

Coccidioidomycosis: epidemiology

Jennifer Brown¹
Kaitlin Benedict²
Benjamin J Park²
George R Thompson III^{1,3}

¹Department of Internal Medicine, Division of Infectious Diseases, University of California, Davis Medical Center, Sacramento, CA, USA; ²Mycotic Diseases Branch, Division of Foodborne, Waterborne, and Environmental Diseases, US Centers for Disease Control and Prevention, Atlanta, GA, USA;

³Department of Medical Microbiology and Immunology, One Shields Avenue, Tupper Hall, Coccidioidomycosis Serology Laboratory, University of California, Davis, CA, USA

Abstract: Coccidioidomycosis consists of a spectrum of disease, ranging from a mild, self-limited, febrile illness to severe, life-threatening infection. It is caused by the soil-dwelling fungi, *Coccidioides immitis* and *C. posadasii*, which are present in diverse endemic areas. Climate changes and environmental factors affect the *Coccidioides* lifecycle and influence infection rates. The incidence of coccidioidomycosis has risen substantially over the past two decades. The vast majority of *Coccidioides* infections occur in the endemic zones, such as California, Arizona, Mexico, and Central America. Infections occurring outside those zones appear to be increasingly common, and pose unique clinical and public health challenges. It has long been known that elderly persons, pregnant women, and members of certain ethnic groups are at risk for severe or disseminated coccidioidomycosis. In recent years, it has become evident that persons with immunodeficiency diseases, diabetics, transplant recipients, and prisoners are also particularly vulnerable.

Keywords: coccidioidomycosis, *Coccidioides*, epidemiology, incidence, risk factors, geography

Introduction

Coccidioides spp. are dimorphic, soil-dwelling, fungi known to cause a broad spectrum of disease, ranging from a mild febrile illness to severe pulmonary manifestations or disseminated disease.^{1,2} The genus *Coccidioides* is comprised of two genetically distinct species: *Coccidioides immitis* and *C. posadasii*. These two species cause similar clinical diseases, however they are present in different geographic regions of endemicity.³⁻⁵ This review addresses the history of *Coccidioides spp.* and summarizes the current knowledge of their ecologic environment and geographic distribution. The risk factors for disease acquisition and recent epidemiologic trends, including coccidioidomycosis in non-endemic areas, the enlarging at-risk populations, and current public health challenges are highlighted.

History

The causative agent of coccidioidomycosis was first identified by a medical intern, Alejandro Posadas, in Buenos Aires in 1892.⁶ During his medical training, Posadas evaluated an Argentine soldier who had skin lesions previously attributed to mycosis fungoides. Microscopic examination of the soldier's skin biopsy specimens revealed organisms that appeared similar to the protozoan *Coccidia*, leading to a misconception that the microbe was a parasite. A few years later, a manual laborer from the San Joaquin Valley, California, was evaluated for similar skin lesions.⁷ Upon his death,

Correspondence: Benjamin J Park
US Centers for Disease Control
and Prevention, 1600 Clifton Rd NE,
Mailstop C09, Atlanta, GA 30333, USA
Email bip5@cdc.gov

George R Thompson III
Department of Medical Microbiology
and Immunology, Department
of Medicine, Division of Infectious
Diseases, University of California,
Davis, One Shields Ave, Tupper Hall,
Rm 3138, Davis, CA 95616, USA
Tel +1 530 752 3545
Email grthompson@ucdavis.edu

autopsy specimens from multiple organs were notable for granulomas containing the same type of “protozoal” organisms. A collaborative evaluation of this patient’s specimens by Emmet Rixford, a surgeon at Cooper Medical College in San Francisco, and T Caspar Gilchrist and CW Stiles, pathologists at Johns Hopkins Medical School, resulted in the decision that the organism was, indeed, a protozoan. Consequently, in 1896, it was given the name *Coccidioides* (derived from its morphologic appearance of “resembling *Coccidia*”) *immitis* (“not mild”, because it was believed that the organism caused lethal disease).^{6,7} Just a few years later, William Ophüls and Herbert C Moffitt, accurately classified *C. immitis* as a fungus.⁸ These investigators satisfied Koch’s postulates by inoculating material from a case patient into guinea pigs, which subsequently developed signs of infection, culturing the fungal organism from the animals’ organs, and ultimately injecting mycelia from this culture into a rabbit, which developed *Coccidioides*-containing nodules in tissue.

