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PREVALENCE OF HUMAN IMMUNODEFICIENCY VIRUS
AMONG REPORTED TUBERCULOSIS CASES,
TEXAS 2000-2010

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M.P.H., Emory University, 2012
B.S., Emory University, 2006

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An abstract of
A Thesis submitted to the Faculty of the
Rollins School of Public Health of Emory University
in partial fulfillment of the requirements of the degree of
Master of Public Health in the Career MPH program
2012

Abstract

PREVALENCE OF HUMAN IMMUNODEFICIENCY VIRUS AMONG REPORTED TUBERCULOSIS CASES, TEXAS 2000-2010

BY

Kathryn Clelia Pezzi

The objective of this analysis was to understand the association between residence in U.S.-Mexico border region of Texas and TB and HIV coinfection. Globally, human immunodeficiency virus (HIV) is the leading contributor to the tuberculosis (TB) epidemic. In the United States the TB epidemic disproportionately affects foreign born; Mexican-born were the largest proportion of foreign born TB cases in 2010. The border region between the United States and Mexico is a socially and demographically unique region of the country with many potential risk factors for TB and HIV. This analysis used Texas statewide TB surveillance data from 2000 through 2010. Multivariable logistic regression was used to test the association between border residence and TB and HIV coinfection while controlling for a number of sociodemographic risk factors. Of 11,282 TB cases, 14.22% were infected with HIV at the time of TB reporting. TB cases reported in the Texas border region were significantly less likely to have concurrent HIV infection. Males, African Americans, injection and non-injection drug users, unemployed persons, and people age 30-49 are more likely to have concurrent TB and HIV infections. The large proportion of reported TB cases with unknown HIV status in Texas indicates that HIV screening in TB patients could be improved, and ongoing efforts to reduce TB and HIV coinfection in the border region should be continued, supported, and expanded.

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ACKNOWLEDGEMENTS

- Thank you to Dr. Kevin Sullivan, for agreeing to be my committee chair and for patiently fostering me through this process.
- Thank you to Dr. Miguel Escobedo for all his help with connecting me to the public health community in Texas, mentoring me through much of this process, and for the lovely tour of El Paso, Texas.
- Thank you to numerous state and local public health personnel in Texas who assisted with this work.

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Abstract

The objective of this analysis was to understand the association between residence in U.S.-Mexico border region of Texas and TB and HIV coinfection. Globally, human immunodeficiency virus (HIV) is the leading contributor to the tuberculosis (TB) epidemic. In the United States the TB epidemic disproportionately affects foreign born; Mexican-born were the largest proportion of foreign born TB cases in 2010. The border region between the United States and Mexico is a socially and demographically unique region of the country with many potential risk factors for TB and HIV. This analysis used Texas statewide TB surveillance data from 2000 through 2010.. Multivariable logistic regression was used to test the association between border residence and TB and HIV coinfection while controlling for a number of sociodemographic risk factors. Of 11,282 TB cases, 14.22% were infected with HIV at the time of TB reporting. TB cases reported in the Texas border region were significantly less likely to have concurrent HIV infection. Males, African Americans, injection and non-injection drug users, unemployed persons, and people age 30-49 are more likely to have concurrent TB and HIV infections. The large proportion of reported TB cases with unknown HIV status in Texas indicates that HIV screening in TB patients could be improved, and ongoing efforts to reduce TB and HIV coinfection in the border region should be continued, supported, and expanded.

Introduction

Problem Statement

Worldwide, tuberculosis (TB) is one of the most common causes of morbidity and mortality among patients infected with human immunodeficiency virus (HIV). In 2000, an estimated 9% of incident TB cases worldwide were directly attributable to HIV infection(1). In the U.S., the HIV epidemic in the 1980's and 90's along with a deteriorating TB control infrastructure led to a concurrent resurgence of TB incidence rates (2). Improved TB control programs and enhanced HIV treatment and prevention helped reverse the upward TB trend among the US-born population. However, among the foreign-born population, this decrease has occurred at a much slower rate (3).

In 2010, incidence rates of TB were the lowest they have been in the United States since national reporting started in 1953; however, foreign-born made up 60.5% (6,707) of the cases of TB in 2010 among individuals with a reported country of birth. The TB incidence rate among foreign born was 18.1 per 100,000 population TB compared to a rate of 1.6 cases per 100,000 population among US-born. Foreign-born from Mexico accounted for 23% (1,539) of all TB incidence among individuals born outside the United States, more than any other country of origin, and more cases of TB were reported among Hispanics than in any other race or ethnic group in the US (4).

In 2009, an analysis was conducted in San Diego County, California to determine trends in tuberculosis (TB) and HIV co-infection and to identify risk factors for TB/HIV co-infection. This investigation found that within the last decade, the TB and HIV co-infection burden decreased in non-Hispanic whites and African Americans, but remained constant in the Hispanic population. Also, the incidence of TB-HIV co-infection in San Diego County was almost double the national rate (0.87 per 100,000 population in San Diego; 0.5 per 100,000 population nationally). In San Diego, Hispanics had gone from accounting for 42% of co-infection cases in 1993 to representing 82% of cases in 2007 (5). These data indicate that current preventive measures are not benefitting the Hispanic population in San Diego County. The authors behind the study pointed out that this could be representative of what was happening in other US-Mexico border cities with large Hispanic populations, using El Paso and Texas as an example, but noted that similar analyses along the border had not been done for comparison.

A 2009 report from CDC's Division of Tuberculosis Elimination (DTBE) reported that in 2009, 17% of TB cases in Texas were co-infected with HIV (6). The same report shows that among foreign born TB cases in Texas, nearly 50% were born in Mexico. Data from the Texas Department of State of Health Services shows that 78.4% of TB cases in border counties are foreign born (7, 8). According to estimates based on census data from 2000, 26.8% of border residents are foreign born and 94.1% of the foreign born border residents were born in Mexico (9).

Theoretical Framework

There is an association between foreign birth, HIV status, and TB disease in the US. This association appears to have had a disproportionate impact on Hispanic immigrant and minority populations. The trends and risk factors for TB/HIV co-infection identified by Rodwell in his analysis could indicate that TB/HIV prevention and education needs in San Diego County, a border county with a large binational Hispanic population, are not being met. The same may be true in the Texas border region, an area with some comparable demographics and migration patterns, but a similar analysis has not been conducted in Texas. Further analysis of factors associated with TB/HIV co-infection in the border region, such as population mobility and proximity to Mexico could yield insights that could result in improved TB and HIV control efforts in the border region.

Purpose Statement

The purpose of this investigation is to determine the proportion of TB cases in Texas with a reported HIV co-infection; to compare the odds of HIV coinfection of TB patients in the Texas border region to the odds of reported coinfection in the rest of Texas, and to characterize clinical and sociodemographic factors associated with TB/HIV co-infection present in this population. This analysis could reveal whether or not there are factors associated with TB/HIV co-infection specific to the border region of Texas when compared to the rest of the state.

Research Objectives

This analysis has two objectives: 1) determine if TB cases in Texas border region have higher odds of HIV co-infection and 2) characterize the clinical and socio-demographic factors associated with HIV co-infection among individuals diagnosed with active TB in Texas and in Texas counties that border Mexico.

Significance Statement

This inquiry will analyze data collected during tuberculosis case investigations in Texas in order to compare TB/HIV co-infection in border counties and the rest of Texas. The findings of this investigation may allow increased insight into population-specific risks in Texas that have not been previously identified. Results will be shared with Texas PH officials to help develop recommendations for targeted public health education and screening.

Definition of Terms

A case of tuberculosis can be confirmed according to clinical or laboratory findings. The CDC/CSTE clinical tuberculosis case definition for tuberculosis disease refers to a person who meets all of the following criteria: has a positive tuberculin skin test or positive gamma release assay for *Mycobacterium tuberculosis*, clinical symptoms consistent with TB (abnormal chest radiograph or other imaging study, or clinical evidence of disease), treatment with two or more TB medications, AND has a completed diagnostic evaluation. Laboratory criteria for diagnosis includes: isolation of *M. tuberculosis* from a clinical specimen (smear positive) OR demonstration of *M. tuberculosis* from a clinical specimen by nucleic acid amplification test (culture positive) OR demonstration of acid-fast bacilli in a clinical specimen when a culture could be obtained, is a false negative, or contaminated(10). A confirmed TB case meets the clinical case definition or is laboratory confirmed. A case of TB may also be referred to as active TB disease which may be infectious or non-infectious.

Latent tuberculosis infection (LTBI) occurs when a patient has been exposed to the *M. tuberculosis* bacilli and develops a positive response to the tuberculin skin test, but has no clinical symptoms of TB. They are infected with inactive TB organisms that could reactivate and cause TB disease at a later date. Persons with latent TB infection have negative chest radiographs and cannot transmit TB (11).

The CDC/CSTE case definition for an adult case of human immunodeficiency virus (HIV) refers to a person greater than or equal to 13 years of age who has a positive HIV antibody screening test and a positive confirmatory supplemental HIV antibody test OR a positive result or report of detectable levels of HIV virus from the following HIV virologic tests: HIV nucleic acid detection test, HIV p24 antigen test, or HIV isolation (viral culture). Tuberculosis disease is one of 26 AIDS-defining conditions identified by CDC/CSTE. Under this designation, a person diagnosed with both confirmed HIV infection and TB disease would be considered an AIDS case (12).

For the purpose of this analysis, a case of TB/HIV co-infection includes individuals who have active tuberculosis disease according to the CDC/CSTE case definitions given, as well as a report of HIV disease (according to CDC/CSTE definition) at the time of TB reporting. In addition, individuals with active TB disease who have an HIV positive status in the data shared from Texas will be considered TB/HIV coinfection cases.

A binational TB case refers to an individual who meets either the US or Mexican case definition for active TB disease and one of the following criteria: 1) the ideal approach to case management would involve collaboration with TB control programs or healthcare providers on the opposite side of the border OR 2) the case is a contact of a binational TB case or is TB source case for contacts on the opposite side of the US-Mexico border. Binational cases are not necessarily border crossers or people living in border regions. This definition is consistent with the definition used by the Texas Department of State Health Services.

The border region of Texas includes 32 counties identified as geographic border counties by the La Paz Agreement in 1983 ([Appendix A](#)). The decision to use this definition for public health purposes is based on recommendations from Texas State Health Services personnel who are well acquainted with the health and demographic profile of Texas.

Review of Literature

Tuberculosis Infection and Disease

The etiologic agent for tuberculosis disease spectrum is *Mycobacterium tuberculosis*, an acid-fast bacillus. TB is spread from person to person through aerosolized microscopic, 1-5 μm particles called droplet nuclei. The small particles can stay aloft in an enclosed environment for a long time, and transmission mostly occurs through air (13). Transmission depends on a number of factors including the duration and environment of exposure, the infectiousness of the source case, and the susceptibility of the exposed individual. The disease is moderately contagious, with an estimated 30% risk of infection for household contacts of TB cases (11). Individuals with cavitory lesions visible on a chest radiograph or sputum smears with visible acid-fast bacilli are most infectious since these individuals have high bacterial burdens in their tissues and respiratory secretions. These bacteria can be spread to others when infectious people sing, cough, sneeze, or speak (11, 14). Contacts are most at risk of infection when they spend extended periods of time, generally days or weeks, in enclosed environments with a person with pulmonary TB disease (11). While pulmonary TB is the most common form of TB, extrapulmonary TB (EPTB) can occur in all organs and tissues. Between 1993- 2006, 19% of reported TB cases in the United States were EPTB cases, and up to 50% of TB cases in HIV patients is EPTB (15, 16). Patients with late-stage HIV infection ($\text{CD4} > 200/\text{mm}^3$) are more likely to have extrapulmonary TB (17, 18).

In the United States, populations with an elevated risk of TB disease include persons recently infected with TB and individuals with medical conditions that weaken the immune system. Recently infected individuals include: close contacts of TB cases, the homeless, injection drug users, recent migrants to the United States from areas with high TB rates, young children with TB infection, and persons who live or work in settings where TB transmission is high, such as hospitals, long-term care facilities, correctional facilities, nursing homes, and homeless shelters. Medical conditions that increase TB disease risk by compromising the immune system include diabetes mellitus, severe kidney disease, substance abuse, organ transplant recipients, conditions requiring the use of immunosuppressant drugs, low body weight, young age, and HIV, which is the single greatest risk factor for TB disease (19). Individuals within these risk populations or with these medical conditions are generally the focus of targeted screening programs.

Common symptoms of pulmonary TB disease include prolonged cough with or without hemoptysis, fatigue, fever, night sweats, weight loss, and chest pain. Hemoptysis, the coughing up of bloody sputum, can occur when small blood vessels are ruptured in the lungs as the disease progresses. If left untreated, TB disease can permanent lung damage or death (19).

When an individual inhales the TB droplet nuclei organisms enter the lung aveoli, sub-clinical infection can occur locally as small microscopic lesions with viable TB organisms form in the lungs and then heal (19). In most healthy individuals there is an estimated lifetime risk of around 10% of developing active TB disease from an initial TB infection and, of those individuals who develop TB disease, half will develop disease within the 2 years following their initial infection (19, 20). Certain subpopulations are at greater risk of experiencing a rapid progression to TB disease, particularly young children and immunocompromised individuals(19). There is a vaccine available for TB, the Bacille Calmette-Guérin (BCG) vaccine, but its protection varies widely among populations and is not complete (21). It is generally given to protect young children against tuberculosis meningitis and disseminated (miliary) tuberculosis, the two most deadly forms of childhood TB, but its efficacy against pulmonary TB is unreliable. The BCG vaccine is not routinely given in the United States and some countries in Europe where TB rates are low. However, global BCG coverage is quite high. The WHO estimated that global vaccination coverage of BCG was 89% in 2007 and the reported BCG coverage in Mexico was 98% in 2010 (22).

Tuberculosis Testing

TB infection can be detecting using multiple methods. A common method of screening for TB infection is the tuberculin skin test (TST). In this test, a small measured amount of purified protein derivative (PPD) tuberculin is injected just under the skin (Mantoux method). The skin reaction at the test site read within 48-72 hours to determine if the person is infected; the reaction should be measured in millimeters of induration (swelling) at the site of the PPD injection. The interpretation of the test depends on individual factors associated with the person being tested. For example, healthy adults without known TB risk factors would need to have 15 mm or more induration in order for their TSTs to be considered positive. A TST of 5 mm or more of induration is considered positive in individuals with HIV infection or other medical conditions that compromise the immune system, close contacts with known TB cases, and patients with chest x-rays suggestive of prior TB disease. The test in individuals with other known risk factors,

such as recent immigrants from high-burden countries, injection drug users, young children and lab personnel would be considered positive with 10 mm or more of induration (23). Both false negative and false positive reactions to the TST can occur, and one of the contributing factors to a false negative test is HIV infection; individuals with HIV infection may be anergic, i.e. unresponsive to antigens used in the Mantoux test. In these individuals, the use of PPD would not produce any response even if the patient was infected with TB. In a study of hospitalized patients, 63% of the HIV positive patients were anergic (24). In addition, false positives to the TST can occur in individuals previously vaccinated with BCG vaccine or infected with non-tubercular mycobacteria (14, 25). However, previous BCG vaccination is not a contraindication to TST and a positive TST reading in someone with previous BCG vaccination is presumed to indicate LTBI (26).

In addition to the TST, there are two FDA-approved interferon-gamma release assay (IGRA) tests for identifying TB infection, the QuantiFERON® -TB Gold In-Tube test (GFT-GIT) and the T-SPOT®.TB (27). Testing with these methods requires only one visit to a physician since testing is conducted on a blood sample and no follow-up reading is required. Another benefit of this testing is its specificity; individuals with the BCG vaccine or most non-tuberculosis mycobacteria will not test positive when using either of these IGRA tests (27). CDC recommendations allow for the use of IGRA testing in place of a TST in all situations in which TSTs would be used to diagnose TB infection. CDC further recommends that IGRA testing be used in populations that have historically low rates of return to clinics for TST readings (such as the homeless) and in persons who have received the BCG vaccine (27). In a number of studies that compared TST tests to IGRA tests, IGRAs appeared to be a viable and cost effective alternative to TST screening when used for in low-prevalence countries like the U.S. for targeted TB testing of high-risk populations such as healthcare workers, foreign born from high prevalence countries, and close contacts of TB cases (28, 29). In addition, there is evidence that IGRA tests may be useful tools for identifying extrapulmonary TB disease (30).

These tests each provide evidence that an individual is infected with tuberculosis, but none of them distinguish between infection and active disease. To diagnose active TB disease, a positive culture of *M. tuberculosis* must be obtained from tissue or secretion samples. In a suspected case of pulmonary TB, three sputum samples are collected for laboratory confirmation of the presence of TB bacilli; smears can be done on directly on clinical samples or on amplified specimens.

Samples are stained by the Ziehl-Neelsen or fluorescent techniques for acid-fast bacteria (AFB) and examined microscopically. Smears are considered positive when there are visible acid-fast bacteria, although these smears are not specific for *M. tuberculosis* since the stains will also adhere to non-tuberculosis mycobacteria. In addition, the sensitivity of the test is low; up to 50% of patients with pulmonary TB can have negative AFB smears, and HIV positive patients often have negative smears (31). For this reason, smear results alone do not allow a definitive diagnosis, but the test is a good screening tool, with results in as little as 24 hours, and an assessment of number of bacilli present in the smear gives an indication of how infectious a patient might be (26).

