# MAJOR ARTICLE







# Glycemic Control and the Prevalence of Tuberculosis Infection: A Population-based Observational Study

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**Background.** Several cohort studies demonstrate that diabetics are at increased risk for active tuberculosis, and poor glycemic control may exacerbate this risk. A higher prevalence of tuberculosis infection at baseline among diabetics may partially explain these results; however, no population-based studies have investigated this association. Furthermore, whether glycemic control modifies the relationship between diabetes and tuberculosis infection, as it does with active tuberculosis, is unknown.

**Methods.** Diabetics were diagnosed through physician evaluation and using 3 laboratory tests including hemoglobin  $A_{\rm IC}$  (HbA $_{\rm IC}$ ), fasting plasma glucose (FPG), or 2-hour plasma glucose (PG). Tuberculosis infection was diagnosed through tuberculin skin tests, and glycemic control was assessed linearly and categorically using recommended targets.

**Results.** Among 4215 participants, the prevalence of tuberculosis infection was 4.1%, 5.5%, and 7.6% in nondiabetic, prediabetic, and diabetic participants ( $P_{\rm trend} = .012$ ). In multivariate analysis, diabetes was associated with tuberculosis infection (adjusted odds ratio [AOR], 1.5; 95% confidence interval [CI], 1.0–2.2). Compared to nondiabetics, diabetics who were undiagnosed (AOR, 2.2 and 1.2 in diagnosed diabetics), FPG >130 mg/dL (AOR, 2.6 and 1.3 in diabetics with FPG ≤130 mg/dL), or not on insulin (AOR, 1.7 and 0.8 in diabetics on insulin) had elevated tuberculosis infection rates. In a linear dose–response analysis, increasing values of FPG (AOR, 1.02 per 1-mg/dL; 95% CI, 1.01–1.04), and HbA $_{\rm 1C}$  (AOR, 1.13 per 1%; 95% CI, 1.04–1.22) all predicted tuberculosis infection.

**Conclusions.** Our results suggest glycemic control may modify the relationship between tuberculosis infection and diabetes. **Keywords.** *Mycobacterium tuberculosis*; diabetes; glycemic control; tuberculosis.

The World Health Organization has set an ambitious target to decrease the incidence of active tuberculosis globally by 90% from 2015 to 2035 [1]. Currently, the incidence of tuberculosis is decreasing at only 1%–2% per year [2]. Individuals with diabetes have a high risk of active tuberculosis, and this has the potential to substantially impact future tuberculosis control efforts [3, 4] because the global prevalence of diabetes is increasing, especially in low-income settings [5] where tuberculosis is endemic. Furthermore, almost half of diabetics are undiagnosed globally, suggesting that this comorbidity's impact on tuberculosis rates may be substantially higher than is currently known [6]. Despite these ominous signs, tuberculosis control policy that deals with this coepidemic has been hampered by a lack of evidenced-based

research, and only conditional recommendations, pending more evidence, have been given by global health organizations [7–9]. A further understanding of the mechanisms that allow diabetes and tuberculosis to interact is necessary in order to develop innovative interventions for preventing and controlling tuberculosis in areas where both diseases are endemic [7, 9].

Several high-quality cohort studies have shown that diabetics are at an increased risk of developing active tuberculosis compared to the general population and that those with poor diabetic control may be especially vulnerable [10-13]. One explanation for these results is that diabetics are more likely, once infected, to develop primary progressive disease or to reactivate an old infection. Alternatively, diabetics in these studies may have a higher underlying prevalence of tuberculosis infection at baseline compared to the general population. However, since tuberculosis infection was not measured in these studies, the cause of this elevated risk is unclear and has not been thoroughly investigated [14]. There have been no population-based studies to investigate whether diabetics have increased susceptibility to tuberculosis infection [15]. Furthermore, whether glycemic control modifies the relationship between diabetes and tuberculosis infection, as it does with active tuberculosis, is unknown [15].

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To attempt to inform this knowledge gap we had 2 objectives. First, we investigated whether diabetics were at increased risk for tuberculosis infection in a large, population-based cohort. Second, we examined whether glycemic control modified this relationship.

## **METHODS**

## **Data Collection and Study Design**

The National Health and Nutrition Examination Survey (NHANES) conducts complex multistage probability surveys among a representative sample of the noninstitutionalized civilian population in the United States. We used NHANES data from 2011 through 2012 for all analyses. Data were collected from participants through standardized questionnaires, biological samples, and physical examinations. Participants were interviewed in their homes to ascertain demographic characteristics and at a mobile examination center to ascertain possible risks factors and exposures for various diseases. All participants provided written informed consent for this evaluation.