Important advances in the understanding of *Coccidioides* pathogenesis and natural history were made during the first half of the twentieth century. In 1929, the conventional notion that coccidioidal infections were rare and consistently fatal was questioned after Harold Chope, a medical student, accidentally inhaled *Coccidioides* spores from a culture plate in the Stanford University laboratory of Ernest Dickson.⁶ Chope developed pneumonia and, despite the grim expectation that death was imminent, he survived. This surprising outcome, along with the fact that the fungus was isolated from Chope’s respiratory specimen, sparked a surge of investigations by Dickson and others. In 1938, Dickson linked “San Joaquin Fever” or “Valley Fever”, the self-resolving illness of cough, chest pain, fever, and erythema nodosum, to dust exposure in patients who had positive reactions to *Coccidioides* skin testing.^{6,9} A few years later, Charles E Smith collected information from over 400 patients with a history of Valley Fever and found that the majority of infections were mild and that there was no evidence for human-to-human transmission.¹⁰ A decade later, Smith and others published the details of a cluster of coccidioidal infections in a group of students in Kern County, California.¹¹ It was determined that the students had been exposed to *Coccidioides* while digging a rattlesnake out of a ground squirrel hole. These reports led to the present-day understanding that coccidioidomycosis is acquired via inhalation of contaminated dust or soil and that it generally does not cause lethal disease.

Ecology

Coccidioides spp. are found in warm, arid, desert regions in the Western Hemisphere. They thrive in an ecological zone that has hot summers, gentle winters without harsh freezes, and annual rainfall of 10–50 cm.^{5,12} The organisms are found within alkaline, sandy soil, commonly about 10–30 cm beneath the surface, though they can have a sporadic or patchy distribution within a given locality.^{5,13}

The *Coccidioides spp.* lifecycle is closely intertwined with changes in climate conditions. The fungal mycelia require moisture in the soil to grow.^{14,15} Hence the fungus is usually recovered in greatest abundance in the spring after the heavy, winter rainfall has ended. The hyphae then need a dry period to promote desiccation and maturation into arthroconidia, which can be aerosolized and inhaled (Figure 1).^{5,14–16} This requirement partially explains why the rate of *Coccidioides* infections tends to rise in the drier months of the year. Other subtle factors in the ecological milieu, such as variations in the chemical components of the soil, may also affect fungal subsistence. In an early study, Elconin and colleagues evaluated the soil in a parcel of land in the San Joaquin Valley where coccidioidomycosis was common; more than 5000 soil samples were collected at monthly intervals over an 8-year period (1955–1962).¹⁷ After controlling for other variables, including yearly temperature and rainfall levels, the authors found that a higher concentration of soluble salts (eg, sodium, calcium, magnesium, sulfates, and chlorides) in the soil was significantly correlated with the presence of *C. immitis*. Thus, differences in soil constitution may account for the patchy distribution of these two *Coccidioides* species within a particular region, as well as for their divergence in geographical distribution.

Ecology – role of an animal vector?

Interestingly, some reports describe higher concentrations of the organisms around archaic Indian burial sites or animal burrows.^{18,19} The later observation has led to speculation that there may be a rodent host reservoir for *Coccidioides*; both the kangaroo rat and the Arizona pocket mouse have been proposed as possible animal reservoirs. However, to date, zoonotic transmission to humans has not been reported. Although carcasses or excretions from infected rodents have been hypothesized to play a role in the environmental life cycle of the fungus,^{14,18} the extent to which rodents or other animals such as bats²⁰ or armadillos²¹ act as a reservoir for *Coccidioides* or influence its geographic distribution is still unknown. Comparative whole genome sequencing