Nucleic acid amplification (NAA) techniques are additional testing tools that can identify TB bacilli directly from clinical specimens without the need for culture. Culture takes longer than the other means of testing; depending on whether liquid or solid media is used it can take between 1-8 weeks to detect mycobacterial growth (32). In patients with suspect TB, NAA testing produces results in 24-48 hours. In addition, NAA tests have a positive predictive value (PPV) of >95% in individuals with AFB positive smears and a PPV 50%-80% in individuals with AFB negative smears. CDC considers a specimen that is smear- and NAA-positive to be a confirmed case of TB, but TB diagnosis cannot be ruled out solely on the basis of a negative NAA test (32). The gold standard in tuberculosis diagnostics remains culturing samples, and cultures should be done for any suspected case of tuberculosis. Cultures are also necessary for drug-susceptibility testing, speciation, and genotyping, all of which should be done to determine the best treatment course for a particular patient. If a patient has positive cultures, they are considered a confirmed active case of TB. In patients with extrapulmonary TB, diagnosis is more challenging. It may be difficult to obtain specimens of suspected TB disease from some sites in the body and specimens may lack adequate bacilli for testing (16). In individuals with EPTB, NAA testing may offer improved diagnostic capabilities for physicians when combined with clinical evaluation (16).

Treating Latent Tuberculosis

When an individual is diagnosed with latent TB infection, treatment is recommended to eliminate the infection and to prevent future progression to TB disease. Without treatment, TB infection could eventually lead to TB disease in around 10% of healthy adults (immunocompromised subgroups have higher risk of progression to TB disease). There are multiple recommended treatment regimens for individuals with LTBI; the preferred regimen consists of 9 months of daily

doses of isoniazid (INH) which is suitable for both HIV positive and negative individuals. Alternative treatments include different durations of treatment with INH and the use of other drugs, i.e. rifampin, and rifampin combined with pyrazinamide, and different durations of treatment, though the shortest treatment duration lasts at least two months. Non-INH-based treatments have unproven efficacy, may be more toxic, and are administered in the setting of INH resistance or intolerance. Some TB treatments can be administered concurrently with HIV treatments. However, treatment of TB/HIV co-infected patients is complicated and requires oversight of a healthcare provider throughout the treatment regimen to ensure patient compliance with treatment and to prevent dangerous drug interactions or side effects to treatment (33).

Treating Tuberculosis Disease

Treatment for tuberculosis disease is more complicated and consists of two phases; first, the initial phase that lasts around 8 weeks during which patients take a combination of 3-4 drugs. After this, patients enter the continuation phase where they continue taking two of the initial drugs. There are several regimens made up of different drugs or different dosing schedules, but most still involve multiple doses of the drugs each week. The continuation phase can last between 18 and 31 weeks. Lapses in drug treatment can lead to the development of drug resistant TB disease which is more difficult to treat. Due to the long duration of TB treatment, the need to prevent treatment interruptions and the need to monitor patients for potential side effects from drug toxicity, directly observed therapy (DOT) is recommended for all TB patients. In this form of treatment, healthcare practitioners watch TB patients take their doses of TB medications. This method improves treatment adherence and allows healthcare workers to assess patients for signs of adverse drug reactions, treatment failure, or non-compliance (34). Non-compliance with TB treatment is associated with an increased likelihood of the development of multi-drug resistant (MDR) TB which is more difficult and costly to treat (35). Treatment of TB in HIV patients is even more complicated and must be monitored closely to prevent drug interactions and to monitor events of immune reconstitution inflammatory syndrome (IRIS), an initial worsening of TB symptoms in patients whose immune systems return after HIV (36).

Testing for HIV Co-infection

In 2006 CDC released its latest guidelines on HIV testing in a healthcare setting. These guidelines recommend HIV testing become a routine part of all clinical care, including for patients being seen in a TB clinic. Under these new guidelines, all TB patients with unknown

HIV status should receive HIV testing when they enter the TB clinic unless they explicitly opt-out of testing (33). CDC guidelines also recommend that all HIV patients, regardless of risk, receive TB testing upon their HIV diagnosis. In addition, the guidelines recommend that HIV patients be tested annually for TB infection if they are at increased risk of exposure to active TB due to occupational, lifestyle, or demographic factors. The rationale behind these recommendations is to identify and treat TB infection in HIV patients since this population is at particular risk of progression to TB disease; and to identify TB disease as early as possible to improve treatment outcomes (36, 37).

TB/HIV Syndemic

Screening HIV patients for TB infection and disease is necessary to control the TB/HIV syndemic that has resulted in a global increase in the burden of TB and significant morbidity and mortality. Syndemics are synergistic interactions between two or more diseases that results in an excess burden of disease in a population. HIV impairs the cell-mediated immune system of its host, which contributes to an increase in individuals in the population susceptible to TB infection. HIV accelerates the progression of TB infection to TB disease; TB disease likewise speeds the progression of HIV to AIDs (17, 37, 38). Among individuals with HIV, the lifetime risk of TB disease increases considerably. Individuals without HIV infection have an estimated 10% lifetime risk of developing TB disease; among people infected with HIV the estimated annual risk of developing TB disease is 5-10% (39).

In the United States, an estimated 26% of TB cases are attributable to HIV infection, higher than the estimated attributable risk in other industrialized countries, even those countries that have higher prevalence of both TB and HIV infection. This is likely due to the fact that HIV and TB infection tend to concentrate within the same high-risk subgroups in low-prevalence countries like the U.S., disproportionately affecting these populations (1). Groups particularly at risk of TB/HIV co-infection in the U.S.-Mexico border region include drug-abusers and the homeless (40). A study in Alameda County, California calculated that 93% of the incident TB cases among HIV positive individuals were attributable to HIV infection (41). Testing TB patients for HIV and HIV patients for TB are key steps to controlling the TB/HIV syndemic in the United States; routine testing can identify TB infection early, before it progresses to disease; and the disease can be treated to prevent progression and spread throughout vulnerable populations like HIV patients.

TB/HIV and the U.S. Mexico Border

The 100 km wide area on either side of the border between the United States and Mexico is a unique region with distinct characteristics differing from the non-border regions of the United States and Mexico. Poverty, strained infrastructure, social problems, lack of insurance and affordable healthcare, and movement back and forth across the border are all factors that may contribute to the higher incidence of a number of infectious diseases in the border region compared to non-border regions of both the U.S. and Mexico (5, 42). Poverty rates are higher in U.S. side the border region, and the population is highly mobile, moving back and forth between the two countries for work, school, family and healthcare, compared to non-border regions of the United States. In addition, the population on the U.S. side of the border region has increased significantly in the past decade, causing strain on the infrastructure of the border region, including healthcare resources (9, 43, 44). In a 2010 report from the Border Health Commission, Texas border counties had the lowest ratio of health providers per 100,000 population of any of the other borders states; the ratio of health professions in Texas border counties was less than half the ratio for the U.S. (9). Drug use and prostitution are also significant problems along the border since numerous drug and human trafficking routes run through this region from South and Central America and the interior of Mexico to the United States (45). While free TB treatment and HIV testing are readily available in both Mexico and the United States, undocumented residents living in the U.S. border region face challenges to accessing affordable healthcare in the United States and may avoid seeking testing or care for disease for fear of deportation (45). The deportation of migrants back to Mexico often results in populations of homeless migrants living in Mexican border cities; these populations are more likely to engage in HIV-related risk behaviors (45, 46).

Mobility back and forth across the border likely plays a significant role in infectious disease transmission in the region. In one study that compared characteristics of foreign born Hispanic TB patients who lived in U.S. border counties to Hispanic TB patients not living in border counties, Mexican-born border TB patients were significantly more likely to return to Mexico, and more likely to have visited Mexico within the 12 months prior to their TB diagnosis. In addition, more than a third of the border-county TB patients traveled into Mexico on a daily or weekly basis (44). Another study conducted in Texas comparing mobile to non-mobile border dwellers on the U.S. side of the border found that 20% of the population crossed the border at least once a week, and 30% of the population crossed the border at least once a month. Of the population identified as mobile, 62.3% said they were crossing the border for medical care or

supplies. Mobile border residents had a lower socio-economic than non-mobile border residents (47). In a telephone survey conducted in El Paso County, Texas found that 27% of the respondents reported crossing the border in the previous two years for some kind of health care (43). Health insurance coverage is low among foreign born Hispanics in the border region is particularly low compared to other populations; 62% lack health insurance coverage (48).

Deportation is another factor associated with increased occurrence of TB and HIV risk behaviors(45). Undocumented migrants apprehended by law enforcement in the U.S. are often repatriated across the border to cities like Tijuana and Ciudad Juarez with few resources; these migrants often opt to stay in these areas to make a later attempt to return to the United States (45). Repatriation of Mexican migrants to cities on the Mexican side of the border has been linked to increased HIV risk behaviors and transmission; in one study, male injection drug users who were deported from the United States had four times the odds of HIV infection than male IDUs without a history of deportation (49). Repatriation may also be associated with increased odds of TB infection; in another study on TB in injection drug users in Tijuana, individuals with a self-reported TB history were significantly more likely to have been deported from the United States than those without TB. In addition, individuals with TB history were six times more likely to have traveled to the United States than those without a history of TB (50).

The United States immigration policies may play a role in this relationship between deportation and TB; when undocumented migrants are detained in U.S. Immigration and Customs Enforcement (ICE) detention facilities they are screened for TB. ICE procedures focus on law enforcement as opposed to public health practices, and detainees may be deported before the results of their TB test are received, their TB treatment can be completed or continuity of care measures are arranged with Mexico. ICE is legally mandated to repatriate migrants within 3 months of their apprehension, an insufficient time to ensure TB treatment is completed. Public health authorities may request an ICE stay-of-removal consideration to ensure treatment completion in special situations, but this is difficult to carry out. (Personal communication, July 2012) In addition, residence in a detention center is a risk factor for tuberculosis, so detainees who may not have had prior TB disease may be exposed to TB disease while in detention (51). Texas county public health offices might receive a report of a TB case in a detainee only to find that the detainee was deported back to Mexico and lost to follow-up; these detainees may never find out they have TB disease (Personal communication, July 2012). This issue was cited as a

significant challenge for TB case management in the U.S.-Mexico border region by the Texas State TB Control Program Manager at the 2010 U.S.-México Border Tuberculosis Consortium (52).

In 2010, the reported incidence rate of TB in the United States was 4.1 per 100,000; prevalence was 4.8 per 100,000. In Mexico, the reported 2010 TB incidence and prevalence were 16 and 18 per 100,000, respectively. Incidence of TB co-infection in HIV positive individuals is more than twice as high in Mexico compared to the United States. In 2010, TB incidence in HIV positive individuals in Mexico was 0.81 per 100,000; in the United States it was 0.36/100,000 (53, 54). In 2010, in the U.S. more cases of TB were reported among Mexican-born than any other country of origin; 23% of foreign born TB cases were born in Mexico(55). In 2010 the TB incidence rate (9.9/100,000) for the Texas border region, with its highly mobile population and significant Mexican-born population, was almost twice as high as the TB incidence rate for the state of Texas (5.5/100,000)(56). In Mexico, screening for TB infection in populations at high risk of TB/HIV co-infection is uncommon, and the use of IGRA testing has been recommended over the use of TST in Mexico because the population's high BCG coverage that makes TST less useful at distinguishing true TB infection (40, 57).

While the reported incidence and prevalence of TB are higher in Mexico than in the U.S., the converse is true for HIV (53, 54, 58). According to the 2010 UNAIDS Global Report, Mexico is considered a low prevalence country for HIV (prevalence = 0.3%); the United States has twice the estimated prevalence of HIV infection as Mexico (0.6%) (58). However, there is ample evidence that subepidemics exist in regions and populations of Mexico where higher rates of HIV risk behaviors such as intravenous drug use and sex work take place and overlap (45, 49, 59-62). In Ciudad Juarez and Tijuana, the large Mexican border cities directly across from El Paso and San Diego, the prevalence of HIV are among the highest in Mexico, particularly in injecting drug users, men who have sex with men (MSMs), female sex workers, and female sex workers using injection drugs (45, 61). In a study that examined HIV prevalence in Tijuana and Ciudad Juarez, 2.8% of the IDUs tested in both cities were HIV positive (59).

Though TB and HIV prevalence in the U.S. and Mexico are both fairly low compared with burden of these diseases in other regions of the world, in countries with low prevalence of TB and HIV, subpopulations such as the homeless and drug users are at increased risk of TB/HIV

coinfection (40). A recent study found that the prevalence of TB infection, HIV infection, and TB/HIV coinfection in high-risk populations (injecting drug users, non-injecting drug users, prostitutes, and homeless) in Tijuana, Mexico was 57%, 4.2% and 2.2%, respectively (40). These prevalences were higher than those reported for the rest of the state and country, and indicate that there is an unmet need to provide screening to identify TB infections earlier in these high risk populations (63). In addition, there is evidence that this HIV sub-epidemic in high-risk populations has spilled over into the general public; in a 2006 study on women delivering infants at Tijuana General Hospital, 1.12% of screened labor/delivery patients were HIV positive, a higher proportion than previously established estimates in Mexico (64). In the United States there is evidence that the TB/HIV co-infection is disproportionately affecting Hispanics in the U.S.-Mexico border region; an analysis of co-infection in San Diego indicated that TB/HIV co-infection rates in all populations except for Hispanics had decreased between 1993-2007; co-infection rates among Hispanics did not change over the same time period. The majority of new TB/HIV cases in San Diego currently occur in Hispanics born in Mexico (5).

These statistics paint a picture of a region on either side of the border that is vulnerable to the TB/HIV syndemic. The higher HIV prevalence on the US side of the border, coupled with the higher prevalence of TB in Mexico, high level of cross-border mobility, vulnerability of populations on both sides of the border, and presence of HIV subepidemics in high-risk populations on the Mexican side of the border combine to create conditions that could theoretically contribute to the TB/HIV syndemic in the border region of Texas. Are individuals living in the border region of Texas more likely to have HIV infection alongside TB disease than non-border residents of Texas?

Methods

Population and Sample

This analysis used data obtained from Texas Department of State Health Services (DSHS). Active tuberculosis disease is a nationally reportable disease in the United States and case reporting of TB is required by law in Texas. When a suspected case of tuberculosis is identified in the state, local public health agencies use the standard, statewide forms (Texas 400A and 400B, [Appendix B](#)) to collect demographic and clinical data from the patient, including HIV status and other known TB risk factors. Texas currently uses the NEDSS Base System (NBS) Tuberculosis

Program Area Module (TB PAM) as its data management and reporting system for TB data. The TB PAM system permits direct reporting of TB cases in Texas to CDC.

Data collected prior to 2009 were stored in the Tuberculosis Information Management System (TIMS), a national surveillance and case management software application system that is no longer in use. A request was made to the Texas DSHS for access to the records for all confirmed cases of TB disease captured in either the TIMS or TB PAM databases between January 2000 and December 2010. When the request was granted, the Texas DSHS data manager merged datasets from both the TIMS and TB PAM databases to align the common variables from the two datasets and create a complete dataset for the appropriate time period. The data was shared with investigators after personal identifiers such as names, social security numbers, specific addresses, and case ID numbers were removed.

The population for this investigation includes all confirmed cases of TB disease according to the official CDC case definition that were reported in Texas between January 1, 2000 and December 31, 2010 in individuals 18 years or older at time of report (n=15,430). Tuberculosis cases included in this analysis had to be confirmed by laboratory, or by clinical provider diagnosis. This population does not include individuals with latent TB infection. Individuals were identified as co-infected if they had a 'Positive' HIV status in the TB case dataset shared by Texas.

In 2007, HIV testing of TB cases in Texas became standard and the majority of TB cases were referred for HIV testing; however, HIV testing for TB cases is not mandatory and TB patients may refuse testing. HIV status is captured as positive, negative, indeterminate, not offered, refused, unknown and missing in the database. For the purposes of this analysis, individuals with indeterminate, not offer, refused, unknown and missing HIV status were combined into a single 'Unknown' category and were not included in the multivariable analysis, though it is likely that this category included some co-infection cases. The final population for this analysis consisted of 11,282 reported cases of tuberculosis.

Research Design

Because the source of the data used in this analysis is reports from a TB program, the population and medical risk factor data collected primarily includes variables known to be associated with TB disease, not HIV disease, though there is some overlap in the risk factors and risk populations

for both diseases. Demographic variables included gender, race, ethnicity, age, county where case was reported, country of birth, and length of time in the US. Smaller ethnic groups ('Multi-racial,' 'Other,' 'American Indian or Alaskan Native,' 'Native Hawaiian or Other Pacific Islander' and 'Unknown') were also excluded from the analysis; each excluded race category had less than 35 records total (78 records were removed in total). TB risk factors included homelessness, drug and alcohol abuse, and residence in a prison or long-term care facility within the 12 months prior to TB diagnosis. These variables were coded as 'yes,' or 'no.' Employment status was coded as 'unemployed' or 'employed.' Individuals with unknown employment status (n=4) were excluded from the analysis. Year of diagnosis was included as well as clinical information on the site of TB disease, coded 'pulmonary,' 'extrapulmonary,' and 'both.'