We included demographic and clinical characteristics from various questionnaires as well as information from tuberculosis and diabetes surveys. Tuberculosis infection and diabetes were evaluated through several laboratory assessments. In 2011–2012, NHANES measured tuberculosis infection using 2 tests: the tuberculin skin test (TST) and the QuantiFERON Gold In-Tube test (QFT-GIT). The TST was administered on the forearm using purified protein derivative (Tubersol, Sanofi, Bridgewater, New Jersey). Trained health workers measured the TST induration reaction 46–76 hours after administration. The QFT-GIT was administered on the same day as the TST and was performed following manufacturer guidelines.

Several diabetes diagnostic tests were used. During the home visit, participants reported whether they had a physician-based diabetes diagnosis. During laboratory testing, hemoglobin  $A_{\rm 1C}$  (HbA $_{\rm 1c}$ ) was administered and measured on all eligible participants. Fasting plasma glucose (FPG) and 2-hour plasma glucose (PG) tests were performed on all participants examined in a morning session after a 9-hour fast. Participants were randomly assigned to a morning examination session (or an afternoon or evening session); therefore, only a random subset of eligible participants took these 2 tests. The oral glucose tolerance test was administered using a calibrated dose of 75 g of glucose and a venipuncture 2 hours later.

## **Definitions**

## **Diabetes and Prediabetes**

We defined diabetes through criteria given by the American Diabetes Association [16] and the World Health Organization [17, 18] and using methods from recent diabetes prevalence surveys [19, 20]. During the home visit, participants self-reported whether they had a physician-based diabetes diagnosis. Participants who responded positively to this question

were classified as diagnosed diabetes; undiagnosed diabetes was defined as participants reporting never previously diagnosed with diabetes but with 1 of the following test results:  $HbA_{1C} \ge 6.5\%$ ,  $FPG \ge 126$  mg/dL, or  $PG \ge 200$  mg/dL.

Using the same guidelines, we performed a similar process to diagnose prediabetes for all individuals who did not meet the diabetes criteria listed above. Prediabetics had 1 of the following: previously told by a physician they had prediabetes,  $HbA_{1C}$  5.7%–6.4%, FPG 100–125 mg/dL, or PG 140–199 mg/dL.

## Glycemic Control and Diabetes Severity

Glycemic control was assessed based on glycemic targets for diabetic patients set by the American Diabetes Association in 2016 [21]. Cutoffs used to classify poor glycemic control were HbA $_{1c} \ge 7\%$  and FPG >130 mg/dL. We also examined the doseresponse relationship between tuberculosis infection and each of the 3 laboratory diagnostic methods (HbA $_{1c}$ , FPG, PG) linearly. We fit a logistic regression model testing the probability of tuberculosis infection for every 1-mg/dL increase in FPG or PG and for every 1% increase of HbA $_{1c}$ .

Diabetes severity was evaluated using answers from several sociodemographic questionnaires administered by NHANES. Survey-based methods used as proxies for disease severity included undiagnosed diabetes, time since diabetes diagnosis, current insulin use, and use of diabetic pills to lower blood sugar.

## **Tuberculosis Infection**

In NHANES 2011–2012, both TST and QFT-GIT were administered to participants in order to assess tuberculosis infection. The sensitivity of the QFT-GIT in diabetic patients is debated, and studies have been inconsistent [22–24]. Because of this, we used the TST in our primary analysis to diagnose tuberculosis infection. Tuberculosis infection was defined as an induration  $\geq$ 10 mm, as seen in prior NHANES studies [25, 26]. To assess the robustness of our results, we conducted a sensitivity analyses using the QFT-GIT. To provide proper validation, we included only participants who took both the TST and QFT-GIT in all analyses. A QFT-GIT was considered positive when the tuberculosis antigen minus nil value was  $\geq$ 0.35 IU/mL, with the tuberculosis response  $\geq$ 25% of the nil value and the nil value  $\leq$ 8.0 IU/mL, per guidelines [27].

## **Statistical Analytical Plan**

Statistical weights were used in all prevalence and modeling analyses to account for unequal probabilities of selection and nonresponse and thus provide estimates representative of the civilian, noninstitutionalized US population. The weighted prevalence of tuberculosis infection was estimated using standard 2  $\times$  2 contingency tables and was stratified by demographic and clinical participant characteristics. Proportions were compared with Pearson  $\chi^2$  statistics with a Rao and Scott

second-order correction using an F statistic with noninteger degrees of freedom, accounting for the survey design [28, 29]. A binary logistic regression model was used to calculate odds ratios (ORs), and these were compared by demographic and risk factors in univariate and multivariate analyses. Variables suggestive of an association with tuberculosis infection (P < .2) and those shown to have an association in prior studies were included in multivariate analyses. Cochran-Armitage tests were used to test for trends between multiple categories in 1 variable.