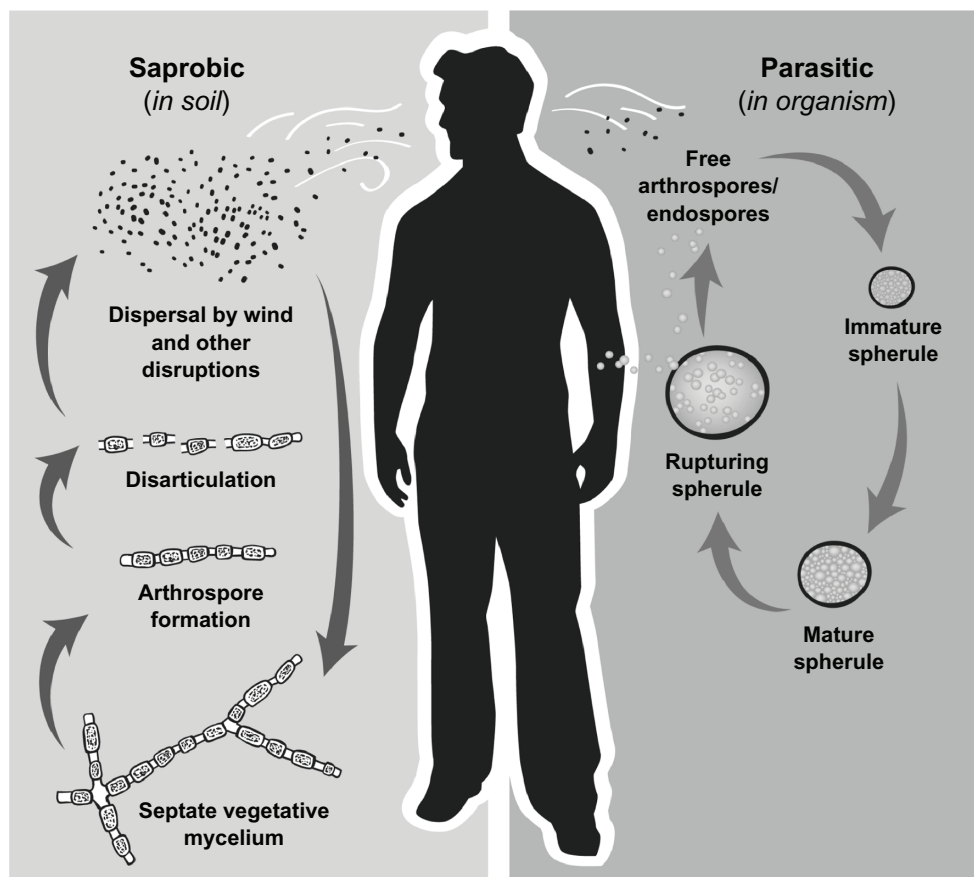


Figure 1 Life cycle of *Coccidioides* spp.

Notes: In the environment *Coccidioides* exists as a mycelium and following periods of low precipitation arthroconidia are formed and are easily aerosolized when disturbed. Arthroconidia can be inhaled or return to the soil and again grow to vegetative mycelia. However if inhaled, arthroconidia undergo a morphologic change and become immature spherules which divide internally until filled with endospores and subsequently rupture. Endospores are dispersed into the surrounding tissue and are then able to form new spherules and repeat the cycle.

data suggest that *Coccidioides* have evolved in response to interaction with an animal host.²²

Coccidioidomycosis has been shown to affect other non-human mammals, including domestic and non-domestic animals in the wild and in captivity.²³ It is especially common among domestic dogs, with an estimated annual incidence of 4% among dogs in Pima and Maricopa Counties, Arizona.²⁴ Because dogs presumably share similar exposures to their human counterparts, studies of canine coccidioidomycosis may be useful for assessing the risk for human infection, particularly in suspected or known, but broadly-defined, endemic areas.²⁵

Geographic range

Historically, methods to isolate *Coccidioides* from the soil across wide-ranging regions have been neither feasible nor practical. As a result, the geographic range for *Coccidioides* spp. has been largely extrapolated from epidemiologic

studies of persons diagnosed with coccidioidomycosis or from population surveys via spherulin or coccidioidin skin testing. One early investigation, that used skin testing to map the distribution of *Coccidioides* in the USA, was conducted by Edwards and Palmer in 1957.²⁶ The results of this and similar studies established that the south-central valley of California and the deserts of southern Arizona, were the most highly endemic for *Coccidioides*.^{3,18,27} In the future, culture-independent methods, such as multiplex polymerase chain reaction testing of bulk soil samples, may prove to be useful tools to identify *Coccidioides* locations.²⁸

Presently, it is known that *Coccidioides* spp. are endemic to specific regions in the Western Hemisphere, primarily those located between the north and south 40° latitudes (Figure 2).^{5,29} The two species, *C. immitis* and *C. posadasii*, populate two distinct and divergent geographic regions. *C. immitis* is found in central and southern California, with the San Joaquin Valley being the region of greatest endemicity.^{3,5,13}