Data Analysis

All analyses were performed using SAS software, Version 9.2/3 of the SAS System for Windows. Frequencies were calculated to describe the demographic characteristics of the study population. For the purposes of this analysis, HIV status was the outcome of interest, and border region residence, determined by county of disease report, was the exposure of interest. Chi-square tests from univariate analyses were used to identify factors significantly related to HIV infection for later inclusion in the multivariable models. Two-sided statistical tests were considered significant at $P < 0.05$.

Bivariate logistic regression identified significant ($P < 0.05$) effect modifiers in the relationship between border residence and HIV infection among reported TB cases. Effect modification was assessed using the Breslow-Day test for interaction; if the test had a p-value less than 0.05 the test was considered significant and the variable was included in the multivariable model as an interaction term. Multivariable logistic regression models were developed using backward elimination of terms; independent variables in the model included interaction terms identified from the bivariate analysis and all other variables assessed in the initial modeling. Confounding was assessed in the multivariable model by comparing the gold standard OR (from the multivariable model that included all variables but no non-significant interaction terms) to the OR obtained after dropping potential confounders that were not statistically significant predictors of the outcome, controlling for all remaining variables in the model. A variable was considered a confounder if removing it from the model resulted in a 10% or larger difference in the subsequent model's OR compared to the gold standard OR. The Hosmer-Lemeshow test for goodness of fit

was used to evaluate the model's fit. The final model was also run using the PROC GENMOD procedure in SAS to estimate the prevalence ratio to evaluate how accurately the odds ratio produced by the logistic regression model assessed the association between exposure and outcome. Output from the multivariable regression modeling is included in [Appendix D](#). This project was submitted to the Emory University Institution Review Board (IRB) for clearance as well as the Centers for Disease Control and Prevention IRB. Both reviews approved this project and returned a non-research determination.

Limitations

The data used in this analysis is cross-sectional in nature, so it is impossible to draw conclusions about whether HIV infection preceded TB infection. Individuals who may have been diagnosed with HIV infection prior to their TB diagnosis may not disclose their HIV infection to TB staff and can refuse testing, so their HIV status would remain unknown in the TB reporting. As a result, an unknown quantity of co-infection cases may not have been captured through the TB surveillance data alone, so these data do not represent all cases of TB/HIV co-infection in Texas.

This analysis uses data collected over 10 years of the Texas TB surveillance program. During that time period TB testing protocol, case definition, the Texas reporting system and the variables it collects, and HIV testing recommendations for TB patients took place. Changes in how TB cases foreign-born and foreign visitor TB cases could be counted by a reporting jurisdiction also changed; both of these of these categories would likely have an impact on the border region compared to other areas of the United States with a less mobile foreign-born population or fewer foreign visitors receiving treatment for TB (65).

Delimitations

This analysis was limited to individuals 18 years of age and older. A significant proportion of the TB cases (26.3%) were excluded from the analysis due to unknown HIV status. In addition, not all variables of interest were routinely collected or reported through the TB reporting system for every TB case and some of those cases were excluded from the multivariable analysis. A number of variables that are particularly interesting in the context of studying TB infection in the border, including variables designating a case a binational case or capturing visa status upon arrival to the U.S., were rarely collected and could not be included in this analysis. These factors all limit the conclusions that can be drawn from the study. Specifically, data on the HIV status of

TB patients should be interpreted with caution because these data are not representative of all TB/HIV co-infection cases for the state of Texas.

Results

Demographics

There were total of 16,895 reported cases of TB in Texas between January 1, 2000, and December 31, 2010, with 15,430 cases of TB disease reported in individuals 18 years or older. Of these cases, 1,618 (10.49%) of the cases were co-infected with HIV disease, and 4,063 (26.3%) cases did not have a known HIV status; the remaining reports were HIV negative. The border region had a lower proportion of TB cases with an unknown HIV status; this was true for all races and ethnicities. The cases with unknown HIV status were excluded from further analyses resulting in a total of 11,282 reported cases of TB.

Of the reported cases of TB disease included in the analysis, 1,604 (14.22%) cases were HIV positive (Table 1). Of all TB patients, 2,105 (18.66%) were reported from counties in the border region of Texas. Slightly more than half of Texas TB cases included in this analysis were Hispanic (5,755, 51.01%). There were nearly equal numbers of foreign born (5,599, 49.63%) and native (5,527, 50.37%) TB cases.

HIV prevalence was lower among border county residents (7.22%) than among non-border residents (15.80%). In the state of Texas, HIV prevalence was greatest among individuals between the ages of 30-39 (23.63%) and 40-49 (20.85%), and higher in men than women (16.15% vs. 9.85%). Prevalence of HIV was also lower in Hispanics (10.31%) compared to non-Hispanics (17.97%), and African Americans (26.99%) had the highest HIV prevalence compared to Asians (3.52%) and whites (10.90%). HIV prevalence was also higher in foreign born, injection drug users, non-injection drug users, the homeless, and individuals with extrapulmonary TB.

Trends

Over the ten year time period, the proportion of each race and ethnic group with unknown HIV status decreased (Figures 2-6, [Appendix C](#)). African-Americans consistently had the lowest proportion of unknown HIV status, but over the period of this analysis the proportion of unknown

decreased from 19.76% to 13.3%. Hispanics had one of the largest decreases in the proportion of unknown HIV status; this population began the decade with 30.77% of the population reporting an unknown status; by 2010 down to 20.96% of Hispanics diagnosed with TB had an unknown HIV status. This proportion was lower than non-Hispanics with unknown HIV test results in 2010 (24.01%). Asians had this highest proportion of TB cases with unknown HIV status; 40.85% of this population had unknown HIV status in 2000. In 2010, the proportion of TB cases in Asians with unknown HIV status had decreased to 31.84%, but that was still the highest proportion of unknowns compared to all other race and ethnic groups.

Multivariable Regression

The bivariate analysis identified five potential interactions (Breslow-Day Test for interaction p-value was less than or equal to 0.05) with border residence: foreign birth, excess alcohol consumption in the past 24 months, length of time in the United States, correctional facility residence at time of diagnosis, and location of TB (pulmonary, extrapulmonary, or both). These interaction terms were included in the final multivariable model but none retained their significance and they were all dropped. After the interaction terms were dropped from the model, the remaining variables in the model were assessed for confounding using the technique described in the methods section; none of the variables removed from the model were confounders but the decision was made to report the results of the ‘gold standard’ model that contained all non-interaction terms. Most of the variables included in the model were frequently reported in the literature as being related to TB/HIV coinfection and were of interest in this analysis.

When controlling for the variables listed in Table 3 in a multivariable logistic regression model, a case of TB/HIV co-infection was less likely to be reported in the Texas border region (AOR 0.63; 95% CI: 0.511,0.776) (Table 2). In addition, a case of TB/HIV co-infection in Texas was more likely to be male (AOR 2.24; 95% CI:1.94, 2.59), African-American (AOR 2.04; 95% CI: 1.705, 2.441), and unemployed (AOR 1.98; 95% CI: 1.74, 2.24). Age was significantly associated with co-infection, with HIV positive cases most likely to be between 30 and 30 years old (AOR 12.68; 95% CI: 8.87, 18.15). Foreign birth, length of time in the US, and Hispanic ethnicity were not significantly associated with increased or decreased likelihood of HIV co-infection among TB patients in Texas.

The sole clinical variable included in the analysis, location of TB infection, was significant; cases TB/HIV co-infection were more likely to have extrapulmonary TB (AOR 2.45; 95% CI 2.12, 2.82) or both extrapulmonary and pulmonary TB (AOR 3.36; 95% CI 2.76, 4.09).

Among risk factors associated with HIV and TB, injection drug use was significantly associated with HIV co-infection in TB patients (AOR 2.37; 95% CI 1.83, 3.07). Non-injection drug users were also more likely to have HIV (AOR 1.59; 95% CI 1.32, 1.91) than non-drug users.

Correctional facility residents and heavy alcohol users were less likely to have a positive HIV test than people not reporting these risk factors in Texas. Long term care facility residence and homelessness were not associated with co-infection with HIV when controlling for all other variables.

The Hosmer-Lemeshow test for goodness of fit indicated a good fit ($\chi^2=7.56$, 8 d.f., $p=0.48$). In addition, the border prevalence ratio estimated from fitting the same model using PROC GENMOD indicated that the OR from the logistic regression did not overstate the association between border residence and HIV co-infection among TB patients (PR 0.63; 95% CI 0.51, 0.78).

Table 1. Characteristics of Reported TB Cases and Prevalence of HIV by Characteristic Among TB Cases, Texas, 2000-2010.*

Characteristic	TB Cases		HIV Prevalence	
	n=11,282	%	n=1604	14.22%
HIV Status				
Positive	1604	14.22	-	-
Negative	9678	85.78	-	-
Border County Resident				
Yes	2105	18.66	152	7.22
No	9177	81.34	1452	15.80
Age at Diagnosis				
18-29	2384	21.13	200	8.38
30-39	2374	21.04	562	23.63
40-49	2592	22.97	541	20.85
50-64	2660	23.58	264	9.92
65+	1272	11.27	37	2.91
Sex				
Female	3453	30.61	340	9.85
Male	7829	69.39	1264	16.15
Ethnicity				
Hispanic	5755	51.01	570	10.31
Non-Hispanic	5527	48.99	1034	17.97
Race				
Asian	1250	11.08	44	3.52
African American	2897	25.68	782	26.99
White	7135	63.24	778	10.90
Foreign Born				
Yes	5599	49.63	566	9.96
No	5683	50.37	1038	18.54
Length of Time in the US (n=5683)				
<1 year	1014	9.01	97	9.57
1-4 years	1375	12.21	142	10.33
5-9 years	857	7.61	93	10.85
10-19 years	1100	9.77	112	10.18
20+ years	1313	11.66	117	8.91
missing	24			
Year of Diagnosis				
2000	943	8.36	154	16.33
2001	973	8.62	169	17.37
2002	918	8.14	184	20.04
2003	980	8.69	163	16.63
2004	1136	10.07	144	12.68
2005	989	8.77	132	13.35
2006	1067	9.46	157	14.71
2007	1129	10.01	143	12.67
2008	1162	10.3	137	11.79
2009	1045	9.26	121	11.58
2010	940	8.33	100	10.64
Injection Drug Use**				
Yes	371	3.29	126	33.96
No	10911	96.71	1478	13.55
Non Injection Drug Use**				
Yes	1055	9.35	248	23.51
No	10227	90.65	1356	13.26
Excessive Alcohol Use**				
Yes	2555	22.65	367	14.36
No	8727	77.35	1237	14.17
Homeless**				
Yes	728	6.45	158	21.70
No	10554	93.55	1446	13.70

Correctional facility resident†					
	Yes	1146	10.17	162	14.14
	No	10122	89.83	1440	14.21
	<i>missing</i>	14			
Long Term Care Facility Resident†					
	Yes	186	1.69	33	17.74
	No	10841	98.31	1551	14.31
	<i>missing</i>	255			
Employment**					
	Employed	4951	43.88	1052	16.62
	Not Employed	6331	56.12	552	11.15
TB Location					
	Pulmonary	8589	76.13	964	11.22
	Extrapulmonary	1967	17.43	439	22.32
	Both	726	6.44	201	27.69

*Excludes TB case reports with unknown HIV status

**Engaged in this risk behavior in the 24 months prior to TB diagnosis

†At time of diagnosis

Table 2. Final Full Model - Adjusted Odds of HIV among TB Cases in Texas with Known HIV Status, 2000-2010 (n=10,989*).

Characteristic	Adjusted Prevalence Odds Ratio	95% Confidence Intervals		p-value
Border				
Yes	0.63	0.51	0.78	<0.01
No	ref			
Foreign born				
Yes	1.23	0.97	1.58	0.09
No	ref			
Ethnicity				
Hispanic	0.83	0.67	1.03	0.09
Non-Hispanic	ref			
Race				
Asian	0.21	0.15	0.31	<0.01
Black	2.04	1.71	2.44	<0.01
White	ref			
Sex				
Female	ref			
Male	2.24	1.94	2.59	<0.01
Age				
18-29	3.74	2.56	5.45	<0.01
30-39	12.68	8.87	18.15	<0.01
40-49	8.94	6.27	12.77	<0.01
50-64	3.65	2.54	5.25	<0.01
65+	ref			
Length of Time in the US				
<1 year	0.79	0.57	1.09	0.15
1-4 years	0.78	0.58	1.04	0.09
5-9 years	0.87	0.64	1.20	0.40
10-19 years	0.83	0.61	1.11	0.20
20+ years	ref			
Injection Drug Use**				
Yes	2.37	1.83	3.07	<0.01
No	ref			
Non-Injection Drug Use**				
Yes	1.59	1.32	1.91	<0.01
No	ref			
Excessive Alcohol Use**				
Yes	0.65	0.56	0.76	<0.01
No	ref			
Homeless†				
Yes	0.93	0.75	1.16	0.53
No	ref			
Correctional facility resident†				
Yes	0.71	0.58	0.89	<0.01
No	ref			
Long Term Care Facility Resident†				
Yes	1.15	0.74	1.78	0.55
No	ref			
Location				
Extrapulmonary	2.45	2.12	2.82	<0.01
Both	3.36	2.76	4.09	<0.01
Pulmonary	ref			
Employment†				
Not Employed	1.98	1.74	2.24	<0.01
Employed	ref			

*Number of observations (records) included in this model

**Engaged in this risk behavior in the 24 months prior to TB diagnosis

†At time of diagnosis

Discussion

This analysis sought to compare the likelihood of prevalent HIV infection in TB cases reported in the U.S.-Mexico border region of Texas to non-border regions of Texas, controlling for common TB risk factors. The objective of making this comparison was to determine if the residence in unique environment of the U.S-Mexico border region contributes to increased odds of HIV co-infection. The population assessed for this analysis included all TB cases reported in Texas between 2000 and 2010. These data only include HIV infection reported in the context of the Texas Department of State Health Service's TB reporting program; cases of HIV reported through ongoing HIV surveillance and data collection in Texas were not included in the analysis.

Conclusions

The results of this analysis indicate that HIV infection in TB patients is less likely to be reported in the 32 border counties of Texas when compared with TB reporting in the rest of the state. In addition, Hispanics were no more likely to have HIV co-infection when compared to non-Hispanics. Foreign birth was also not significantly associated with HIV infection.

Almost 50% of all the TB and HIV co-infection cases reported for this 10 year span of data were reported in African-Americans (48.27%) even though they make up a smaller proportion of the Texas population (12%) than Hispanics (38%) or Non-Hispanic whites (44.8%) (66). African-Americans were also disproportionately affected by TB in Texas; African-Americans made up 23.1% of reported TB cases over the 10 year period. While Hispanics had lower odds of HIV than non-Hispanics, they are disproportionately affected by TB, making up 48.3% of TB cases during the same period.

The decision to leave the unknown HIV status population out of this analysis was a potential source of bias. The population with unknown status likely contained a combination of HIV positive and negative TB cases; the proportion of HIV positive TB cases among the unknowns likely changed over the ten year time period as testing recommendations and practices changed and HIV testing became more consistent in the general population. The different proportions of unknown HIV status between the various race/ethnic groups indicates that TB programs in Texas may either not offer HIV testing to some populations as frequently as others, and/or that HIV testing acceptability is lower among some populations compared to others. While the proportion of all race and ethnic groups with unknown HIV status decreased over the time period, there were

still fairly large proportions of Asians (over 30%), whites (~24%), and Hispanics (~21%) who had unknown HIV status, particularly in non-border regions of Texas. However, this could also be an issue of reporting; individuals may be getting tested for HIV, but reporting of HIV testing results through the TB reporting system may be incomplete.

In the case of Texas's border population, border residence may be protective against HIV infection, possibly because the close proximity to Mexico could facilitate more frequent travel to Mexico to see family and seek healthcare. A number of studies of migrant populations in the United States note that HIV risk is high for migrants to the United States compared to non-migrants in Mexico for many reasons – the U.S. has a higher HIV prevalence compared to Mexico, many migrants lack of access to healthcare and HIV testing, the social isolation of migration contributes to increased HIV risk behaviors such as use of male sexual partners, greater number of sexual partners, prostitution, and drug use (67-69). Migrants living in the border may not have as many of these risk factors because they are less isolated from family and protective cultural factors; frequent travel to visit family may be reduce some of the risk behaviors associated with HIV transmission. In addition, those migrants who are able to travel back and forth between the U.S. and Mexico likely represent migrants who have documents and are less vulnerable than undocumented migrants. At the same time, frequent visits to Mexico may also increase migrants' risk of exposure to TB infection.