We calculated the prevalence of tuberculosis infection for each measurement method of glycemic control and disease severity and then compared the corresponding prevalence of tuberculosis infection in nondiabetic participants. Univariate and multivariate logistic regression models were then constructed to determine whether effect modification was present. Statistical significance was assessed using 95% confidence intervals (CIs) in all models. Data were analyzed using Stata version 14.1 statistical software.

#### **RESULTS**

## **Study Population and Diabetes Prevalence**

Participants aged  $\geq$ 20 years were included (n = 5560). We excluded participants who only took the TST (n = 28), only took the QFT-GIT (n = 747), or did not take either test (n = 552).

We also excluded participants with an indeterminate QFT-GIT (n = 18). This left 4215 participants for the final analysis (Figure 1).

The prevalence of diagnosed diabetes in our study population was 9.2% (95% CI, 7.9–10.7; N = 535), 4.0% (95% CI, 3.3–4.9; N = 241) for undiagnosed diabetes, and 33.7% (95% CI, 31.0–36.4; N = 1441) for prediabetes (Table 1). Among diabetic patients, 30.4% (95% CI, 25.6–35.6) were undiagnosed. Diabetics were more likely to be older, have a lower education level, and have a high body mass index (P < .01). There was a statistically increasing trend between nondiabetic, prediabetic, and diabetic patients in regard to their median PG, FPG, and HbA<sub>1c</sub> (all  $P_{\rm trend} < .01$ ).

#### **Prevalence of Tuberculosis Infection**

The overall weighted prevalence of tuberculosis infection was 5.0% (95% CI, 3.5–7.1; N = 404 infected; Table 2). Tuberculosis infection prevalence was elevated in individuals who did not complete high school (11.3% infected), individuals who were in household contact with a tuberculosis case (14.6% infected), or individuals who were in a household with 6 or more people (10.4% infected). Tuberculosis infection prevalence was 4.1%, 5.5%, and 7.6% in nondiabetic, prediabetic, and diabetic participants ( $P_{\rm trend}$  = .012). When we stratified by the diagnostic method used to detect diabetes (Figure 2), there was an

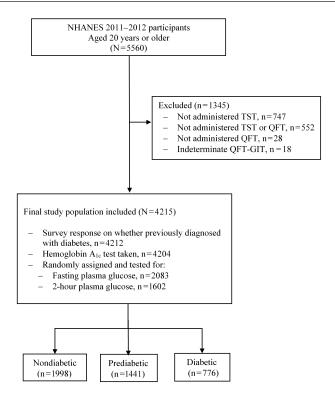


Figure 1. Eligibility and enrollment of included participants. Abbreviations: NHANES, National Health and Nutrition Examination Survey; QFT-GIT, QuantiFERON Gold In-Tube test TST, tuberculin skin test.

increasing prevalence of tuberculosis infection from nondiabetic, prediabetic, and diabetic patients for FPG ( $P_{\rm trend}$  = .0265), PG ( $P_{\rm trend}$  = .0009), and HbA $_{\rm lc}$  ( $P_{\rm trend}$  = .022) tests but not for self-reported diabetes ( $P_{\rm trend}$  = .5958).

In a multivariate analysis controlling for age, sex, birthplace, household tuberculosis contact, cigarette smoking status, and family size, diabetes was significantly associated with tuberculosis infection (adjusted OR [AOR], 1.5; 95% CI, 1.0–2.2; Table 3).

Table 1. Demographic and Clinical Characteristics of Adults Aged ≥ 20 Years in the United States, 2011–2012, Stratified by Diabetes Status