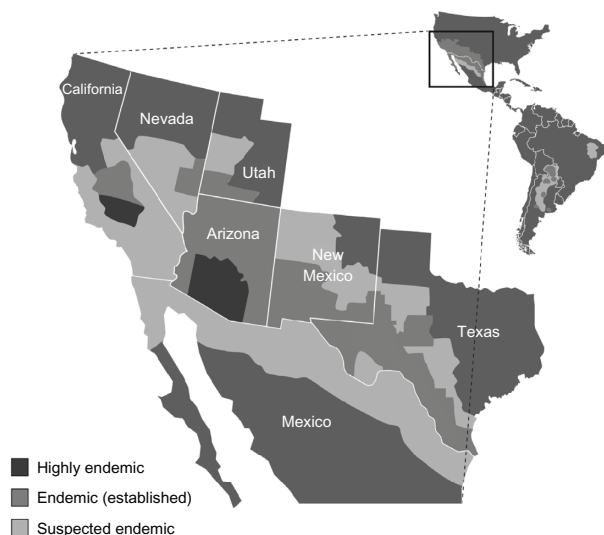


Figure 2 Endemic areas for coccidioidomycosis.

The rate of positive skin tests ranges as high as 50%–70% in Kern County, including the city of Bakersfield, and its neighboring Tulare and Kings Counties.²⁶ Compared with *C. immitis*, *C. posadasii* has a broader region of endemicity, from central and southern Arizona to western Texas and southern New Mexico. The most concentrated region for *C. posadasii* is in Arizona, where the majority of positive skin tests and coccidioidomycosis cases occur in Maricopa County (including the city of Phoenix), Pima County (including the city of Tucson), and Pinal County.³ *C. posadasii* is also present in sporadic sites in southern Utah and Nevada.^{13,30}

Beyond the USA, *C. posadasii* is present in parts of Mexico and Central and South America.^{4,5,31–33} On the basis of surveys using coccidioidin testing conducted in the 1960s in Mexico, it was concluded that *Coccidioides*-endemic regions include the Northern zone bordering the USA, (eg, the Mojave, Sonoran, and Chihuahuan Desert regions), the Pacific Coast zone (eg, Nayarit, Jalisco, and Michoacan), and the Central Zone (eg, Durango).^{5,33} More limited skin surveys, performed in the 1990s, found rates of positivity ranging from 40% to over 90% in the state of Coahuila.³⁴ The mapping of *Coccidioides* endemicity in Central and South America is less complete. This reflects the fact that, in most countries, coccidioidomycosis is not considered a reportable disease, and nationwide skin testing surveys have not been conducted. In Central America, skin testing has identified regions of endemicity in the Montagua River Valley in Guatemala and in Comayagua, Honduras.^{4,5,31,33} In South America, endemic regions have been found in northern Venezuela and northeastern Brazil.^{5,33} Argentina, Bolivia, Paraguay, and Nicaragua also may have some endemic

areas.^{4,5,33} The precision of these surveys is unknown, though, since skin tests may cross-react with other mycoses such as *Histoplasma capsulatum*, *Blastomyces dermatitidis*, and *Paracoccidioides brasiliensis*, which may also be endemic in these regions.

Coccidioidomycosis in non-endemic areas

While the vast majority of the burden of coccidioidomycosis exists in Arizona and California, there is evidence to suggest that the disease also poses specific clinical and public health challenges outside the endemic areas. The majority of cases observed outside the southwestern USA occur among persons who visit or temporarily relocate to endemic areas and seek medical care after returning to their permanent residence. In many instances, travelers become symptomatic shortly after returning home, although latent or reactivation infection presenting months to years later can also occur in immunosuppressed persons.^{35,36} As the infectious dose of *Coccidioides* arthroconidia is presumed to be very small, even a very brief exposure in an endemic area may result in symptomatic infection.

Clusters of coccidioidomycosis cases among groups of travelers have been observed, including clusters in Pennsylvania³⁷ and Washington State³⁸ among church group members who had visited Mexico. Another cluster occurred in US Marine reserves from Tennessee who completed 3 weeks of training exercises in southern California.³⁹ In each of these reports, nearly all case-patients had extensive exposure to soil or dusty environments. In addition, the travelers were unaware of their risk for acquiring coccidioidomycosis, and many case-patients were initially misdiagnosed with bacterial or viral respiratory infections.^{38,39} Several series of isolated coccidioidomycosis cases (not occurring as part of a cluster) outside endemic areas have been described, including 23 cases at an Ohio hospital during 1980–1998,⁴⁰ 161 cases identified from New York State hospital discharge records during 1992–1997,⁴¹ two cases with musculoskeletal involvement in Chicago,⁴² and four cases in New Orleans.⁴³ Travel-associated coccidioidomycosis cases have also occurred among visitors from other countries,⁴⁴ including an outbreak after a model airplane flying event in Kern County, California in 2001, which over 300 persons from more than 30 countries attended.^{45,46} These reports of coccidioidomycosis in non-endemic areas highlight the importance of educating travelers and healthcare providers in non-endemic areas about the disease so that delays in diagnosis and treatment can be avoided or minimized.