There is extensive information and research reported in the literature on both TB and HIV in high-risk populations in Tijuana and San Diego. Less information is available in the literature on Ciudad Juarez and other border cities, and very little is present in the literature on the TB/HIV syndemic along other areas along the nearly 2,000 mile long U.S.-Mexico border. While much of the border runs through rural areas with extremely low population, there are still many cross-border sister cities throughout the region that have dynamic populations and may be host to the same risk factors for TB and HIV as those documented in the larger, well-studied cross-border metropolitan areas. In addition, traditional high risk groups for TB and HIV have been fairly well studied (sex workers, MSMs, IDUs) but less information is available on the TB/HIV syndemic in the general border population beyond these traditional risk groups. One of the few studies that looked at a non-high risk population (pregnant and delivering women in a Tijuana hospital) indicated that HIV prevalence was higher in that population than previously estimated (62). In a study of reported TB cases in San Diego, evidence that HIV co-infection disproportionately affects the general Hispanic population is another indication that the broader border population is

at increased risk from the TB/HIV syndemic (5), though the conclusions from this analysis did not find that Hispanic or border resident TB cases in Texas had a great odds of concurrent HIV infection compared to non-border or non-Hispanic Texas TB cases.

Limitations

This analysis used data collected by a surveillance system based on legally mandated notifiable disease reporting; TB is a notifiable disease in the state of Texas and any suspect or confirmed case of TB must be reported to the Texas Department of State Health Services (DSHS) (70).

While there are some components of the Texas TB surveillance programs that target high-risk populations in the state of Texas for TB screening and testing, the system is a passive reporting system that relies on notifications from healthcare providers, laboratories, schools, and others. This type of system likely does not identify all cases of TB in the state of Texas, particularly in those individuals who may seek treatment for their illness another state or country. In addition, completeness can be a problem when following up on TB case reports; many of the records contained in the dataset used in this analysis did not include complete information for all variables. Because of TB's lengthy and complicated treatment regimen, TB surveillance data is reported and counted by the county that will handle at least 90 days of the treatment of the TB case and not necessarily the county in which the case acquired TB (4). If a TB case is identified in Texas but is treated elsewhere, the case would not be counted for Texas. This may result in an under- or overestimate of TB disease in some areas of Texas depending on where individuals are diagnosed vs. treated.

The data used in this analysis is cross-sectional in nature, so it is impossible to draw conclusions about whether HIV infection preceded TB disease, which does not allow for any conclusions to be made about whether or not HIV could have contributed to the TB disease. In addition, these data are reporting data, not clinical data, and they do not capture TB patients who eventually test positive for HIV after their TB infection is reported to the state. Also, no data from the Texas HIV surveillance system was included in the report. Theoretically, any case of TB identified through routine testing of HIV patients should also have been reported to the state of Texas, but this analysis did not cross-reference the TB and HIV reporting systems in Texas to determine if such reporting was done through the HIV program and to link patients reported to both disease reporting systems. HIV positive patients diagnosed with TB may know their HIV status prior to their TB diagnosis but choose to withhold that information from TB care providers and decline further HIV testing. Patients may also have been diagnosed with HIV outside of Texas (in

another state or country) and may not divulge HIV status to the Texas TB program at the time of their TB diagnosis. As a result, an unknown quantity of prevalent HIV cases may not have been captured, and these data do not represent all cases of TB/HIV co-infection in Texas. Finally, in the dataset used for this analysis, there was no way to distinguish between HIV status based on lab report or self-report, or how advanced the HIV disease was in the patient at the time of TB diagnosis.

This analysis was susceptible to selection bias; almost a third (26.3%) of all the reported TB cases were excluded from the analysis due to unknown HIV status. In addition, the decision to use the 'gold standard' model including all variables resulted in the further elimination of observations missing some of these variables, which could also introduce selection bias. Finally, racial groups with small numbers (>80 records total) present in the Texas reporting data were excluded from the analysis as well; including them would have resulted in small number or zero cells and hampered the analysis. A number of variables that are particularly interesting in the context of studying TB infection in the border, including variables designating a case a binational case or capturing visa status upon arrival to the U.S., were not consistently collected over the study period and could not be included in this analysis. These factors all limit the conclusions that can be drawn from the study. Specifically, data on the HIV status of TB patients should be interpreted with caution because these data are not representative of all TB/HIV co-infection cases for the state of Texas.

Also, HIV testing may not be as widely available in some areas of Texas compared to others. Differential availability of HIV testing may impact some populations more than others. For example, if rural counties do not have the capacity to routinely test for HIV in TB patients and most of the population they see is Hispanic, this could have resulted in fewer Hispanic TB patients being tested for TB compared with African-Americans whose populations are primarily concentrated in urban areas of the state where HIV testing is routinely available.

Recommendations

A number of recommendations can be made despite the narrow conclusions and significant limitations of this study. First, given the high proportion of the population that had unknown HIV test status at the time of TB reporting, continued effort needs to be made to comprehensively test and report HIV status of all TB cases when they present for TB care, regardless of historical status as a risk group. This may be improved by increasing targeted,

culturally sensitive education to specific groups to increase acceptance of HIV testing. In addition, it may be beneficial to provide better education to all healthcare providers diagnosing, reporting, and treating TB in Texas on the need to routinely screen all patients for HIV regardless of perceived HIV risk. The use of IGRA testing over TST testing should be emphasized in foreign born patients who commonly receive the BCG vaccine and populations at high risk of not returning to the clinic to have their TSTs read (homeless, migrants, drug users). The benefits will be twofold; patients will not have to return to clinics for a follow-up reading of the TST, reducing the burden on clinics and the number of patients lost to follow up. In addition, detection of LTBI will be better in populations (such as the Mexican-born) with high-levels of BCG vaccination.

Also, efforts to improve cross-border and binational case management for patients with TB and TB/HIV should be continued and strengthened. In addition to traditional health partners on both sides of the border, Immigrations and Customs Enforcement medical staff should continue to be involved in TB case management of detainees in custody to ensure that patients with TB and TB/HIV are identified prior to release from ICE custody and to improve planning for case management and treatment during and after deportation.

While this analysis found that individuals with TB disease in the border region were less likely to have concurrent HIV infection, the true extent of the TB/HIV syndemic in the region is not known, and could not be determined by this investigation. This leaves room for further exploration of the TB/HIV syndemic in this dynamic and epidemiologically distinct region.

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Appendix A. Border Counties Defined by the 1983 La Paz Agreement



Figure 1. 32 Texas Border Counties

Map source: <http://www.dshs.state.tx.us/borderhealth/countyfacts.shtm>

In 1983, a federal designation was made to define “border” counties in U.S. It was called the “La Paz Agreement”. This agreement states that a “border” county is any county which touches the international mark or touches a county which touches the international mark. There are 32 Texas border counties defined under this agreement and are comprised of:

- Brewster
- Brooks
- Cameron
- Crockett
- Culberson
- Dimmit
- Duval
- Edwards
- El Paso
- Frio
- Hidalgo
- Hudspeth
- Jeff Davis
- Jim Hogg
- Kenedy
- Kinney
- La Salle
- Maverick
- McMullen
- Pecos
- Presidio
- Real
- Reeves
- Starr
- Sutton
- Terrell
- Uvalde
- Val Verde
- Webb
- Willacy
- Zapata
- Zavala

Appendix B. Texas TB Reporting Forms

TB 400A

**Texas Department of Health
Tuberculosis Elimination Division
Report of Case and Patient Services**

Date reported to health department: ____/____/____
Date form sent to region: ____/____/____
Date form sent to central office: ____/____/____

Initial Report Hospital Admission
 Address Change Name Change (show new name and draw single line through old) Other Change (please circle)

SSN _____ Medicaid # _____ ID# _____ DOB ____/____/____
MM DD YY

Name _____ AKA _____
(Last) (First) (Middle)

Street _____ Apt# _____ City _____ County _____ Zip Code _____ Patient's Tel# _____

Facility/Care Provider Name _____
Initial Reporting Source: Health Dept Private Physician Public Hospital VA Hospital Name of person completing this form: _____
 Military Hospital TDCJ Other (Specify) _____

Country of Birth If foreign born, Date of entry into U.S. ____/____/____	Notice of Arrival of Alien with TB Class <input type="checkbox"/> A <input type="checkbox"/> B1 <input type="checkbox"/> B2 <input type="checkbox"/> B3	Reported at Death <input type="checkbox"/> Yes <input type="checkbox"/> No If yes, Death Date ____/____/____ Was TB cause of death <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unk	Reported Out of State or Country <input type="checkbox"/> Yes Specify _____ <input type="checkbox"/> No
Preferred Language _____	ETHNICITY <input type="checkbox"/> Unknown <input type="checkbox"/> SEX <input type="checkbox"/> Hispanic or Latino <input type="checkbox"/> Male <input type="checkbox"/> Not Hispanic or Latino <input type="checkbox"/> Female		

RACE (check all that apply)
 White Native Hawaiian or Pacific Islander
 Black or African American American Indian or Alaskan Native
 Asian Unknown

OCCUPATION (within past 2 years)
 Unemployed during last 2 yrs Unknown
 Employed (If employed, check all that apply)
 Migrant/Seasonal Worker Student Retiree Child Disabled
 Health Care Worker (Specify) _____ Homemaker
 Correctional Emp Other Occupation Institutionalized

Resident of Correctional Facility at Time of Dx
If Yes: Federal Prison State Prison County Jail No City Jail Unknown Incarceration Date ____/____/____
 Juvenile Correctional Facility ICE Other

Resident of Long Term Care Facility at Time of Dx
If Yes: Nursing Home Yes No Unknown Hospital-Based Facility Residential Facility Mental Health Residential Facility
 Alcohol/Drug Treatment Facility Other Long Term Care Facility

Testing activities to find latent TB infections
 Patient referred, TB infection Project targeted testing Individual targeted testing Administrative: Not at risk for TB

POPULATION RISKS <input type="checkbox"/> Low Income <input type="checkbox"/> Inner-city resident <input type="checkbox"/> Foreign born <input type="checkbox"/> Binational (US-Mexico) <input type="checkbox"/> *Within past 2 years <input type="checkbox"/> Correctional employee* <input type="checkbox"/> Health care worker* <input type="checkbox"/> Prison/Jail Inmate* <input type="checkbox"/> Long-term facility for elderly/resident* <input type="checkbox"/> Health care facility/resident* <input type="checkbox"/> Shelter for homeless persons* <input type="checkbox"/> Migrant farm worker* <input type="checkbox"/> None of the above risks apply	MEDICAL RISKS <input type="checkbox"/> Diabetes mellitus <input type="checkbox"/> Alcohol Abuse (within past year) <input type="checkbox"/> Tobacco use <input type="checkbox"/> Siliocosis <input type="checkbox"/> Corticosteroids or other immunosuppressive therapy <input type="checkbox"/> Gastrectomy or jejunioileal bypass <input type="checkbox"/> Recent exposure to TB (Contact to TB case) <input type="checkbox"/> Contact to MDR-TB case <input type="checkbox"/> Weight at least 10% less than ideal body weight <input type="checkbox"/> Chronic malabsorption syndromes	<input type="checkbox"/> Leukemia <input type="checkbox"/> Lymphoma <input type="checkbox"/> Cancer of head <input type="checkbox"/> Cancer of neck <input type="checkbox"/> Drug abuse within past year: <input type="checkbox"/> Injecting <input type="checkbox"/> Non-injecting <input type="checkbox"/> Unknown if injecting <input type="checkbox"/> HIV seropositive (check only if laboratory confirmed) <input type="checkbox"/> Tuberculin skin test conversion within 2 years <input type="checkbox"/> Fibrotic lesions (on chest x-ray) consistent with old, healed TB
		<input type="checkbox"/> Chronic renal failure <input type="checkbox"/> Organ Transplant <input type="checkbox"/> Other _____ <input type="checkbox"/> None of these medical risks apply

HIV TEST RESULTS
Date HIV Test ____/____/____
 Positive Negative
 Pending Refused
 Not Offered
Date CD4 Count ____/____/____
Results CD4 Count _____

TUBERCULIN SKIN TEST Documented history of positive TST? Yes No
____/____/____ mm Positive Negative Not Read
____/____/____ mm Positive Negative Not Read

PRIOR LTBI TREATMENT Yes No
Start Date ____/____/____
Stop Date ____/____/____

FOR TREATMENT OF LTBI ONLY
DOPT: Yes, totally observed No, self-administered Both
DOPT site: Clinic or medical facility Field
Frequency: Daily Twice Weekly Three X's Weekly
Date Normal Chest X-ray ____/____/____ Weight _____ Height _____
AT5 Classification
 0 No M. TB Exposure, Not TB Infected
 1 M. TB Exposure, No Evidence of TB Infection
 2 M. TB Infection, No Disease
 4 M. TB, No Current Disease
Date Regimen Start ____/____/____ Date Regimen Stop ____/____/____
Date Restart ____/____/____ Date Regimen Stop ____/____/____
 Isoniazid _____ mgs Other (specify) _____ mgs
 Rifampin _____ mgs Other (specify) _____ mgs
 B6 _____ mgs Prescribed for: _____ months Maximum refills authorized: _____
Physician Signature _____ Date _____

CLOSURE: Date ____/____/____
 Completion adequate therapy _____ # months on Rx _____ # months recommended
 Lost to followup Patient chose to stop Deceased (Cause) _____
 Adverse Drug Reaction Moved out of state/country to: _____
Provider decision: Pregnant Non-TB Other: _____

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Texas Department of Health
TB-400A (11/03)

**Texas Department of Health
Tuberculosis Elimination Division
Report of Case and Patient Services**

Date reported to health department: ____/____/____
Date form sent to region: ____/____/____
Date form sent to central office: ____/____/____

Initial Report Drug Resistance Followup or Medical Review Hospital Admission or Discharge

Name _____ (Last) _____ (First) _____ (Middle) DOB ____/____/____
MM DD YY

Street _____ Apt# _____ City _____ County _____ Zip Code _____
Facility/Care Provider Name _____
Facility responsible for patient care: Public Health Clinic Private Physician Hospital
 Other (Specify) _____ Name of person completing this form _____

Signs/Symptoms at DX Fever <input type="checkbox"/> Y <input type="checkbox"/> N Chills <input type="checkbox"/> Y <input type="checkbox"/> N Cough <input type="checkbox"/> Y <input type="checkbox"/> N Productive Cough <input type="checkbox"/> Y <input type="checkbox"/> N Hemoptysis <input type="checkbox"/> Y <input type="checkbox"/> N Night Sweats <input type="checkbox"/> Y <input type="checkbox"/> N Weight Loss (> 10%) <input type="checkbox"/> Y <input type="checkbox"/> N Other: _____	Chest X-Ray Date: ____/____/____ Results: <input type="checkbox"/> Normal <input type="checkbox"/> Abnormal <input type="checkbox"/> Not Done <input type="checkbox"/> Unk If Abnormal, check abnormality: _____ <input type="checkbox"/> Cavitary Status: <input type="checkbox"/> Stable <input type="checkbox"/> Non-cavitary, consistent with TB <input type="checkbox"/> Worsening <input type="checkbox"/> Non-cavitary, not consistent with TB <input type="checkbox"/> Improving Comments: _____ <input type="checkbox"/> Unknown	If Pediatric TB Case (<15 Years Old) Country of birth for primary guardians: Guardian 1) _____ Guardian 2) _____ Patient lived outside US for > 3 months <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown If yes, Country: _____
--	--	---

Status: New Recurrent Reopen
 Prior Therapy: Yes No Start Date: ____/____/____ Stop Date: ____/____/____

ATIS Classification

<input type="checkbox"/> 0	No M. TB Exposure, Not TB Infected
<input type="checkbox"/> 1	M. TB Exposure, No Evidence of TB Infection
<input type="checkbox"/> 2	M. TB Infection, No Disease
<input type="checkbox"/> 3	M. TB Infection, Current Disease
<input type="checkbox"/> 4	M. TB, No Current Disease
<input type="checkbox"/> 5	M. TB Suspect, Diagnosis Pending

Predominant Site: (Class 3, 4)

Significant Sites other than Predominant

<input type="checkbox"/> 00	Pulmonary	30	<input type="checkbox"/>	Bone and/or Joint
<input type="checkbox"/> 10	Pleural	40	<input type="checkbox"/>	Genitourinary
<input type="checkbox"/> 20	Lymphatic	50	<input type="checkbox"/>	Miliary/Disseminated
<input type="checkbox"/> 21	Cervical	60	<input type="checkbox"/>	Meningeal
<input type="checkbox"/> 22	Intrathoracic	70	<input type="checkbox"/>	Peritoneal
<input type="checkbox"/> 23	Other	80	<input type="checkbox"/>	Other (Specify) _____

AFB Smear Results
 Current: ____/____/____ Negative Positive
 Pending Not done
 Specimen type: sputum urine bronchial washing
 biopsy other
 If biopsy or other, list anatomic site of specimen: _____
 If other than sputa, type of exam: _____
 Collection date of initial positive AFB smear: ____/____/____
 Collection date of first consistently negative AFB smear: ____/____/____

Nucleic Acid Amplification Test
 Current: ____/____/____ Negative Positive
 Indeterminate Not done

Culture Results
 Current: ____/____/____ Negative Positive for M. TB
 Positive for Non-M. TB Pending Not done
 Specimen type: sputum urine bronchial washing
 biopsy other
 If biopsy or other, list anatomic site of specimen: _____
 Collection date of initial positive MTB culture: ____/____/____
 Collection date of first consistently negative MTB culture: ____/____/____
 Sputum culture conversion documented? Yes No NA
 If no, then reason: _____