Variable	Nondiabetic, % (n)	Prediabetic, % (n)	Diabetic, % (n)	All Participants, % (n)	P Value <sup>a</sup>
N	53.1 (1998)	33.7 (1441)	13.2 (776)	100 (4215)	
Demographic characteristics					
Age group, y					
20–29	27.6 (586)	9.4 (141)	1.7 (16)	18.0 (743)	<.0001
30–39	21.7 (463)	14.3 (228)	7.4 (49)	17.3 (740)	
40–49	20.1 (343)	20.1 (258)	16.0 (107)	19.6 (708)	
≥50	30.6 (606)	56.2 (814)	74.9 (604)	45.1 (2024)	
Sex					
Male	44.1 (904)	52.4 (777)	49.8 (394)	47.6 (2075)	.002
Female	55.9 (1094)	47.6 (664)	50.1 (382)	52.4 (2140)	
Birthplace					
United States	83.4 (1437)	84.4 (1029)	79.9 (528)	83.3 (2994)	.124
Foreign	16.6 (560)	15.7 (411)	20.1 (248)	16.7 (1219)	
Education					
Beyond high school	69.6 (1258)	60.2 (739)	46.6 (327)	63.4 (2324)	<.0001
High school	17.7 (378)	22.1 (350)	26.0 (177)	20.3 (905)	
Less than high school	12.7 (362)	17.7 (351)	27.4 (271)	16.3 (984)	
Missing					
Household tuberculosis contact					
Yes	2.1 (54)	3.5 (56)	3.4 (35)	2.8 (145)	.1168
No	97.9 (1933)	96.5 (1374)	96.6 (738)	97.3 (4045)	
Smoking					
Current	17.9 (392)	23.1 (326)	17.4 (133)	19.6 (851)	.0067
Former	22.6 (375)	24.2 (354)	32.3 (248)	24.4 (977)	
Never	59.5 (1231)	52.3 (761)	50.3 (395)	55.6 (2387)	
Family size					
Fewer than 3	51.2 (943)	54.7 (714)	60.1 (438)	53.5 (2095)	.045
3–5	42.7 (871)	38.9 (604)	32.9 (259)	40.1 (1734)	
6 or more	6.2 (184)	6.5 (123)	7.0 (79)	6.4 (386)	
Clinical characteristics					
Median BMI (IQR), kg/m <sup>2</sup>	26.3 (23.0–30.2)	28.6 (24.9–33.1)	31.1 (26.8–36.4)	27.8 (24.1–32.5)	<.0001
BMI, kg/m <sup>2</sup>					
<18.5	2.3 (57)	0.9 (18)	0.6 (3)	1.6 (78)	<.0001
18.5–24.9	35.5 (745)	22.5 (340)	12.6 (113)	28.1 (1198)	
25.0–29.9	35.6 (649)	35.3 (488)	25.1 (216)	34.2 (1353)	
≥30.0	26.6 (523)	41.2 (583)	61.8 (431)	36.1 (1537)	
Median 2-hour plasma glucose (IQR), mg/dL <sup>b</sup>	94 (78–108)	117 (96–141)	223 (201–261)	108 (88–135)	<.0001
Median fasting plasma glucose (IQR), mg/dL <sup>b</sup>	91.5 (87–95)	103 (99–108)	129 (113–163)	100 (92–110)	<.0001
Median hemoglobin A <sub>1c</sub> (IQR), % <sup>b</sup>	5.3 (5.1–5.5)	5.8 (5.6–6.0)	6.7 (6.2–7.8)	5.5 (5.2–5.9)	<.0001
Hepatitis B, surface antigen					
Positive	0.3 (14)	0.4 (14)	0.2 (4)	0.3 (32)	.5246
Negative	99.7 (1965)	99.6 (1414)	99.8 (755)	99.7 (4134)	
Hepatitis C, antibody					
Positive	1.1 (29)	1.9 (28)	3.4 (24)	1.7 (18)	.168
Negative	98.9 (1943)	98.1 (1395)	96.6 (734)	98.3 (4072)	

Abbreviations: BMI, body mass index; IQR, interquartile range.

Data are weighted percentages of participants in each subsample. The absolute number of participants is shown in parentheses. The absolute number of participants does not perfectly correspond to percentages because the percentages are weighted, adjusting for the survey design. Percentages may not total 100% because within column percentages were rounded to the nearest integer. Column totals vary across different characteristics because of missing values for some participants.

<sup>&</sup>lt;sup>a</sup>P value is an F statistic with noninteger degrees of freedom by using a second-order Rao and Scott correction, taking into account the survey design. This P value is based on Pearson  $\chi^2$  statistic for 2-way contingency tables.

<sup>&</sup>lt;sup>b</sup>Not every participant took all 3 tests.

Tuberculosis infection was also associated with current smoking (AOR, 1.7; 95% CI, 1.0–2.8 compared to never smokers), household tuberculosis contact (AOR, 3.2; 95% CI, 1.5–6.9), foreign-born status (AOR, 14.9; 95% CI, 8.5–26.3 compared to US born), a family size  $\geq$ 6 (AOR, 1.8; 95% CI, 1.0–3.4 compared to <3 family members), and being aged  $\geq$ 50 years (AOR, 1.7; 95% CI, 1.0–2.9 compared to 20–29 years). In a separate multivariate model, adding time since arrival in the United States, the association seen among tuberculosis infection and diabetes remained unchanged (AOR, 1.5; 95% CI, 1.0–2.3).

