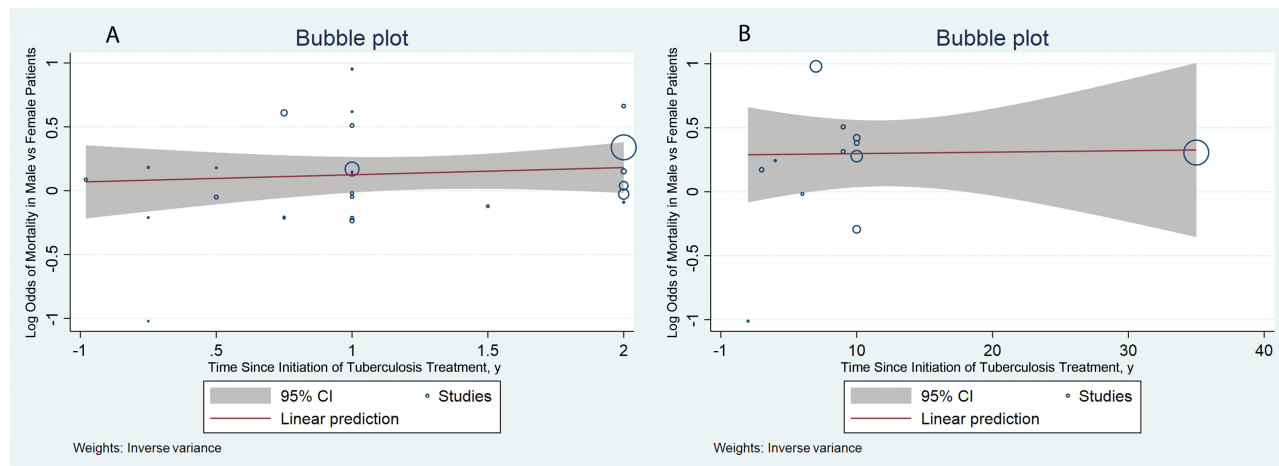
**Table 4. Subgroup Analysis of Pooled Effect Sizes for the Association of Male Sex With All-Cause Mortality Among Patients With Tuberculosis**

Subgroup Characteristics	Pooled ORs <sup>a</sup>			Pooled HRs <sup>a</sup>				
	No. of Studies	Unadjusted OR (95% CI)	$\hat{\rho}$ Statistic	$P$ Value <sup>b</sup>	No. of Studies	Unadjusted HR (95% CI)	$\hat{\rho}$ Statistic	$P$ Value <sup>b</sup>
<b>Income status of country</b>								
Low	18	1.19 (1.03–1.38)	27.6	.06	7	1.17 (1.02–1.35)	44.7	<b>.02</b>
Middle	60	1.19 (1.09–1.32)	75.4		21	1.06 (.94–1.19)	55.9	
High	41	1.39 (1.26–1.52)	80.1		13	1.34 (1.20–1.56)	48.3	
<b>Tuberculosis burden</b>								
High	65	1.19 (1.09–1.30)	74.2	<b>.03</b>	23	1.08 (.98–1.20)	57.1	<b>.031</b>
Low	54	1.37 (1.26–1.49)	74.9		18	1.30 (1.14–1.49)		
<b>HIV-tuberculosis burden</b>								
High	66	1.19 (1.09–1.36)	73.9	<b>.03</b>	24	1.10 (.99–1.21)	58.2	.08
Low	53	1.37 (1.25–1.50)	75.3		17	1.28 (1.12–1.47)	56.8	
<b>Continent</b>								
Africa	44	1.19 (1.08–1.30)	43.1	<b>&lt;.001</b>	17	1.14 (1.03–1.25)	48.9	<b>.004</b>
Asia	48	1.32 (1.19–1.47)	76.9		17	1.19 (1.01–1.41)	68.5	
Europe	11	1.41 (1.35–1.47)	0		2	2.35 (1.04–5.28)	22.5	
North America	9	1.16 (.96–1.40)	23.6		4	1.38 (1.17–1.62)	0	
South America	6	0.97 (.94–1.01)	0		3	0.78 (.61–1.00)	0	
Australia/Pacific	2	2.13 (1.72–2.63)	0		...	...	...	
<b>Drug resistance pattern of tuberculosis</b>								
Drug sensitive	24	1.31 (1.13–1.52)	62.6	.88	1	1.75 (1.43–2.15)	0	.05
Drug resistant	23	1.18 (1.03–1.34)	46.5		9	1.23 (1.04–1.61)	10	
Multidrug resistant	11	1.26 (1.01–1.57)	50.2		6	1.23 (.98–1.62)	15	
Extensively drug resistant	1	1.19 (.64–2.21)	...		1	0.86 (.54–1.37)	...	
<b>Tuberculosis-HIV coinfection</b>								
HIV negative	4	1.55 (1.12–2.16)	32.15	<b>.03</b>	3	1.53 (1.22–1.90)	24.4	<b>.005</b>
HIV positive	21	0.97 (.94–1.01)	42.98		14	1.04 (.89–1.21)	44.0	
<b>Tuberculosis site</b>								
Pulmonary	29	1.36 (1.16–1.59)	76.4	.29	12	1.29 (1.07–1.56)	68.7	.12
Extrapulmonary	11	1.13 (.94–1.36)	0		2	1.00 (.77–1.30)	0	
<b>DOTS implementation in study country</b>								
Implemented	111	1.33 (1.15–1.43)	74.9	.09	35	1.15 (1.07–1.26)	58.7	.60
Not implemented	19	1.47 (1.21–1.60)	66.9		5	1.25 (.93–1.68)	73.2	