Susceptibility Results
 Date initial susceptibility culture was collected: ____/____/____
 Initial culture was resistant to: Isoniazid Rifampin Ethambutol
 Date last positive culture was collected: ____/____/____
 Last culture was resistant to: Isoniazid Rifampin Ethambutol
 Other quinolone(s) _____
 Other(s) _____

Reason Therapy Extending > 12 months: _____
 Hospitalization Advised: Yes No Control Order: ____/____/____
 Quarantine Advised: Yes No Court Action: ____/____/____
 Return for chest x-ray: ____/____/____ Compliant: Yes No
 Collect next sputum on: ____/____/____ Other lab studies: ____/____/____
 Return to MD clinic on: ____/____/____
 Return to Nurse clinic on: ____/____/____

Prescribed for: _____ months Maximum refills authorized: _____
 Closure: _____
 Date: ____/____/____ % doses taken by DOT: _____
 # doses taken: _____ # doses recommended: _____
 # months on Rx: _____ # months recommended: _____

Completion of adequate therapy Lost to followup
 Patient chose to stop Adverse drug reaction
 Deceased (Cause) _____
 Moved out of state/country to: _____
 Date referral sent to Austin: ____/____/____
 Provider decision: Pregnant Non-TB Other: _____

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 Texas Department of Health
 TB-400B (11/03)

Appendix C. Trends in HIV Status

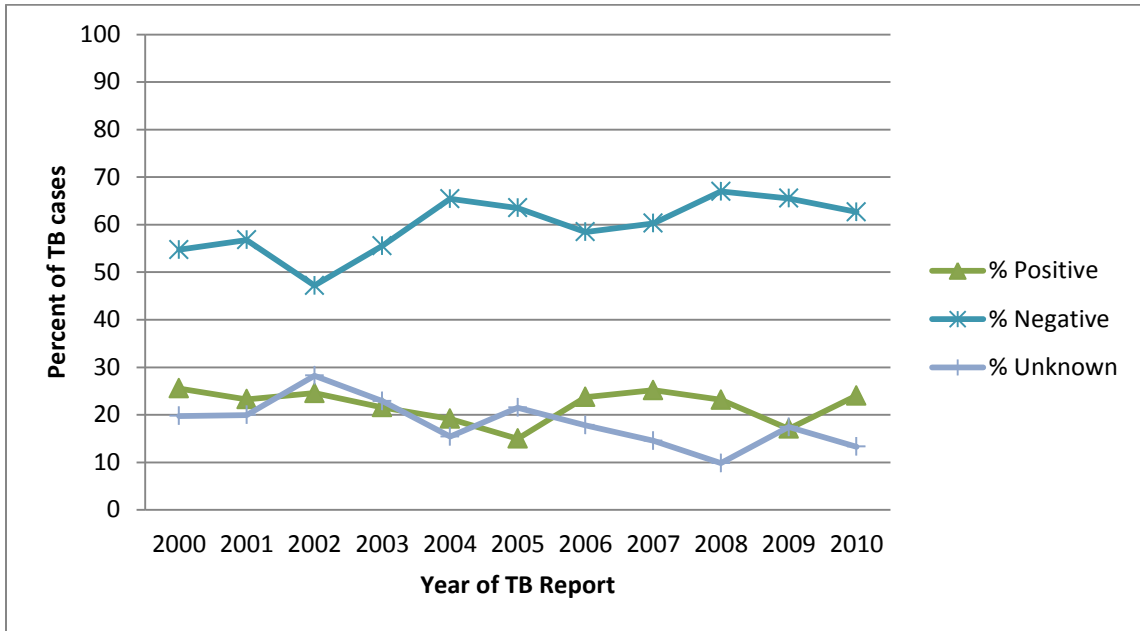


Figure 2. Change in Reported HIV Status Proportions in African American TB Cases in Texas, 2000-2010

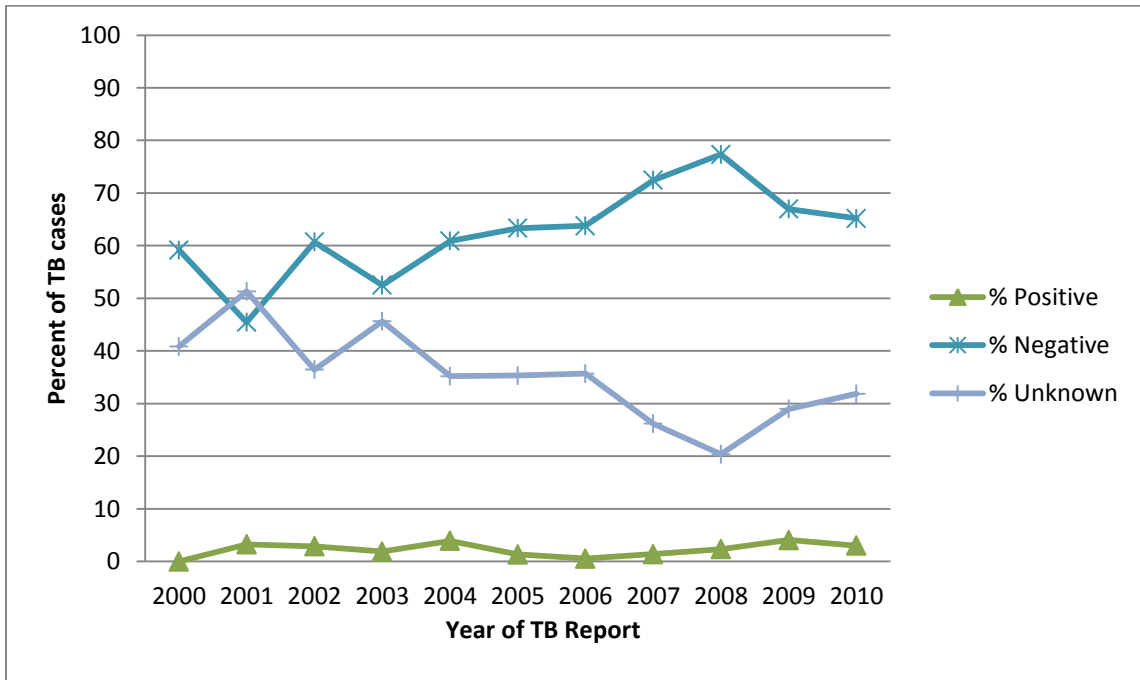


Figure 3. Change in Reported HIV Status Proportions in Asian TB Cases in Texas, 2000-2010

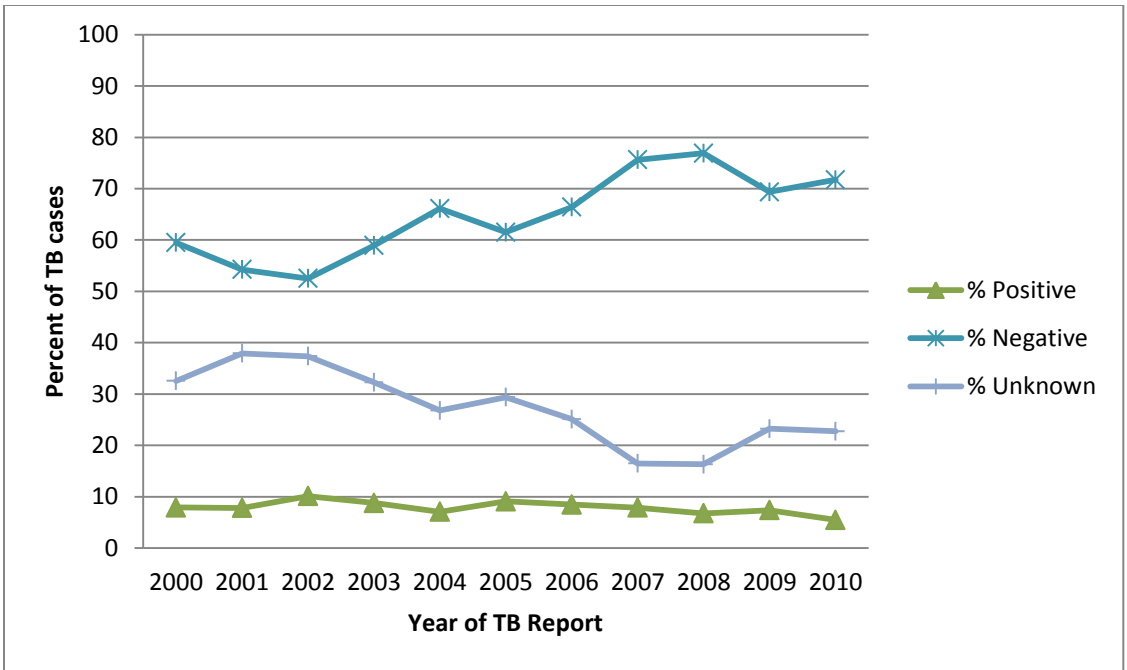


Figure 4. Change in Reported HIV Status Proportions Caucasian TB Cases in Texas, 2000-2010

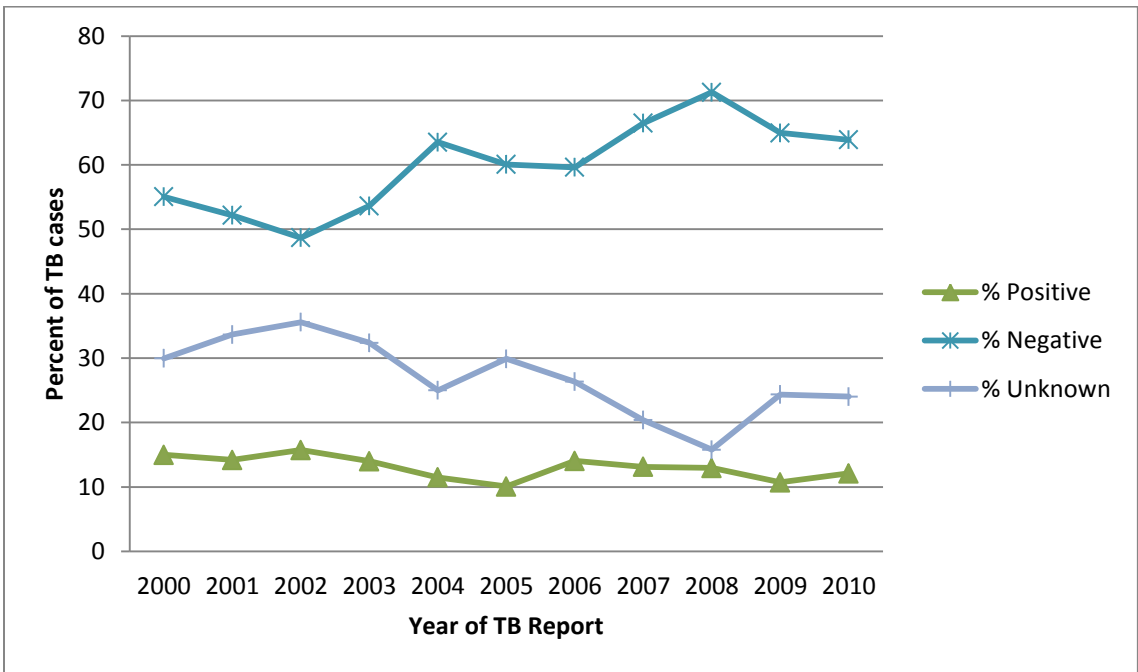


Figure 5. Change in Reported HIV Status Proportions in Non-Hispanic TB Cases in Texas, 2000-2010

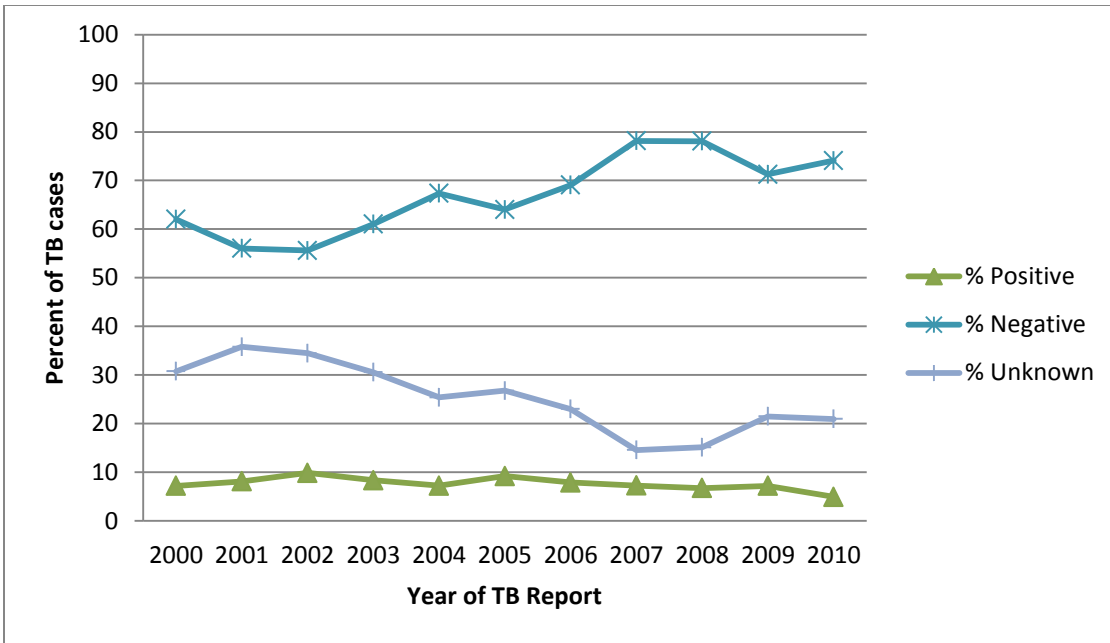


Figure 6. Change in Reported HIV Status Proportions in Hispanic TB Cases in Texas, 2000-2010

Appendix D. SAS Output from Analysis

*This model is my first logistic regression model containing all two-way interaction terms identified in the previous bivariate analyses. In addition, all variables used in the analysis (significant or not in the univariate and bivariate analyses) are included in this model. I used this model to start a backwards elimination of non-significant variables, starting with the two-way interaction terms. Once I removed all non-significant interaction terms I assessed the goodness-of-fit of the model using the chi-square from the Hosmer-Lemeshow test.

Step 1 Full Model with Interaction Terms

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The LOGISTIC Procedure

Model Information

Data Set	TBHIV.LOGISTIC2
Response Variable	HIV
Number of Response Levels	2
Model	binary logit
Optimization Technique	Fisher's scoring

Number of Observations Read	11282
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Number of Observations Used	10989
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Response Profile

Ordered		Total
Value	HIV	Frequency

1	0	9412
2	1	1577

Probability modeled is HIV=1.

NOTE: 293 observations were deleted due to missing values for the response or explanatory variables.

Class Level Information

Class	Value	Design Variables								
FB	0	0								
	1	1								
year_counted 0 0 0 0 0 0 0 0	2000	1	0	0	0	0	0	0	0	0
	2001	0	1	0	0	0	0	0	0	0
	2002	0	0	1	0	0	0	0	0	0
	2003	0	0	0	1	0	0	0	0	0
	2004	0	0	0	0	1	0	0	0	0
	2005	0	0	0	0	0	1	0	0	0
	2006	0	0	0	0	0	0	1	0	0
	2007	0	0	0	0	0	0	0	1	0

0	2008	0	0	0	0	0	0	0	0	1
1	2009	0	0	0	0	0	0	0	0	0
0	2010	0	0	0	0	0	0	0	0	0
age1	25	1	0	0	0					
	35	0	1	0	0					
	45	0	0	1	0					
	55	0	0	0	1					

The LOGISTIC Procedure

Class Level Information

Class	Value	Design Variables				
	65	0	0	0	0	0
USYEARS	0	1	0	0	0	0
	1	0	1	0	0	0
	2	0	0	1	0	0
	7	0	0	0	1	0
	15	0	0	0	0	1
	20	0	0	0	0	0
race2	Asian	1	0			
	Black	0	1			
	White	0	0			
location	1	0	0			
	2	1	0			
	3	0	1			
Occupation	0	1				
	1	0				

Ethnicity	0	0
	1	1

Model Convergence Status

Convergence criterion (GCONV=1E-8) satisfied.

Model Fit Statistics

Criterion	Intercept	Intercept
	Only	and Covariates
AIC	9041.095	7482.697
SC	9048.400	7796.797
-2 Log L	9039.095	7396.697

Step 1 Full Model with Interaction Terms

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The LOGISTIC Procedure

Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	1642.3983	42	<.0001
Score	1585.6167	42	<.0001
Wald	1224.6354	42	<.0001

Type 3 Analysis of Effects

Effect	DF	Wald	
		Chi-Square	Pr > ChiSq
BORDER	1	17.2056	<.0001
FB	1	2.2380	0.1347
age1	4	400.6157	<.0001
year_counted	10	36.7572	<.0001
sex	1	118.7751	<.0001
Ethnicity	1	2.2346	0.1350
race2	2	199.1764	<.0001
IDU	1	42.0527	<.0001
USYEARS	4	4.0463	0.3998
NoInject	1	22.2425	<.0001

alcohol	1	33.0347	<.0001
homeless	1	0.3319	0.5645
prison	1	5.3360	0.0209
LTCRes	1	0.4329	0.5106
location	2	201.2183	<.0001
Occupation	1	105.5128	<.0001
BORDER*FB	1	0.0773	0.7811
BORDER*alcohol	1	3.2464	0.0716
BORDER*USYEARS	4	3.4812	0.4807
BORDER*prison	1	0.9764	0.3231
BORDER*location	2	5.4910	0.0642

*Based on these results, I decided to drop Border*FB (foreign birth) because this was the least significant interaction term in the model(p= 0.78).