#### **Diabetes Severity and Tuberculosis Infection**

When we stratified diabetes by several distinct markers of glycemic and diabetic control (Figure 3), some characteristics modified the relationship between diabetes and tuberculosis infection. In both univariate and multivariate analyses, increasing values for HbA<sub>1c</sub> (AOR, 1.13 per 1% increase in HbA<sub>1c</sub>; 95% CI, 1.04-1.22), FPG (AOR, 1.02 per 1-mg/dL increase in FPG; 95% CI, 1.01-1.03), and PG test (AOR, 1.02 per 1-mg/dL increase in PG; 95% CI, 1.01-1.04) results predicted tuberculosis infection (Figure 3). After categorizing these laboratory tests by glycemic control targets for diabetic patients, diabetic patients with an FPG >130 mg/dL had a higher prevalence of tuberculosis infection compared to diabetics with an FPG ≤130 mg/dL and nondiabetics (9.9% vs 6.5% and 4.1%). In a multivariate model, diabetics with an FPG >130 mg/dL continued to have a significantly elevated risk of tuberculosis infection (AOR, 2.6; 95% CI, 1.5-4.6) compared to nondiabetics, while diabetics with an FPG ≤130 mg/dL (AOR, 1.3; 95% CI, 0.8-3.6) were not at increased risk. In univariate analysis, diabetic patients with HbA<sub>1</sub>, test results ≥7% were at an increased tuberculosis infection risk (9.1% prevalence in diabetics with  $HbA_{1c} \ge 7\%$ , 6.7% in diabetics with  $HbA_{1c} < 7\%$ , and 4.1% in nondiabetics;  $P_{\text{trend}} = .003$ ). However, these differences were no longer present in a multivariate model (compared to nondiabetic reference; AOR, 1.7, 95% CI, 0.9-3.0 for diabetics with  $HbA_{1c} \ge 7\%$  and AOR, 1.5, 95% CI, 0.9–2.5 for diabetics with  $HbA_{1c}$  < 7%).

Among survey-based proxies for diabetic control, high rates of tuberculosis infection were seen in diabetics who were undiagnosed (12.9% vs 5.4% in diagnosed cases and 4.1% in nondiabetic individuals) or were not currently taking insulin (8.8% prevalence vs 3.3% in diabetics currently taking insulin and 4.1% in nondiabetic individuals). In a multivariate model (Figure 3), undiagnosed diabetics (AOR, 2.2; 95% CI, 1.3–3.6) and diabetic patients not currently taking insulin (AOR, 1.7; 95% CI, 1.2–2.6) remained associated with tuberculosis infection, but diagnosed diabetics (AOR, 1.2; 95% CI, 0.7–2.0) and those taking insulin (AOR, 0.8; 95% CI, 0.3–2.3) were not. Other markers for disease severity, such as the time since diabetes diagnosis or diabetic pill use, did not modify the relationship between diabetes and tuberculosis infection.

## **Sensitivity Analysis**

When we used the QFT-GIT, the prevalence of tuberculosis infection in the study population was 5.6% (95% CI, 4.6–6.7). Test agreement and kappa values were 90.6% and 0.46, respectively. The prevalence of tuberculosis infection was 4.4% (95% CI, 3.5–5.4), 5.5% (95% CI, 4.2–7.3), and 10.5% (95% CI, 7.4–10.5) in nondiabetic, prediabetic, and diabetic participants ( $P_{\rm trend}=.0001$ ; Supplementary Materials Table E1). After multivariate adjustment, diabetics were 1.7 times (95% CI, 1.0–2.9) more likely to have tuberculosis infection compared to nondiabetic participants (Supplementary Materials Table E2). When we stratified by the diagnostic method used to detect diabetes (Supplementary Materials Figure E1), there was an increasing prevalence of tuberculosis infection from nondiabetic, prediabetic, and diabetic patients for PG ( $P_{\rm trend}=.0088$ ), HbA $_{\rm 1c}$  ( $P_{\rm trend}=.0003$ ), and self-report ( $P_{\rm trend}=.0037$ ), as well as a suggestive trend for FPG ( $P_{\rm trend}=.0987$ ).

## **DISCUSSION**

In this population-based sample with more than 4000 participants and 700 diabetics, we found that several biomarkers, both categorical and linear, of glycemic control modified the relationship between tuberculosis infection and diabetes. Several cohort studies have demonstrated that diabetics with poor diabetic control have especially high risk for active tuberculosis [10, 12, 13]. Our results suggest these diabetics may also have increased susceptibility to tuberculosis infection.