Abbreviations: CI, confidence interval; DOTS, directly observed treatment, short-course; HIV, human immunodeficiency virus; HR, hazard ratio; OR, odds ratio.

<sup>a</sup>Studies reporting outcomes in intensive care unit settings were excluded from the subgroup analyses.

<sup>b</sup> $P$  values pertain to the difference in the ORs and HRs for mortality in male patients among the subgroups.



**Figure 3.** Bubble plots showing the association between log odds of mortality in male compared with female patients and the time of assessment of mortality, during tuberculosis treatment (A) and during follow-up (B). Abbreviation: CI, confidence interval.

**Table 5. Association of Male Sex With Sputum Smear and Sputum Culture Positivity Among Patients With Tuberculosis, Using Random Effects Meta-analysis**

Outcome and Time of Assessment	Unadjusted ORs				Adjusted ORs			
	No. of studies	Effect Size (95% CI)	$I^2$ Statistic	<i>P</i> Value <sup>a</sup>	No. of studies	Effect Size (95% CI)	$I^2$ Statistic	<i>P</i> Value <sup>a</sup>
<b>Sputum culture for AFB</b>								
All studies	13	1.60 (1.28–1.99)	24.6	...	4	1.81 (1.39–2.34)	0	...
End of intensive phase	13	1.60 (1.28–1.99)	24.6		4	1.81 (1.39–2.34)	0	
End of treatment	...	...	...		...	...	...	
<b>Sputum smear for AFB</b>								
All studies	29	1.40 (1.24–1.58)	21.06	.06	6	1.45 (1.31–1.61)	0	.48
End of intensive phase	6	1.58 (1.41–1.77)	0		3	1.42 (1.26–1.60)	0	
End of treatment	22	1.23 (1.02–1.48)	28.27		3	1.51 (1.16–1.98)	26.06	

Abbreviations: AFB, acid-fast bacilli; CI, confidence interval; OR, odds ratio.

<sup>a</sup>*P* values pertain to the difference in the ORs for male patients between the subgroups (end of intensive phase and end of treatment).

Despite evidence of earlier diagnostic testing and treatment initiation in male compared with female patients [7, 28], male patients have significantly higher rates of hemoptysis, abnormal chest radiographs, and more severe lung damage at computed tomography and higher odds of tuberculosis-related and all-cause mortality [7, 28]. In addition, compared with women, men show delayed radiological response with computed tomography after ATT [7]. These observations are consistent with the hypothesis that men exhibit more severe tuberculosis disease at presentation and have worse outcomes despite better healthcare access and prompt initiation of treatment.

In our cohort, men had disproportionately high rates of smoking and cavitory disease than women. Prior studies have shown higher proportions of cavitory disease in men compared with women with similar smoking histories [6, 29], which may be explained by increased expression of matrix metalloproteinase in the lungs in men [30]. In our cohort, men had higher odds of 2-month sputum culture positivity, after adjustment for smoking and presence of cavitory disease at baseline. Our systematic review also showed higher pooled unadjusted and adjusted odds of sputum AFB smear and culture positivity in male patients after tuberculosis treatment initiation. These findings highlight the potential biological role of male sex in unfavorable microbiological outcomes.