NOTE: The following parameters have been set to 0, since the variables are a linear combination

of other variables as shown.

$$\text{USYEARS0} = \text{Intercept} - \text{FB1}$$

$$\text{USYEARS0BORDER} = \text{BORDER} - \text{FB1BORDER}$$

Step 1 Full Model with Interaction Terms

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The LOGISTIC Procedure

Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept		1	-5.0511	0.2400	443.1074	<.0001
BORDER		1	-0.9238	0.2227	17.2056	<.0001
FB	1	1	0.2155	0.1440	2.2380	0.1347
age1	25	1	1.3181	0.1926	46.8570	<.0001
age1	35	1	2.5355	0.1829	192.1593	<.0001
age1	45	1	2.1824	0.1816	144.3766	<.0001
age1	55	1	1.2871	0.1851	48.3370	<.0001
year_counted	2000	1	0.2937	0.1558	3.5509	0.0595
year_counted	2001	1	0.3560	0.1532	5.4011	0.0201
year_counted	2002	1	0.6301	0.1522	17.1329	<.0001
year_counted	2003	1	0.3603	0.1539	5.4800	0.0192
year_counted	2004	1	0.0504	0.1550	0.1059	0.7448
year_counted	2005	1	0.1067	0.1585	0.4531	0.5009
year_counted	2006	1	0.2681	0.1541	3.0284	0.0818
year_counted	2007	1	0.2364	0.1557	2.3038	0.1291
year_counted	2008	1	0.0276	0.1563	0.0313	0.8596
year_counted	2009	1	0.1023	0.1645	0.3870	0.5339
sex		1	0.8064	0.0740	118.7751	<.0001
Ethnicity	1	1	-0.1621	0.1084	2.2346	0.1350

race2	Asian	1	-1.5195	0.1913	63.1165	<.0001
race2	Black	1	0.7113	0.0915	60.5022	<.0001
IDU		1	0.8598	0.1326	42.0527	<.0001
USYEARS	0	0	0	.	.	.
USYEARS	1	1	-0.2226	0.1882	1.3998	0.2368
USYEARS	2	1	-0.2980	0.1663	3.2092	0.0732
USYEARS	7	1	-0.2191	0.1813	1.4601	0.2269
USYEARS	15	1	-0.3051	0.1736	3.0901	0.0788
NoInject		1	0.4508	0.0956	22.2425	<.0001
alcohol		1	-0.4825	0.0840	33.0347	<.0001
homeless		1	-0.0635	0.1101	0.3319	0.5645
prison		1	-0.2773	0.1201	5.3360	0.0209
LTCRes		1	0.1485	0.2257	0.4329	0.5106
location	2	1	0.8337	0.0768	117.9539	<.0001
location	3	1	1.1884	0.1065	124.4833	<.0001
Occupation	0	1	0.6712	0.0653	105.5128	<.0001
BORDER*FB	1	1	0.0805	0.2898	0.0773	0.7811
BORDER*alcohol		1	0.3739	0.2075	3.2464	0.0716
BORDER*USYEARS	0	0	0	.	.	.
BORDER*USYEARS	1	1	0.1375	0.4111	0.1118	0.7381
BORDER*USYEARS	2	1	0.3528	0.3696	0.9111	0.3398
BORDER*USYEARS	7	1	0.5216	0.4046	1.6619	0.1973
BORDER*USYEARS	15	1	0.5720	0.3515	2.6487	0.1036
BORDER*prison		1	-0.3279	0.3318	0.9764	0.3231
BORDER*location	2	1	0.5422	0.2314	5.4909	0.0191
BORDER*location	3	1	0.1704	0.3250	0.2750	0.6000

Step 1 Full Model with Interaction Terms

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The LOGISTIC Procedure

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
age1	25 vs 65	3.736	2.562	5.449
age1	35 vs 65	12.623	8.820	18.066
age1	45 vs 65	8.868	6.212	12.659
age1	55 vs 65	3.622	2.520	5.206
year_counted	2000 vs 2010	1.341	0.988	1.820
year_counted	2001 vs 2010	1.428	1.057	1.927
year_counted	2002 vs 2010	1.878	1.393	2.531
year_counted	2003 vs 2010	1.434	1.060	1.939
year_counted	2004 vs 2010	1.052	0.776	1.425
year_counted	2005 vs 2010	1.113	0.815	1.518
year_counted	2006 vs 2010	1.308	0.967	1.769
year_counted	2007 vs 2010	1.267	0.933	1.719
year_counted	2008 vs 2010	1.028	0.757	1.396
year_counted	2009 vs 2010	1.108	0.802	1.529
sex		2.240	1.937	2.589
Ethnicity	1 vs 0	0.850	0.688	1.052
race2	Asian vs White	0.219	0.150	0.318
race2	Black vs White	2.037	1.702	2.437
IDU		2.363	1.822	3.064

NoInject		1.570	1.301	1.893
homeless		0.939	0.756	1.165
LTCRes		1.160	0.745	1.805
Occupation	0 vs 1	1.957	1.721	2.224

Association of Predicted Probabilities and Observed Responses

Percent Concordant	79.5	Somers' D	0.593
Percent Discordant	20.2	Gamma	0.595
Percent Tied	0.4	Tau-a	0.146
Pairs	14842724	c	0.796

Partition for the Hosmer and Lemeshow Test

Group	Total	HIV = 1		HIV = 0	
		Observed	Expected	Observed	Expected
1	1099	18	12.62	1081	1086.38
2	1102	29	28.28	1073	1073.72
3	1100	47	44.33	1053	1055.67
4	1099	59	62.43	1040	1036.57
5	1099	69	87.24	1030	1011.76
6	1099	128	118.14	971	980.86
7	1098	148	159.48	950	938.52

Step 1 Full Model with Interaction Terms

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The LOGISTIC Procedure

Partition for the Hosmer and Lemeshow Test

Group	Total	HIV = 1		HIV = 0	
		Observed	Expected	Observed	Expected
8	1102	226	221.18	876	880.82
9	1098	319	317.13	779	780.87
10	1093	534	526.17	559	566.83

Hosmer and Lemeshow Goodness-of-Fit Test

Chi-Square	DF	Pr > ChiSq
9.1101	8	0.3331

*The chi-square test for fit is not significant, so this model fits the data.

The LOGISTIC Procedure

Model Information

Data Set	TBHIV.LOGISTIC2
Response Variable	HIV
Number of Response Levels	2
Model	binary logit
Optimization Technique	Fisher's scoring

Number of Observations Read	11282
Number of Observations Used	10989

Response Profile

Ordered Value	HIV	Total Frequency
1	0	9412
2	1	1577

Probability modeled is HIV=1.

NOTE: 293 observations were deleted due to missing values for the response or explanatory variables.

Class Level Information

Class	Value	Design Variables								
FB	0	0								
	1	1								
year_counted 0	2000	1	0	0	0	0	0	0	0	0
	2001	0	1	0	0	0	0	0	0	0
	2002	0	0	1	0	0	0	0	0	0
	2003	0	0	0	1	0	0	0	0	0
	2004	0	0	0	0	1	0	0	0	0
	2005	0	0	0	0	0	1	0	0	0
	2006	0	0	0	0	0	0	1	0	0
	2007	0	0	0	0	0	0	0	1	0
	2008	0	0	0	0	0	0	0	0	1
	2009	0	0	0	0	0	0	0	0	0
1	2010	0	0	0	0	0	0	0	0	0
age1	25	1	0	0	0					
	35	0	1	0	0					
	45	0	0	1	0					

55	0	0	0	1
65	0	0	0	0

The LOGISTIC Procedure

Class Level Information

Class	Value	Design Variables				
USYEARS	0	1	0	0	0	0
	1	0	1	0	0	0
	2	0	0	1	0	0
	7	0	0	0	1	0
	15	0	0	0	0	1
	20	0	0	0	0	0
race2	Asian	1	0			
	Black	0	1			
	White	0	0			
location	1	0	0			
	2	1	0			
	3	0	1			
Occupation	0	1				
	1	0				
Ethnicity	0	0				
	1	1				

Model Convergence Status

Convergence criterion (GCONV=1E-8) satisfied.

Model Fit Statistics

Criterion	Intercept	Intercept
	Only	and Covariates
AIC	9041.095	7482.697
SC	9048.400	7796.797
-2 Log L	9039.095	7396.697

Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	1642.3983	42	<.0001
Score	1585.6167	42	<.0001
Wald	1224.6354	42	<.0001

The LOGISTIC Procedure

Type 3 Analysis of Effects

Effect	DF	Wald	
		Chi-Square	Pr > ChiSq
BORDER	1	10.5998	0.0011
FB	1	2.2380	0.1347
age1	4	400.6157	<.0001
year_counted	10	36.7572	<.0001
sex	1	118.7751	<.0001
Ethnicity	1	2.2346	0.1350
race2	2	199.1764	<.0001
IDU	1	42.0527	<.0001
USYEARS	4	4.0463	0.3998
NoInject	1	22.2425	<.0001
alcohol	1	33.0347	<.0001
homeless	1	0.3319	0.5645
prison	1	5.3360	0.0209
LTCRes	1	0.4329	0.5106
location	2	201.2183	<.0001
Occupation	1	105.5128	<.0001
BORDER*alcohol	1	3.2464	0.0716
BORDER*USYEARS	5	6.2207	0.2853
BORDER*prison	1	0.9764	0.3231
BORDER*location	2	5.4910	0.0642

*Based on these results, I decided to drop Border*prison because this was the least significant interaction term remaining in the model(p= 0.32).

NOTE: The following parameters have been set to 0, since the variables are a linear combination

of other variables as shown.

$$\text{USYEARS0} = \text{Intercept} - \text{FB1}$$

Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept		1	-5.0511	0.2400	443.1074	<.0001
BORDER		1	-0.8432	0.2590	10.5998	0.0011
FB	1	1	0.2155	0.1440	2.2380	0.1347
age1	25	1	1.3181	0.1926	46.8570	<.0001
age1	35	1	2.5355	0.1829	192.1593	<.0001
age1	45	1	2.1824	0.1816	144.3766	<.0001
age1	55	1	1.2871	0.1851	48.3370	<.0001
year_counted	2000	1	0.2937	0.1558	3.5509	0.0595
year_counted	2001	1	0.3560	0.1532	5.4011	0.0201
year_counted	2002	1	0.6301	0.1522	17.1329	<.0001
year_counted	2003	1	0.3603	0.1539	5.4800	0.0192
year_counted	2004	1	0.0504	0.1550	0.1059	0.7448

The LOGISTIC Procedure

Analysis of Maximum Likelihood Estimates

Parameter	DF	Estimate	Standard	Wald	Pr > ChiSq	
			Error	Chi-Square		
year_counted	2005	1	0.1067	0.1585	0.4531	0.5009
year_counted	2006	1	0.2681	0.1541	3.0284	0.0818
year_counted	2007	1	0.2364	0.1557	2.3038	0.1291
year_counted	2008	1	0.0276	0.1563	0.0313	0.8596
year_counted	2009	1	0.1023	0.1645	0.3870	0.5339
sex		1	0.8064	0.0740	118.7751	<.0001
Ethnicity	1	1	-0.1621	0.1084	2.2346	0.1350
race2	Asian	1	-1.5195	0.1913	63.1165	<.0001
race2	Black	1	0.7113	0.0915	60.5022	<.0001
IDU		1	0.8598	0.1326	42.0527	<.0001
USYEARS	0	0	0	.	.	.
USYEARS	1	1	-0.2226	0.1882	1.3998	0.2368
USYEARS	2	1	-0.2980	0.1663	3.2092	0.0732
USYEARS	7	1	-0.2191	0.1813	1.4601	0.2269
USYEARS	15	1	-0.3051	0.1736	3.0901	0.0788
NoInject		1	0.4508	0.0956	22.2425	<.0001
alcohol		1	-0.4825	0.0840	33.0347	<.0001
homeless		1	-0.0635	0.1101	0.3319	0.5645
prison		1	-0.2773	0.1201	5.3360	0.0209
LTCRes		1	0.1485	0.2257	0.4329	0.5106

location	2	1	0.8337	0.0768	117.9539	<.0001
location	3	1	1.1884	0.1065	124.4833	<.0001
Occupation	0	1	0.6712	0.0653	105.5128	<.0001
BORDER*alcohol		1	0.3739	0.2075	3.2464	0.0716
BORDER*USYEARS	0	1	-0.0805	0.2898	0.0773	0.7811
BORDER*USYEARS	1	1	0.1375	0.4111	0.1118	0.7381
BORDER*USYEARS	2	1	0.3528	0.3696	0.9111	0.3398
BORDER*USYEARS	7	1	0.5216	0.4046	1.6619	0.1973
BORDER*USYEARS	15	1	0.5720	0.3515	2.6487	0.1036
BORDER*prison		1	-0.3279	0.3318	0.9764	0.3231
BORDER*location	2	1	0.5422	0.2314	5.4909	0.0191
BORDER*location	3	1	0.1704	0.3250	0.2750	0.6000

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
FB	1 vs 0	1.240	0.935	1.645
age1	25 vs 65	3.736	2.562	5.449
age1	35 vs 65	12.623	8.820	18.066
age1	45 vs 65	8.868	6.212	12.659
age1	55 vs 65	3.622	2.520	5.206
year_counted	2000 vs 2010	1.341	0.988	1.820
year_counted	2001 vs 2010	1.428	1.057	1.927

The LOGISTIC Procedure

Odds Ratio Estimates

Effect	Point Estimate	95% Wald Confidence Limits	
year_counted 2002 vs 2010	1.878	1.393	2.531
year_counted 2003 vs 2010	1.434	1.060	1.939
year_counted 2004 vs 2010	1.052	0.776	1.425
year_counted 2005 vs 2010	1.113	0.815	1.518
year_counted 2006 vs 2010	1.308	0.967	1.769
year_counted 2007 vs 2010	1.267	0.933	1.719
year_counted 2008 vs 2010	1.028	0.757	1.396
year_counted 2009 vs 2010	1.108	0.802	1.529
sex	2.240	1.937	2.589
Ethnicity 1 vs 0	0.850	0.688	1.052
race2 Asian vs White	0.219	0.150	0.318
race2 Black vs White	2.037	1.702	2.437
IDU	2.363	1.822	3.064
NoInject	1.570	1.301	1.893
homeless	0.939	0.756	1.165
LTCRes	1.160	0.745	1.805
Occupation 0 vs 1	1.957	1.721	2.224

Association of Predicted Probabilities and Observed Responses

Percent Concordant	79.5	Somers' D	0.593
Percent Discordant	20.2	Gamma	0.595
Percent Tied	0.4	Tau-a	0.146
Pairs	14842724	c	0.796

Partition for the Hosmer and Lemeshow Test

Group	Total	HIV = 1		HIV = 0	
		Observed	Expected	Observed	Expected
1	1099	18	12.62	1081	1086.38
2	1102	29	28.28	1073	1073.72
3	1100	47	44.33	1053	1055.67
4	1099	59	62.43	1040	1036.57
5	1099	69	87.24	1030	1011.76
6	1099	128	118.14	971	980.86
7	1098	148	159.48	950	938.52
8	1102	226	221.18	876	880.82
9	1098	319	317.13	779	780.87
10	1093	534	526.17	559	566.83

The LOGISTIC Procedure

Hosmer and Lemeshow Goodness-of-Fit Test

Chi-Square	DF	Pr > ChiSq
9.1101	8	0.3331

*Goodness-of-Fit test still shows that this model fits my data.

Step 3 * Prison*border removed from the model.

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The LOGISTIC Procedure

Model Information

Data Set	TBHIV.LOGISTIC2
Response Variable	HIV
Number of Response Levels	2
Model	binary logit
Optimization Technique	Fisher's scoring

Number of Observations Read	11282
Number of Observations Used	10989

Response Profile

Ordered Value	HIV	Total Frequency
1	0	9412
2	1	1577

Probability modeled is HIV=1.

NOTE: 293 observations were deleted due to missing values for the response or explanatory variables.

Class Level Information

Class	Value	Design Variables								
FB	0	0								
	1	1								
year_counted 0	2000	1	0	0	0	0	0	0	0	0
	2001	0	1	0	0	0	0	0	0	0
	2002	0	0	1	0	0	0	0	0	0
	2003	0	0	0	1	0	0	0	0	0
	2004	0	0	0	0	1	0	0	0	0
	2005	0	0	0	0	0	1	0	0	0
	2006	0	0	0	0	0	0	1	0	0
	2007	0	0	0	0	0	0	0	1	0
	2008	0	0	0	0	0	0	0	0	1
	2009	0	0	0	0	0	0	0	0	0
1	2010	0	0	0	0	0	0	0	0	0
age1	25	1	0	0	0					
	35	0	1	0	0					
	45	0	0	1	0					

55	0	0	0	1
65	0	0	0	0

The LOGISTIC Procedure

Class Level Information

Class	Value	Design Variables				
USYEARS	0	1	0	0	0	0
	1	0	1	0	0	0
	2	0	0	1	0	0
	7	0	0	0	1	0
	15	0	0	0	0	1
	20	0	0	0	0	0
race2	Asian	1	0			
	Black	0	1			
	White	0	0			
location	1	0	0			
	2	1	0			
	3	0	1			
Occupation	0	1				
	1	0				
Ethnicity	0	0				
	1	1				

Model Convergence Status

Convergence criterion (GCONV=1E-8) satisfied.