In 3 important studies, poorly controlled diabetes (measured through HbA<sub>1c</sub> ≥7% [12], FPG >130 mg/dL [13], and diabetes complications [10]) was associated with active tuberculosis in diabetics. To our knowledge, we are the first to investigate whether glycemic control also modifies the relationship between diabetes and tuberculosis infection. We found that diabetics with poorly controlled disease (as measured by an FPG >130 mg/dL, undiagnosed diabetes, or those not using insulin) had especially high rates of tuberculosis infection. We also found the odds of tuberculosis infection significantly increased for every 1-mg/ dL increase in FPG or PG and for every 1% increase in HbA1. This corresponds well with results from a recent large study from Taiwan that showed an increasing hazard for active tuberculosis for every 10-mg/dL increase in FPG (AOR, 1.1 [95% CI, 1.0-1.1]) [13]. We suggest part of the risk for active tuberculosis among diabetics seen in many studies [11] may relate to a higher rate for tuberculosis infection, especially among those with poorly controlled glucose levels. However, our study was not longitudinal and we did not measure active tuberculosis. Therefore, we cannot state the contributory role of primary progressive disease in this process. Future cohort studies estimating the risk of active tuberculosis among diabetic patients should measure and control for tuberculosis infection at baseline.

In a recent metaanalysis in which AORs from 13 observational studies were pooled, a slightly higher, but significant, rate of

Table 2. Risk Factors for Tuberculosis Infection Among Adults Aged ≥ 20 Years in the United States, 2011–2012

Variable	Total N <sup>a</sup>	Prevalence of Tuberculosis Infection (95% CI) <sup>a</sup>	Crude Odds Ratio (95% CI)ª
N	4215	5.0 (3.5–7.1)	_
Demographic characteristics	1210	0.0 (0.0 7.1)	
Age group, y			
20–29	406	3.4 (1.8–6.5)	1 (Referent)
30–39	1430	5.2 (3.3–8.2)	1.7 (1.0–3.1)
40–49	1470	5.8 (4.0–8.3)	1.6 (1.1–2.3)
≥50	909	3.8 (2.5–5.7)	1.5 (0.9–2.4)
Sex	000	0.0 (2.0 0.7)	(0.0 2.1)
Male	2075	5.4 (3.7–7.9)	1 (Referent)
Female	2140	4.6 (3.3–6.5)	0.8 (0.7–1.08)
Birthplace		(2.12 2.12)	
United States	2994	1.8 (1.0–3.1)	1 (Referent)
Foreign	1219	21.2 (15.8–27.7)	
Education			(0.0,
Beyond high school	2324	3.5 (2.6–4.8)	1 (Referent)
High school	905	4.6 (2.9–7.4)	1.3 (0.9–2.0)
Less than high school	984	11.3 (7.4–16.9)	3.5 (2.5–4.9)
Household tuberculosis contact			
No	4045	4.7 (3.3–6.6)	1 (Referent)
Yes	145	14.6 (8.2–24.6)	3.5 (1.9–6.2)
Cigarette smoking status		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	,
Never	2387	5.0 (3.4–7.2)	1 (Referent)
Former	977	4.8 (3.3–7.0)	1.0 (0.6–1.5)
Current	851	5.4 (3.1–9.3)	1.1 (0.7–1.8)
Family size			
Fewer than 3	2095	3.2 (2.1-4.8)	1 (Referent)
3–5	1734	6.6 (4.2–10.4)	2.2 (1.3–3.8)
6 or more	386	10.4 (6.5–16.1)	3.6 (1.9–6.8)
Clinical characteristics			
2-hour plasma glucose, mg/dLb	1602	-	1.0 (1.0-1.1)
Fasting plasma glucose, mg/dL <sup>b</sup>	2083	-	1.0 (1.0–1.0)
Hemoglobin A <sub>1c</sub> , % <sup>b</sup>	4202	_	1.2 (1.1-1.3)
Diabetes status			
Nondiabetic	1998	4.1 (2.9-5.6)	1 (Referent)
Prediabetic	1441	5.5 (3.2-9.3)	1.4 (0.9–2.1)
Diabetic	776	7.6 (5.8–10.0)	2.0 (1.5–2.6)
BMI, kg/m <sup>2</sup> (continuous)	4215	-	0.99 (0.97–1.01
BMI, kg/m <sup>2</sup>			
<18.5	78	6.4 (2.3–16.8)	1.2 (0.4–3.8)
18.5–24.9	1198	5.3 (3.5–7.8)	1 (Referent)
25.0–29.9	1353	4.8 (3.4–6.8)	0.9 (0.7–1.3)
≥30	1537	5.1 (3.2–7.9)	1.0 (0.6–1.5)
Hepatitis B, surface antigen			
Negative	4134	5.0 (3.5–7.0)	1 (Referent)
Positive	32	21.3 (7.5–47.2)	5.2 (1.4–18.9)
Hepatitis C, antibody			
Negative	4072	5.0 (3.5–7.0)	1 (Referent)
Positive	81	8.9 (2.4–27.5)	1.9 (0.5-7.2)