Recent evidence suggests that some coccidioidomycosis cases occurring outside the endemic area may be locally-acquired, including three cases in eastern Washington (one case of primary cutaneous coccidioidomycosis and two cases of pneumonia, one of which progressed to meningeal disease).⁴⁷ Another case occurred in a 14-year old Chinese boy who had never visited, or been exposed to materials imported from, an endemic area. Interestingly, he had reported choking on seawater,⁴⁸ an environment in which *Coccidioides* can survive.⁴⁹ These occurrences suggest that the geographic range of *Coccidioides* may be expanding or is wider than that previously recognized.

Risk factors for infection

Any person who resides in or travels to a *Coccidioides* endemic region can become infected with *Coccidioides spp.* following inhalation of airborne arthroconidia. Although infections contracted from non-respiratory routes (eg, in utero exposure, animal bites) have been reported, these are rare.^{50,51}

Populations with intense exposure to aerosolized arthroconidia are at greater risk for infection. These groups include agricultural or construction workers, or persons who participate in outdoor activities such as hunting or digging in the soil.^{5,11,52,53} Outbreaks of coccidioidomycosis have been linked to military training exercises, model airplane competitions, earthquakes, windstorms, and armadillo hunting adventures.^{38,54–58} Though uncommon, infections acquired in persons with minimal exposure (eg, changing airplanes at the Phoenix airport or briefly driving through the San Joaquin Valley) have also been reported.^{41,59}

After inhaling *Coccidioides* spores, 60% of infected persons remain asymptomatic. The rest develop mild-to-severe symptomatic pulmonary infections. About 1% of infected persons develop disseminated disease, which can involve the skin, joints, bones, central nervous system, or other organs. Many factors are correlated with an increased risk for severe or disseminated disease (Table 1).

Age

Although coccidioidomycosis affects all age groups, the highest incidence rates are consistently documented among adults, typically among those aged 40–49 years in California⁶⁶ and among those aged ≥ 65 in Arizona.^{60,61} Several studies have examined the factors associated with the disproportionate burden of coccidioidomycosis among the elderly in Arizona. For example, a case-control study showed that among coccidioidomycosis case-patients

Table 1 Risk factors for severe or disseminated coccidioidomycosis

Filipino or African ethnicity
HIV/AIDS
Immunosuppressive medications
Prednisone
TNF- α inhibitors
Chemotherapy
Organ transplantation (tacrolimus, etc)
Diabetes mellitus
Pregnancy
Cardiopulmonary disease

Abbreviations: AIDS, acquired immunodeficiency syndrome; HIV, human immunodeficiency virus; TNF, tumor necrosis factor.

aged ≥ 60 years, the risk for acquiring symptomatic coccidioidomycosis declined by 5% for each year lived in the endemic area.⁶⁰ After controlling for duration of residence, congestive heart failure, corticosteroid use, smoking, cancer, and male sex were independently associated with developing symptomatic coccidioidomycosis among cases compared to geographically-matched controls.⁶⁰ This suggests that coccidioidomycosis among the elderly in Arizona may be related, at least in part, to the high prevalence of seasonal or permanent relocation of retirement-age individuals (many of whom have comorbid conditions) from non-endemic areas.

While an older population at risk may explain the high rates among the elderly, other factors may more fully account for the risk of developing adverse outcomes. A retrospective medical record review of Arizona coccidioidomycosis cases found no significant differences in disease manifestations between patients aged ≥ 60 years and those aged < 60 years when controlling for comorbid conditions; regardless of age or ethnicity, immunosuppression was the only factor associated with poor outcomes including hospitalization, disseminated infection, or death.⁶² Similarly, a study of Kern County, California coccidioidomycosis cases found that older age was independently associated with developing severe pulmonary disease, but not with developing disseminated disease.⁶³ Altogether, these studies suggest that additional research into the role of age as a risk factor for infection and for the development of adverse outcomes is warranted.