Model Fit Statistics

Criterion	Intercept	Intercept
	Only	and Covariates
AIC	9041.095	7481.699
SC	9048.400	7788.494
-2 Log L	9039.095	7397.699

Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	1641.3960	41	<.0001
Score	1583.3554	41	<.0001
Wald	1223.5380	41	<.0001

The LOGISTIC Procedure

Type 3 Analysis of Effects

Effect	DF	Wald	
		Chi-Square	Pr > ChiSq
BORDER	1	11.7055	0.0006
FB	1	2.1520	0.1424
age1	4	400.8396	<.0001
year_counted	10	36.9769	<.0001
sex	1	118.8056	<.0001
Ethnicity	1	2.1587	0.1418
race2	2	199.5249	<.0001
IDU	1	41.5251	<.0001
USYEARS	4	4.0169	0.4037
NoInject	1	22.2336	<.0001
alcohol	1	33.1815	<.0001
homeless	1	0.3305	0.5654
prison	1	8.0599	0.0045
LTCRes	1	0.4253	0.5143
location	2	200.8913	<.0001
Occupation	1	106.5139	<.0001
BORDER*alcohol	1	3.2300	0.0723
BORDER*USYEARS	5	6.1534	0.2916
BORDER*location	2	6.3196	0.0424

*based on these results I decided to drop USYEARS from the model because it was the least significant interaction term left (p-value= 0.29).

NOTE: The following parameters have been set to 0, since the variables are a linear combination

of other variables as shown.

$$\text{USYEARS}_0 = \text{Intercept} - \text{FB1}$$

Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard	Wald	Pr > ChiSq
				Error	Chi-Square	
Intercept		1	-5.0486	0.2400	442.6620	<.0001
BORDER		1	-0.8800	0.2572	11.7055	0.0006
FB	1	1	0.2112	0.1439	2.1520	0.1424
age1	25	1	1.3158	0.1926	46.6894	<.0001
age1	35	1	2.5344	0.1829	191.9536	<.0001
age1	45	1	2.1821	0.1816	144.3139	<.0001
age1	55	1	1.2861	0.1851	48.2560	<.0001
year_counted	2000	1	0.2960	0.1558	3.6070	0.0575
year_counted	2001	1	0.3558	0.1532	5.3926	0.0202
year_counted	2002	1	0.6327	0.1522	17.2725	<.0001
year_counted	2003	1	0.3601	0.1540	5.4701	0.0193
year_counted	2004	1	0.0519	0.1550	0.1119	0.7380
year_counted	2005	1	0.1071	0.1585	0.4566	0.4992

The LOGISTIC Procedure

Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
year_counted	2006	1	0.2666	0.1541	2.9939	0.0836
year_counted	2007	1	0.2375	0.1558	2.3248	0.1273
year_counted	2008	1	0.0265	0.1563	0.0288	0.8653
year_counted	2009	1	0.1035	0.1646	0.3951	0.5296
sex		1	0.8067	0.0740	118.8056	<.0001
Ethnicity	1	1	-0.1592	0.1083	2.1587	0.1418
race2	Asian	1	-1.5191	0.1913	63.0836	<.0001
race2	Black	1	0.7127	0.0914	60.7680	<.0001
IDU		1	0.8520	0.1322	41.5251	<.0001
USYEARS	0	0	0	.	.	.
USYEARS	1	1	-0.2214	0.1882	1.3842	0.2394
USYEARS	2	1	-0.2966	0.1663	3.1797	0.0746
USYEARS	7	1	-0.2173	0.1813	1.4366	0.2307
USYEARS	15	1	-0.3043	0.1736	3.0739	0.0796
NoInject		1	0.4507	0.0956	22.2336	<.0001
alcohol		1	-0.4835	0.0839	33.1815	<.0001
homeless		1	-0.0633	0.1102	0.3305	0.5654
prison		1	-0.3202	0.1128	8.0599	0.0045
LTCRes		1	0.1473	0.2258	0.4253	0.5143
location	2	1	0.8331	0.0767	117.8244	<.0001

location	3	1	1.1866	0.1065	124.1841	<.0001
Occupation	0	1	0.6738	0.0653	106.5139	<.0001
BORDER*alcohol		1	0.3733	0.2077	3.2300	0.0723
BORDER*USYEARS	0	1	-0.0739	0.2899	0.0650	0.7988
BORDER*USYEARS	1	1	-0.00581	0.3862	0.0002	0.9880
BORDER*USYEARS	2	1	0.3146	0.3681	0.7306	0.3927
BORDER*USYEARS	7	1	0.5128	0.4051	1.6027	0.2055
BORDER*USYEARS	15	1	0.5688	0.3518	2.6142	0.1059
BORDER*location	2	1	0.5768	0.2296	6.3081	0.0120
BORDER*location	3	1	0.2085	0.3231	0.4164	0.5187

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
FB	1 vs 0	1.235	0.931	1.638
age1	25 vs 65	3.728	2.556	5.437
age1	35 vs 65	12.609	8.810	18.046
age1	45 vs 65	8.865	6.210	12.656
age1	55 vs 65	3.619	2.518	5.202
year_counted	2000 vs 2010	1.344	0.991	1.825
year_counted	2001 vs 2010	1.427	1.057	1.927
year_counted	2002 vs 2010	1.883	1.397	2.537
year_counted	2003 vs 2010	1.434	1.060	1.939

The LOGISTIC Procedure

Odds Ratio Estimates

Effect	Point Estimate	95% Wald Confidence Limits	
year_counted 2004 vs 2010	1.053	0.777	1.427
year_counted 2005 vs 2010	1.113	0.816	1.519
year_counted 2006 vs 2010	1.306	0.965	1.766
year_counted 2007 vs 2010	1.268	0.934	1.721
year_counted 2008 vs 2010	1.027	0.756	1.395
year_counted 2009 vs 2010	1.109	0.803	1.531
sex	2.240	1.938	2.590
Ethnicity 1 vs 0	0.853	0.690	1.055
race2 Asian vs White	0.219	0.150	0.318
race2 Black vs White	2.039	1.705	2.440
IDU	2.344	1.809	3.038
NoInject	1.569	1.301	1.893
homeless	0.939	0.756	1.165
prison	0.726	0.582	0.906
LTCRes	1.159	0.744	1.804
Occupation 0 vs 1	1.962	1.726	2.229

Association of Predicted Probabilities and Observed Responses

Percent Concordant	79.4	Somers' D	0.593
Percent Discordant	20.2	Gamma	0.595
Percent Tied	0.4	Tau-a	0.146
Pairs	14842724	c	0.796

Partition for the Hosmer and Lemeshow Test

Group	Total	HIV = 1		HIV = 0	
		Observed	Expected	Observed	Expected
1	1099	18	12.57	1081	1086.43
2	1099	29	28.24	1070	1070.76
3	1099	48	44.30	1051	1054.70
4	1100	57	62.43	1043	1037.57
5	1100	71	87.30	1029	1012.70
6	1097	128	118.01	969	978.99
7	1099	147	159.44	952	939.56
8	1100	227	220.55	873	879.45
9	1099	320	316.70	779	782.30
10	1097	532	527.47	565	569.53

The LOGISTIC Procedure

Hosmer and Lemeshow Goodness-of-Fit Test

Chi-Square	DF	Pr > ChiSq
8.9628	8	0.3454

Step 4 *Model with alcohol*USYEARS removed.

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The LOGISTIC Procedure

Model Information

Data Set	TBHIV.LOGISTIC2
Response Variable	HIV
Number of Response Levels	2
Model	binary logit
Optimization Technique	Fisher's scoring

Number of Observations Read	11282
Number of Observations Used	10989

Response Profile

Ordered		Total
Value	HIV	Frequency
1	0	9412
2	1	1577

Probability modeled is HIV=1.

NOTE: 293 observations were deleted due to missing values for the response or explanatory variables.

Class Level Information

Class	Value	Design Variables								
FB	0	0								
	1	1								
year_counted 0	2000	1	0	0	0	0	0	0	0	0
	2001	0	1	0	0	0	0	0	0	0
	2002	0	0	1	0	0	0	0	0	0
	2003	0	0	0	1	0	0	0	0	0
	2004	0	0	0	0	1	0	0	0	0
	2005	0	0	0	0	0	1	0	0	0
	2006	0	0	0	0	0	0	1	0	0
	2007	0	0	0	0	0	0	0	1	0
	2008	0	0	0	0	0	0	0	0	1
	2009	0	0	0	0	0	0	0	0	0
1	2010	0	0	0	0	0	0	0	0	0
age1	25	1	0	0	0					
	35	0	1	0	0					
	45	0	0	1	0					

55	0	0	0	1
65	0	0	0	0

The LOGISTIC Procedure

Class Level Information

Class	Value	Design Variables				
USYEARS	0	1	0	0	0	0
	1	0	1	0	0	0
	2	0	0	1	0	0
	7	0	0	0	1	0
	15	0	0	0	0	1
	20	0	0	0	0	0
race2	Asian	1	0			
	Black	0	1			
	White	0	0			
location	1	0	0			
	2	1	0			
	3	0	1			
Occupation	0	1				
	1	0				
Ethnicity	0	0				
	1	1				

Model Convergence Status

Convergence criterion (GCONV=1E-8) satisfied.

Model Fit Statistics

Criterion	Intercept	Intercept
	Only	and Covariates
AIC	9041.095	7477.700
SC	9048.400	7747.972
-2 Log L	9039.095	7403.700

Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	1635.3954	36	<.0001
Score	1578.6195	36	<.0001
Wald	1221.3175	36	<.0001

The LOGISTIC Procedure

Type 3 Analysis of Effects

Effect	DF	Wald	
		Chi-Square	Pr > ChiSq
BORDER	1	23.6687	<.0001
FB	1	2.5989	0.1069
age1	4	400.6134	<.0001
year_counted	10	37.5892	<.0001
sex	1	118.7003	<.0001
Ethnicity	1	2.9515	0.0858
race2	2	207.6081	<.0001
IDU	1	41.1365	<.0001
USYEARS	4	3.0012	0.5576
NoInject	1	22.5512	<.0001
alcohol	1	32.7800	<.0001
homeless	1	0.3077	0.5791
prison	1	8.1568	0.0043
LTCRes	1	0.4061	0.5240
location	2	200.2052	<.0001
Occupation	1	108.0638	<.0001
BORDER*alcohol	1	2.7236	0.0989
BORDER*location	2	6.2356	0.0443

*Border*alcohol term will be dropped in the next model; it is least significant interaction term (0.10).

NOTE: The following parameters have been set to 0, since the variables are a linear combination

of other variables as shown.

$$\text{USYEARS0} = \text{Intercept} - \text{FB1}$$

Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept		1	-5.0678	0.2398	446.5758	<.0001
BORDER		1	-0.7352	0.1511	23.6687	<.0001
FB	1	1	0.2023	0.1255	2.5989	0.1069
age1	25	1	1.3094	0.1925	46.2675	<.0001
age1	35	1	2.5277	0.1829	191.0784	<.0001
age1	45	1	2.1850	0.1816	144.7231	<.0001
age1	55	1	1.2876	0.1851	48.3812	<.0001
year_counted	2000	1	0.3122	0.1557	4.0220	0.0449
year_counted	2001	1	0.3666	0.1531	5.7330	0.0166
year_counted	2002	1	0.6433	0.1521	17.8803	<.0001
year_counted	2003	1	0.3738	0.1538	5.9102	0.0151
year_counted	2004	1	0.0629	0.1550	0.1649	0.6847
year_counted	2005	1	0.1189	0.1584	0.5629	0.4531
year_counted	2006	1	0.2783	0.1540	3.2669	0.0707

The LOGISTIC Procedure

Analysis of Maximum Likelihood Estimates

Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
year_counted 2007	1	0.2494	0.1557	2.5657	0.1092
year_counted 2008	1	0.0365	0.1562	0.0546	0.8152
year_counted 2009	1	0.1110	0.1646	0.4554	0.4998
sex	1	0.8060	0.0740	118.7003	<.0001
Ethnicity 1	1	-0.1841	0.1072	2.9515	0.0858
race2 Asian	1	-1.5589	0.1899	67.4075	<.0001
race2 Black	1	0.7125	0.0914	60.8081	<.0001
IDU	1	0.8467	0.1320	41.1365	<.0001
USYEARS 0	0	0	.	.	.
USYEARS 1	1	-0.2149	0.1649	1.6969	0.1927
USYEARS 2	1	-0.2345	0.1484	2.4971	0.1141
USYEARS 7	1	-0.1236	0.1624	0.5794	0.4465
USYEARS 15	1	-0.1833	0.1519	1.4556	0.2276
NoInject	1	0.4535	0.0955	22.5512	<.0001
alcohol	1	-0.4801	0.0839	32.7800	<.0001
homeless	1	-0.0611	0.1101	0.3077	0.5791
prison	1	-0.3160	0.1106	8.1568	0.0043
LTCRes	1	0.1438	0.2257	0.4061	0.5240
location 2	1	0.8303	0.0767	117.2121	<.0001
location 3	1	1.1850	0.1065	123.9048	<.0001

Occupation	0	1	0.6776	0.0652	108.0638	<.0001
BORDER*alcohol		1	0.3391	0.2055	2.7236	0.0989
BORDER*location	2	1	0.5657	0.2276	6.1766	0.0129
BORDER*location	3	1	0.2425	0.3205	0.5726	0.4492

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
FB	1 vs 0	1.224	0.957	1.565
age1	25 vs 65	3.704	2.540	5.402
age1	35 vs 65	12.524	8.752	17.923
age1	45 vs 65	8.890	6.228	12.692
age1	55 vs 65	3.624	2.521	5.209
year_counted	2000 vs 2010	1.366	1.007	1.854
year_counted	2001 vs 2010	1.443	1.069	1.948
year_counted	2002 vs 2010	1.903	1.412	2.564
year_counted	2003 vs 2010	1.453	1.075	1.964
year_counted	2004 vs 2010	1.065	0.786	1.443
year_counted	2005 vs 2010	1.126	0.826	1.536
year_counted	2006 vs 2010	1.321	0.977	1.786
year_counted	2007 vs 2010	1.283	0.946	1.741
year_counted	2008 vs 2010	1.037	0.764	1.409
year_counted	2009 vs 2010	1.117	0.809	1.543

The LOGISTIC Procedure

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
sex		2.239	1.937	2.588
Ethnicity	1 vs 0	0.832	0.674	1.026
race2	Asian vs White	0.210	0.145	0.305
race2	Black vs White	2.039	1.705	2.439
IDU		2.332	1.800	3.021
USYEARS	1 vs 20	0.807	0.584	1.115
USYEARS	2 vs 20	0.791	0.591	1.058
USYEARS	7 vs 20	0.884	0.643	1.215
USYEARS	15 vs 20	0.833	0.618	1.121
NoInject		1.574	1.305	1.898
homeless		0.941	0.758	1.167
prison		0.729	0.587	0.906
LTCRes		1.155	0.742	1.797
Occupation	0 vs 1	1.969	1.733	2.238

Association of Predicted Probabilities and Observed Responses

Percent Concordant	79.4	Somers' D	0.591
Percent Discordant	20.3	Gamma	0.593

Percent Tied	0.4	Tau-a	0.145
Pairs	14842724	c	0.796

Partition for the Hosmer and Lemeshow Test

Group	Total	HIV = 1		HIV = 0	
		Observed	Expected	Observed	Expected
1	1101	19	12.84	1082	1088.16
2	1100	29	28.34	1071	1071.66
3	1099	50	44.80	1049	1054.20
4	1099	52	62.38	1047	1036.62
5	1101	78	87.42	1023	1013.58
6	1100	125	118.43	975	981.57
7	1100	143	160.13	957	939.87
8	1099	228	221.12	871	877.88
9	1099	321	317.07	778	781.93
10	1091	532	524.48	559	566.52

The LOGISTIC Procedure

Hosmer and Lemeshow Goodness-of-Fit Test

Chi-Square	DF	Pr > ChiSq
9.6685	8	0.2891

Step 5 *Border*alcohol dropped from this model
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The LOGISTIC Procedure

Model Information

Data Set	TBHIV.LOGISTIC2
Response Variable	HIV
Number of Response Levels	2
Model	binary logit
Optimization Technique	Fisher's scoring

Number of Observations Read	11282
Number of Observations Used	10989

Response Profile

Ordered Value	HIV	Total Frequency
1	0	9412
2	1	1577

Probability modeled is HIV=1.

NOTE: 293 observations were deleted due to missing values for the response or explanatory variables.