Abbreviations: BMI, body mass index; CI, confidence interval

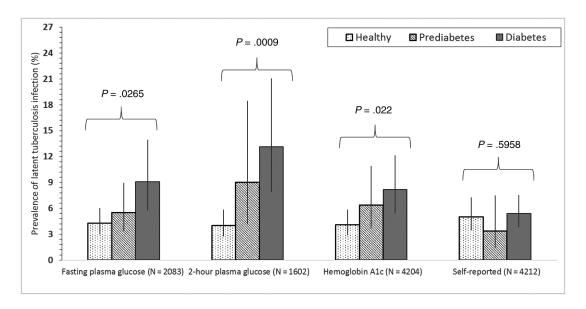
The tuberculin skin test was used to measure tuberculosis infection; a positive test was defined as a skin induration reaction ≥10 mm 46–76 hours after application of the test. A separate analysis was conducted using the QuantiFERON Gold In-Tube test and is included in Supplementary Materials Table E1.

tuberculosis infection among diabetics was found [15]. However, only 2 included studies showed a statistically higher rate of tuberculosis infection among diabetics. There may be several reasons for the differing results from our study. First, since NHANES performs comprehensive laboratory testing, we identified diabetics using physician diagnosis, HbA<sub>1C</sub>, PG, and FPG tests. Past studies identified diabetics through medical records or self-report [15]. Only 1 study [30] used a combination of self-report and laboratory testing to diagnose diabetics. These diagnostic definitions may substantially underestimate diabetes [6, 16, 31]. In a recent metaanalysis of more than 300 000 diabetics, the sensitivity of HbA<sub>1C</sub> to detect undiagnosed diabetes was reported to be 52.8% and 32.7% compared to FPG and PG [32]. We found that undiagnosed diabetics had especially high rates of tuberculosis infection, and this may influence previous study results if these patients are misclassified. Second, many of these studies had sample sizes with few overall participants or few diabetics, leading to low statistical power when tuberculosis infection was analyzed. Last, all prior studies were conducted in high-risk populations (tuberculosis contacts, refugees, hemodialysis patients, or patients with human immunodeficiency virus) [15] that are unlikely to be representative of the general population.

The mechanisms that underlie the association between tuberculosis infection and diabetes seen in this study are unclear and cannot be definitively ascertained from the epidemiological study design used here. Previously, researchers have postulated that this interaction may be influenced by impairment of innate or adaptive immunity through diabetes-induced dysglycemia [30, 33]. In addition, human leukocyte antigen-DQ ß gene alleles (especially at codon 57) may enhance the risk for tuberculous infection and disease progression and confer concurrent risk of diabetes mellitus [34, 35]. Regardless of the cause of this increased risk, policy on screening of tuberculosis infection in diabetic patients should be discussed. Currently, screening of diabetic patients for tuberculosis infection is not recommended by global health organizations [8]. Our study results must be confirmed in further studies, and cost-effectiveness analyses are still needed. However, screening tuberculosis infection in diabetic patients with poor glycemic control may provide an opportunity to effectively target a high-risk population for tuberculosis transmission and primary progressive disease [10, 12, 13] while also limiting expended resources to manageable levels for tuberculosis control programs from low-income settings. Additional research is needed to ascertain the proportion of tuberculosis disease that is attributable to increased

<sup>&</sup>lt;sup>a</sup>Data are weighted prevalence percentages of participants in each subsample and odds ratio (95% confidence interval). The absolute number of participants from each characteristic can be seen on the left hand side.

<sup>&</sup>lt;sup>b</sup>Not all participants took all 3 tests. FPG and PG tests were performed in morning sessions after a 9-hour fast. Participants were randomlyassigned to a morning examination session (or afternoonor evening session); therefore, only a random subset of eligibleparticipants took these 2 tests.



**Figure 2.** Prevalence of tuberculosis infection by the diagnostic method used to diagnose diabetes mellitus. Vertical lines represent 95% confidence intervals. *P* values represent the presence of a trend after accounting for the survey design.

Table 3. Multivariate Logistic Regression Models of Risk Factors for Tuberculosis Infection in the United States, 2011–2012

Variable	Adjusted Odds Ratio (95% Confidence Interval) <sup>a</sup>	<i>P</i> Value <sup>b</sup>	
Age group, y			
20–29	1 (Referent)		
30–39	1.2 (0.6–2.3)	.54	
40–49	1.2 (0.8–1.0)	.31	
≥50	1.7 (1.0–2.9)	.05	
Sex			
Male	1 (Referent)		
Female	0.9 (0.7–1.3)	.85	
Birthplace			
United States	1 (Referent)		
Foreign	14.9 (8.5–26.3)	<.01	
Household tuberculosi	s contact		
No	1 (Referent)		
Yes	3.2 (1.5–6.9)	<.01	
Cigarette smoking stat	tus		
Never	1 (Referent)		
Former	1.2 (0.8–1.9)	.43	
Current	1.7 (1.0–2.8)	.04	
Family size			
Fewer than 3	1 (Referent)		
3–5	1.6 (0.9–2.7)	.07	
6 or more	1.8 (1.0–3.4)	.05	
Diabetes status			
Nondiabetic	1 (Referent)		
Prediabetes	1.3 (0.8–2.1)	.28	
Diabetes	1.5 (1.0–2.2)	.04	