Male sex has been shown to predispose to other infectious diseases [31–36]. Although tuberculosis risk factors causing immune dysregulation, such as HIV [37] and malnutrition [38], have been well characterized, sex as a risk factor for tuberculosis disease has been understudied. Studies have shown more severe disease in male mice after infection with *M. tuberculosis* and nontuberculous mycobacteria, manifested by higher lung bacillary burdens and increased mortality rates [8, 9, 39]. However, the molecular basis for these phenotypic differences remains to be elucidated.

Recently, sex hormones were shown to differentially regulate female and male immune responses to various pathogens [40]. Testosterone was shown to reduce antimicrobial activity and promote mycobacterial growth [41, 42], reduce the expression of Toll-like receptor (TLR) 4 in mouse macrophages [43] and to have immunosuppressive effects by increasing inhibitory cytokines, reducing immunoglobulin production, and inhibiting T-cell and B-cell maturation [44–47]. Androgens are also known to induce an interleukin 4–driven M2 alveolar macrophage differentiation [48]. Signaling through the androgen receptor can increase monocyte recruitment by CCL2 and CXCL1 [49], potentially driving more extensive cavitory disease in men. In contrast, estradiol has been shown to enhance macrophage activation [50]. Sex-based differences in baseline activation of mammalian target of rapamycin complex have been reported [51, 52], and dysregulation of this pathway may be postulated to result in up-regulation of lipid accumulation [53], inhibition of autophagy [54], and increased cell necrosis, leading to a proinflammatory environment and increased cavitation in the lungs of men with tuberculosis. The human X chromosome-linked genes that encode effectors in immune responses, including TLR7, TLR8, interleukin 1 receptor–associated kinase, CD40 ligand, and Forkhead box protein P3 (FOXP3), have been implicated in sex differences in immune responses and tissue damage after infection [55]. It is possible that X-linked genes in females escape silencing, leading to higher expression of host-protective immune-related genes [56].

In our meta-analysis, male patients had a higher pooled OR for mortality among studies including exclusively HIV-negative patients with tuberculosis than in studies including patients with HIV-tuberculosis coinfection. Prior literature has shown that among HIV-negative patients, men had worse outcomes than women, while among tuberculosis-HIV-coinfected patients, women had worse outcomes [57, 58]. Thus, HIV coinfection appears to reduce the potential

immunomodulatory protective effect of female sex on tuberculosis treatment outcomes.

Our study has several strengths. The large sample size of our cohort aided in the analysis of various confounding factors. Because the cohort data were obtained from a single institution, heterogeneity in treatment decisions were minimized. Our systematic review of studies from 81 countries with variable healthcare access, allowed for generalizability of our results. The large number of studies on all-cause mortality in our systematic review enabled us to perform subgroup analyses and to arrive at various inferences.

Several limitations are noted. We were unable to assess parameters such as socioeconomic status, occupational history, and delay in diagnosis in our cohort. We did not have access to additional radiographic information, such as the volume of lung affected as a measure of severity for comparison between the sexes. In our systematic review, antibiotic susceptibilities, study setting, and timing of outcome assessment were not reported consistently among the included studies. Several studies in our review, performed from program registries, did not report data on baseline characteristics stratified by sex. During meta-analysis of adjusted effect sizes, individual studies were adjusted for comparable but different parameters. The treatment regimens followed may not be uniform across the included studies.

Our results have several important therapeutic implications. Recognizing the association between male sex and adverse treatment outcomes should inform medical decision making in tuberculosis treatment programs and motivate further focused research into the biological (immunological and genetic) basis of these sex-based differences, and, in turn, novel host-directed therapies to improve clinical, pathological, and microbiological outcomes during tuberculosis treatment in both sexes.

### Supplementary Data

Supplementary materials are available at *Clinical Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

### Notes

**Author contributions.** V. C. and P. C. K. conceived the idea for the review. V. C., N. L. T., M. G. M., J. B., and P. C. K. designed and undertook the literature review. V. C., N. L. T., M. G. M., S. K. A., A. K., P. N., E. P. W., and E. J. A. screened the articles and extracted the data (with help from S. W. and A. Z.). V. C., N. L. T., and M. G. M. performed the statistical analysis and prepared the figures and appendix. V. C., M. G. M., A. K., and R. K. S. analyzed and interpreted the data. V. C., M. G. M., J. R. C., A. K., R. K. S., and P. C. K. wrote the first draft of the manuscript, and V. C., A. G., J. Y. W., and P. C. K. revised the subsequent drafts. All authors reviewed the final draft of the manuscript.

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