Class Level Information

Class	Value	Design Variables								
FB	0	0								
	1	1								
year_counted 0	2000	1	0	0	0	0	0	0	0	0
	2001	0	1	0	0	0	0	0	0	0
	2002	0	0	1	0	0	0	0	0	0
	2003	0	0	0	1	0	0	0	0	0
	2004	0	0	0	0	1	0	0	0	0
	2005	0	0	0	0	0	1	0	0	0
	2006	0	0	0	0	0	0	1	0	0
	2007	0	0	0	0	0	0	0	1	0
	2008	0	0	0	0	0	0	0	0	1
	2009	0	0	0	0	0	0	0	0	0
1	2010	0	0	0	0	0	0	0	0	0
USYEARS	0	1	0	0	0	0				
	1	0	1	0	0	0				
	2	0	0	1	0	0				

7	0	0	0	1	0
15	0	0	0	0	1

The LOGISTIC Procedure

Class Level Information

Class	Value	Design Variables				
	20	0	0	0	0	0
age1	25	1	0	0	0	
	35	0	1	0	0	
	45	0	0	1	0	
	55	0	0	0	1	
	65	0	0	0	0	
race2	Asian	1	0			
	Black	0	1			
	White	0	0			
location	1	0	0			
	2	1	0			
	3	0	1			
Occupation	0	1				
	1	0				
Ethnicity	0	0				
	1	1				

Model Convergence Status

Convergence criterion (GCONV=1E-8) satisfied.

Model Fit Statistics

Criterion	Intercept	Intercept
	Only	and Covariates
AIC	9041.095	7478.390
SC	9048.400	7741.357
-2 Log L	9039.095	7406.390

Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	1632.7052	35	<.0001
Score	1576.7572	35	<.0001
Wald	1220.8425	35	<.0001

The LOGISTIC Procedure

Type 3 Analysis of Effects

Effect	DF	Wald	
		Chi-Square	Pr > ChiSq
BORDER	1	22.3428	<.0001
FB	1	2.7122	0.0996
age1	4	404.4143	<.0001
year_counted	10	37.2144	<.0001
sex	1	119.6223	<.0001
Ethnicity	1	2.8748	0.0900
race2	2	207.3826	<.0001
IDU	1	42.6628	<.0001
USYEARS	4	3.1344	0.5356
NoInject	1	23.3144	<.0001
alcohol	1	30.4156	<.0001
homeless	1	0.4728	0.4917
prison	1	8.3093	0.0039
LTCRes	1	0.4052	0.5244
location	2	201.4582	<.0001
Occupation	1	108.8718	<.0001
BORDER*location	2	6.0024	0.0497

*Border*location can be dropped; it rounds up to 0.05 and is not significant.

NOTE: The following parameters have been set to 0, since the variables are a linear combination

of other variables as shown.

$$\text{USYEARS0} = \text{Intercept} - \text{FB1}$$

Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept		1	-5.0918	0.2395	451.8612	<.0001
BORDER		1	-0.6202	0.1312	22.3428	<.0001
FB	1	1	0.2065	0.1254	2.7122	0.0996
age1	25	1	1.3176	0.1924	46.8774	<.0001
age1	35	1	2.5389	0.1827	193.0145	<.0001
age1	45	1	2.1923	0.1816	145.7323	<.0001
age1	55	1	1.2919	0.1851	48.7121	<.0001
year_counted	2000	1	0.3086	0.1557	3.9282	0.0475
year_counted	2001	1	0.3616	0.1531	5.5798	0.0182
year_counted	2002	1	0.6403	0.1521	17.7144	<.0001
year_counted	2003	1	0.3729	0.1538	5.8767	0.0153
year_counted	2004	1	0.0633	0.1550	0.1668	0.6830
year_counted	2005	1	0.1175	0.1585	0.5504	0.4582
year_counted	2006	1	0.2794	0.1540	3.2902	0.0697
year_counted	2007	1	0.2494	0.1557	2.5642	0.1093

The LOGISTIC Procedure

Analysis of Maximum Likelihood Estimates

Parameter		DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
year_counted	2008	1	0.0349	0.1562	0.0501	0.8229
year_counted	2009	1	0.1128	0.1646	0.4694	0.4932
sex		1	0.8086	0.0739	119.6223	<.0001
Ethnicity	1	1	-0.1817	0.1072	2.8748	0.0900
race2	Asian	1	-1.5521	0.1899	66.8269	<.0001
race2	Black	1	0.7153	0.0914	61.3073	<.0001
IDU		1	0.8610	0.1318	42.6628	<.0001
USYEARS	0	0	0	.	.	.
USYEARS	1	1	-0.2213	0.1647	1.8041	0.1792
USYEARS	2	1	-0.2382	0.1482	2.5820	0.1081
USYEARS	7	1	-0.1268	0.1623	0.6103	0.4347
USYEARS	15	1	-0.1880	0.1518	1.5338	0.2155
NoInject		1	0.4608	0.0954	23.3144	<.0001
alcohol		1	-0.4348	0.0788	30.4156	<.0001
homeless		1	-0.0755	0.1098	0.4728	0.4917
prison		1	-0.3187	0.1106	8.3093	0.0039
LTCRes		1	0.1435	0.2255	0.4052	0.5244
location	2	1	0.8348	0.0767	118.5264	<.0001
location	3	1	1.1864	0.1065	124.1214	<.0001
Occupation	0	1	0.6799	0.0652	108.8718	<.0001

BORDER*location 2	1	0.5522	0.2265	5.9450	0.0148
BORDER*location 3	1	0.2382	0.3197	0.5555	0.4561

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
FB	1 vs 0	1.229	0.962	1.572
age1	25 vs 65	3.735	2.561	5.446
age1	35 vs 65	12.666	8.853	18.121
age1	45 vs 65	8.956	6.274	12.785
age1	55 vs 65	3.640	2.532	5.231
year_counted	2000 vs 2010	1.361	1.003	1.847
year_counted	2001 vs 2010	1.436	1.063	1.938
year_counted	2002 vs 2010	1.897	1.408	2.556
year_counted	2003 vs 2010	1.452	1.074	1.963
year_counted	2004 vs 2010	1.065	0.786	1.443
year_counted	2005 vs 2010	1.125	0.824	1.534
year_counted	2006 vs 2010	1.322	0.978	1.788
year_counted	2007 vs 2010	1.283	0.946	1.741
year_counted	2008 vs 2010	1.036	0.762	1.406
year_counted	2009 vs 2010	1.119	0.811	1.545
sex		2.245	1.942	2.595
Ethnicity	1 vs 0	0.834	0.676	1.029

The LOGISTIC Procedure

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
race2	Asian vs White	0.212	0.146	0.307
race2	Black vs White	2.045	1.710	2.446
IDU		2.366	1.827	3.063
USYEARS	1 vs 20	0.802	0.580	1.107
USYEARS	2 vs 20	0.788	0.589	1.054
USYEARS	7 vs 20	0.881	0.641	1.211
USYEARS	15 vs 20	0.829	0.615	1.116
NoInject		1.585	1.315	1.911
alcohol		0.647	0.555	0.756
homeless		0.927	0.748	1.150
prison		0.727	0.585	0.903
LTCRes		1.154	0.742	1.796
Occupation	0 vs 1	1.974	1.737	2.242

Association of Predicted Probabilities and Observed Responses

Percent Concordant	79.4	Somers' D	0.591
Percent Discordant	20.3	Gamma	0.593
Percent Tied	0.4	Tau-a	0.145

Pairs 14842724 c 0.796

Partition for the Hosmer and Lemeshow Test

Group	Total	HIV = 1		HIV = 0	
		Observed	Expected	Observed	Expected
1	1100	19	12.90	1081	1087.10
2	1099	31	28.53	1068	1070.47
3	1101	50	44.78	1051	1056.22
4	1098	51	62.48	1047	1035.52
5	1096	79	87.15	1017	1008.85
6	1101	124	118.28	977	982.72
7	1099	144	159.80	955	939.20
8	1099	230	220.25	869	878.75
9	1099	313	316.33	786	782.67
10	1097	536	526.50	561	570.50

Hosmer and Lemeshow Goodness-of-Fit Test

Chi-Square	DF	Pr > ChiSq
9.8948	8	0.2725

*Goodness of fit test is not as good as the full model with all interaction terms, but is still insignificant.

Step 6: Gold Standard Model *this is my Gold Standard Model that includes all variables but doesn't include any interaction terms; I will use the OR from this model to assess confounding (this will be the OR that I compare other ORs to as I drop variables from my model;

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Model Information

Data Set	TBHIV.LOGISTIC2
Response Variable	HIV
Number of Response Levels	2
Model	binary logit
Optimization Technique	Fisher's scoring

Number of Observations Read	11282
Number of Observations Used	10989

Response Profile

Ordered		Total
Value	HIV	Frequency

1	0	9412
2	1	1577

Probability modeled is HIV=1.

NOTE: 293 observations were deleted due to missing values for the response or explanatory variables.

Class Level Information

Class	Value	Design Variables								
FB	0	0								
	1	1								
year_counted 0 0 0 0 0 0 0 0	2000	1	0	0	0	0	0	0	0	0
	2001	0	1	0	0	0	0	0	0	0
	2002	0	0	1	0	0	0	0	0	0
	2003	0	0	0	1	0	0	0	0	0
	2004	0	0	0	0	1	0	0	0	0
	2005	0	0	0	0	0	1	0	0	0
	2006	0	0	0	0	0	0	1	0	0
	2007	0	0	0	0	0	0	0	1	0
	2008	0	0	0	0	0	0	0	0	1

1	2009	0	0	0	0	0	0	0	0	0
0	2010	0	0	0	0	0	0	0	0	0
USYEARS	0	1	0	0	0	0				
	1	0	1	0	0	0				
	2	0	0	1	0	0				
	7	0	0	0	1	0				
	15	0	0	0	0	1				

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Class Level Information

Class	Value	Design Variables				
	20	0	0	0	0	0
age1	25	1	0	0	0	
	35	0	1	0	0	
	45	0	0	1	0	
	55	0	0	0	1	
	65	0	0	0	0	
race2	Asian	1	0			
	Black	0	1			
	White	0	0			

location	1	0	0
	2	1	0
	3	0	1
Occupation	0	1	
	1	0	
Ethnicity	0	0	
	1	1	

Model Convergence Status

Convergence criterion (GCONV=1E-8) satisfied.

Model Fit Statistics

Criterion	Intercept Only	Intercept and Covariates
AIC	9041.095	7480.241
SC	9048.400	7728.600
-2 Log L	9039.095	7412.241

Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	1626.8537	33	<.0001
Score	1574.3059	33	<.0001
Wald	1223.3705	33	<.0001

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The LOGISTIC Procedure

Type 3 Analysis of Effects

Effect	DF	Wald	
		Chi-Square	Pr > ChiSq
BORDER	1	18.7470	<.0001
FB	1	2.8260	0.0928
age1	4	403.9406	<.0001
year_counted	10	37.7005	<.0001
sex	1	118.9423	<.0001
Ethnicity	1	2.9498	0.0859
race2	2	207.3788	<.0001
IDU	1	43.0944	<.0001
USYEARS	4	3.5278	0.4737
NoInject	1	23.4284	<.0001
alcohol	1	30.1381	<.0001
homeless	1	0.4004	0.5269
prison	1	9.4440	0.0021

LTCRes	1	0.3591	0.5490
location	2	245.4490	<.0001
Occupation	1	109.1809	<.0001

NOTE: The following parameters have been set to 0, since the variables are a linear combination of other variables as shown.

$$USYEARS0 = \text{Intercept} - \text{FB1}$$

Analysis of Maximum Likelihood Estimates

Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept	1	-5.1065	0.2396	454.2827	<.0001
BORDER	1	-0.4627	0.1069	18.7470	<.0001
FB	1	0.2102	0.1251	2.8260	0.0928
age1	25	1.3185	0.1925	46.9140	<.0001
age1	35	2.5402	0.1827	193.2306	<.0001
age1	45	2.1909	0.1816	145.6124	<.0001
age1	55	1.2948	0.1850	48.9624	<.0001
year_counted 2000	1	0.3120	0.1557	4.0146	0.0451
year_counted 2001	1	0.3646	0.1531	5.6683	0.0173
year_counted 2002	1	0.6420	0.1521	17.8037	<.0001
year_counted 2003	1	0.3808	0.1538	6.1273	0.0133
year_counted 2004	1	0.0648	0.1550	0.1747	0.6760
year_counted 2005	1	0.1177	0.1585	0.5517	0.4576

year_counted 2006	1	0.2812	0.1541	3.3320	0.0679
year_counted 2007	1	0.2514	0.1558	2.6043	0.1066
year_counted 2008	1	0.0330	0.1562	0.0447	0.8325

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The LOGISTIC Procedure

Analysis of Maximum Likelihood Estimates

Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
year_counted 2009	1	0.1165	0.1646	0.5011	0.4790
sex	1	0.8071	0.0740	118.9423	<.0001
Ethnicity 1	1	-0.1842	0.1072	2.9498	0.0859
race2 Asian	1	-1.5594	0.1900	67.3824	<.0001
race2 Black	1	0.7130	0.0914	60.8137	<.0001
IDU	1	0.8631	0.1315	43.0944	<.0001
USYEARS 0	0	0	.	.	.
USYEARS 1	1	-0.2381	0.1642	2.1042	0.1469
USYEARS 2	1	-0.2525	0.1478	2.9176	0.0876
USYEARS 7	1	-0.1365	0.1620	0.7098	0.3995
USYEARS 15	1	-0.1929	0.1515	1.6211	0.2029
NoInject	1	0.4613	0.0953	23.4284	<.0001
alcohol	1	-0.4327	0.0788	30.1381	<.0001
homeless	1	-0.0695	0.1098	0.4004	0.5269

prison		1	-0.3384	0.1101	9.4440	0.0021
LTCRes		1	0.1351	0.2255	0.3591	0.5490
location	2	1	0.8940	0.0725	151.9128	<.0001
location	3	1	1.2120	0.1008	144.5507	<.0001
Occupation	0	1	0.6806	0.0651	109.1809	<.0001

Odds Ratio Estimates

Effect		Point Estimate	95% Wald Confidence Limits	
BORDER		0.630	0.511	0.776
FB	1 vs 0	1.234	0.966	1.577
age1	25 vs 65	3.738	2.563	5.451
age1	35 vs 65	12.682	8.865	18.145
age1	45 vs 65	8.943	6.265	12.765
age1	55 vs 65	3.650	2.540	5.246
year_counted	2000 vs 2010	1.366	1.007	1.854
year_counted	2001 vs 2010	1.440	1.067	1.944
year_counted	2002 vs 2010	1.900	1.410	2.560
year_counted	2003 vs 2010	1.463	1.083	1.978
year_counted	2004 vs 2010	1.067	0.787	1.446
year_counted	2005 vs 2010	1.125	0.825	1.535
year_counted	2006 vs 2010	1.325	0.979	1.792
year_counted	2007 vs 2010	1.286	0.948	1.745
year_counted	2008 vs 2010	1.034	0.761	1.404
year_counted	2009 vs 2010	1.124	0.814	1.551
sex		2.241	1.939	2.591
Ethnicity	1 vs 0	0.832	0.674	1.026
race2	Asian vs White	0.210	0.145	0.305
race2	Black vs White	2.040	1.705	2.441

The LOGISTIC Procedure

Odds Ratio Estimates

Point	95% Wald		
Effect	Estimate	Confidence Limits	
IDU	2.371	1.832	3.067
USYEARS 1 vs 20	0.788	0.571	1.087
USYEARS 2 vs 20	0.777	0.581	1.038
USYEARS 7 vs 20	0.872	0.635	1.198
USYEARS 15 vs 20	0.825	0.613	1.110
NoInject	1.586	1.316	1.912
alcohol	0.649	0.556	0.757
homeless	0.933	0.752	1.157
prison	0.713	0.575	0.885
LTCRes	1.145	0.736	1.781
location 2 vs 1	2.445	2.121	2.818
location 3 vs 1	3.360	2.758	4.094
Occupation 0 vs 1	1.975	1.738	2.244

Association of Predicted Probabilities and Observed Responses

Percent Concordant	79.3	Somers' D	0.590
Percent Discordant	20.3	Gamma	0.593
Percent Tied	0.4	Tau-a	0.145
Pairs	14842724	c	0.795

Partition for the Hosmer and Lemeshow Test

	HIV = 1	HIV = 0			
Group	Total	Observed	Expected	Observed	Expected
1	1102	20	13.40	1082	1088.60
2	1100	32	29.34	1068	1070.66
3	1099	46	45.12	1053	1053.88
4	1099	50	63.21	1049	1035.79
5	1099	84	87.94	1015	1011.06
6	1099	119	118.04	980	980.96
7	1099	150	158.59	949	940.41
8	1099	224	219.23	875	879.77
9	1101	321	316.21	780	784.79
10	1092	531	525.93	561	566.07

Hosmer and Lemeshow Goodness-of-Fit Test

Chi-Square DF Pr > ChiSq

7.5575 8 0.4778

*Goodness-of-Fit test - the p-value is less significant, indicating a better fit than the models containing interaction terms.

***Gold standard OR = 0.630 (CI 0.511, 0.776)