<sup>a</sup>Data are weighted odds ratio (95% confidence interval [CI]) adjusting for the survey design. The model is adjusted for participant age, gender, household contact with a tuberculosis case, birthplace (United States or foreign born), smoking status, family size, and diabetes status. In a separate multivariate model, adding time since arrival in the United States, the association seen among tuberculosis infection, and diabetes remained unchanged (adjusted odds ratio, 1.5; 95% CI, 1.0–2.3). A sensitivity analysis was conducted using the QuantiFERON Gold In-Tube test and is included in Supplementary Materials Table E1.

tuberculosis infection. In a recent study [36], researchers found that compared to tuberculosis contacts, self-reported diabetics had lower prevalence of interferon-gamma positivity but a higher rate of tuberculosis disease, suggesting that although diabetes may be a risk factor for both tuberculosis infection and disease, the increased risk of disease progression may be especially potent.

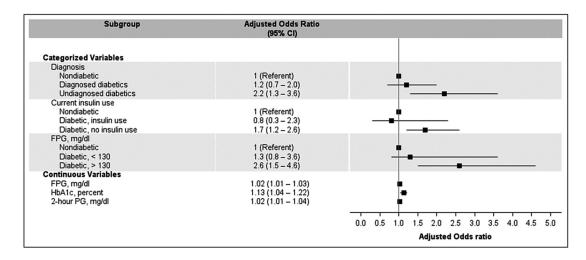
There are limitations to our analyses. First, the prevalence of tuberculosis infection increases with age [25, 37, 38], and we cannot distinguish whether participants acquired tuberculosis infection before or after developing diabetes. Although reverse causality may have occurred in our study, we believe this is unlikely for 2 reasons: (1) tuberculosis infection is unlikely to cause an increased risk of diabetes development; (2) we found a dose-response relationship between tuberculosis infection and differing severities of diabetes. Second, like all observational studies, unmeasured and residual confounding is always a concern. Third, a repeat measurement after a single positive HbA<sub>1,2</sub> FPG, or PG result is recommended to confirm diagnosis [16]. Since NHANES participants only have 1 study visit, this was not possible and diabetes overdiagnosis is possible [19]. However, if present, this bias would bring the association between tuberculosis infection and diabetes toward the null and therefore our estimates may be conservative. Lastly, tuberculosis infection rates were low, limiting our statistical power to assess some markers of diabetes severity.

In conclusion, in a large population-based study, we found that diabetic patients had elevated rates of tuberculosis infection, and diabetics with poor glycemic control modified this relationship.

## **Supplementary Data**

Supplementary materials are available at *Clinical Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted

<sup>&</sup>lt;sup>b</sup>P value is an F statistic with noninteger degrees of freedom by using a second-order Rao and Scott correction, taking into account the survey design.



**Figure 3.** Adjusted logistic regression analysis of tuberculosis infection after stratification by several survey- and laboratory-based proxies of diabetes severity. Data are weighted odds ratios with 95% confidence intervals. Adjusted odds ratios for continuous variables are shown per unit increase (1-mg/dL increase in FPG or PG and for every 1% increase of HbA1C). Diabetic participants were classified as "diagnosed" if they self-reported having been previously diagnosed by a physician or health professional. Undiagnosed diabetics were classified as those who reported not previously been diagnosed with diabetes but had a HbA1C ≥6.5%, an FPG ≥126 mg/dL, or PG ≥200 mg/dL test result. Prediabetics are not shown for categorized variables. Each model is adjusted for participant age, gender, birthplace (United States or foreign born), and smoking status. Abbreviations: CI, confidence interval; HbA<sub>1+</sub>, hemoglobin A<sub>1+</sub>; FPG, fasting plasma glucose; PG, 2-hour plasma glucose.

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. Conception and/or design: L. M.; data analyses: L. M.; interpretation of study results: L. M., L. Z., M. E. C., B. D. H., Q. L., C. C., C. C. W.; writing of first draft: L. M.; subsequent drafting of manuscript for important intellectual content: L. M., L. Z., M. E. C., B. D. H., Q. L., C. C., C. C. W.; review and edit of manuscript: L. M., L. Z., M. E. C., B. D. H., Q. L., C. C., C. C. W.; and approval of final version of the manuscript: L. M., L. Z., M. E. C., B. D. H., Q. L., C. C., C. C. W.

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