Gender

Historically, coccidioidomycosis has been more common among men than women. Surveillance data from Arizona from 1993–2006 have consistently shown a preponderance of cases in men (55%–66%) compared with women, although more cases have occurred among females in Arizona

since 2009.^{60,61,64} In California, from 2000–2007, men had an average annual rate of 7.6 *Coccidioides* infections per 100,000 compared with a rate of 4.0 per 100,000 in women.⁶⁵ This gender distribution may be partially explained by the greater participation by men in high-risk activities or cigarette smoking. Certain human sex hormones, which have been shown to stimulate the growth of *Coccidioides* in vitro, may also contribute to the elevated incidence in males.⁶⁶

Severe or disseminated disease may occur more often in men than in women. In the 1940s, Smith and colleagues performed serial coccidioidin skin testing and clinical evaluations of military personnel at US Army airfields in the San Joaquin Valley.⁶⁷ Though the number of women in the study was small, the frequency of disseminated disease in women was one-fifth the rate found in men. Durry et al evaluated 128 cases of coccidioidomycosis that occurred in 1991 in Tulare County, California, and found that male sex was associated with a higher rate of hospitalization for severe disease.⁶⁸ In contrast, among the aforementioned coccidioidomycosis cases (n = 380) in Kern County, there was no correlation between male gender and severe or disseminated disease.⁶³

Race

Extrapulmonary dissemination of *Coccidioides* develops more frequently in certain ethnic groups, including those of African or Filipino descent.⁶⁹ A study by Smith et al, in the 1940s, found that black persons had rates of dissemination 8–10 times higher than those in white persons.⁶⁷ Though the age of the subjects in these studies was not reported, all subjects worked in similar conditions and received the same nutrition, housing, and medical care. Decades later, after a severe dust storm led to an outbreak of coccidioidomycosis in California, a disproportionate increase in the incidence of disseminated infection was noted in the non-Caucasian population; the rate of dissemination among African American men was 23.8 cases per 100,000 versus 2.5 cases per 100,000 for Caucasian men.⁵³ Similar findings were reported from Kern County where, in 1995–1996, large numbers of coccidioidomycosis cases occurred in Filipinos, Asians, Hispanics, and blacks.⁶³ Although black race was not significantly associated with severe pulmonary disease, it was associated with disseminated coccidioidomycosis. The other ethnic groups were not linked with either elevated rates of severe pulmonary disease or extrapulmonary dissemination, though the low numbers of Filipinos or Asians in the study limited the analysis. Others have confirmed these findings using geographic and demographic strata to control for differences in group-level exposure – making the results more externally valid than prior

epidemiologic studies conducted solely on military personnel, clinic or hospitalized patients, or student population.^{70,71}

The causes for racial disparities for *Coccidioides* infections are not completely understood. Genetic influences, such as the ABO blood group type and human leukocyte antigen (HLA) alleles (eg, the HLA class II DRB1*1301 allele has been associated with the development of severe, disseminated disease), may play a role,^{69,72} although these may simply be surrogate markers for at-risk ethnic populations.⁷³

Pregnancy

Pregnant women are especially vulnerable to coccidioidal infection, and their risk of developing severe or disseminated disease rises when infection is acquired in the later stages of pregnancy.^{74,75} Also, severe coccidioidomycosis is more likely to occur during the immediate postpartum period if infection is acquired in the third trimester.^{74,76} Two retrospective reviews of Kern County birth records from the 1940s to the 1960s were notable for a rate of 7.7–11 coccidioidomycosis cases per 10,000 pregnancies, with a striking rate of death (33%) among women who had disseminated disease.^{77,78} A more recent review of 81 coccidioidomycosis cases in pregnant women found that disseminated disease occurred in 50% of the cases diagnosed in the first trimester, 62% in the second trimester, and 96% in the third trimester.⁷⁵ These extremely high rates of dissemination, which have not been seen in other investigations, may reflect a component of reporting bias. In contrast, Wack et al, in a 1988 review of over 47,000 deliveries from three health care centers in Tucson, Arizona, found only 10 cases of coccidioidomycosis (rate of 2.1 cases per 10,000 pregnancies).⁷⁶ The authors did, however, find an association of severe disease with increasing stage of pregnancy; of the seven cases diagnosed in the first or second trimester, illness resolved in all, whereas two-thirds of the cases diagnosed in the third trimester developed disseminated infection. More recently, among 32 cases of coccidioidomycosis in pregnant women in Kern County, dissemination occurred in about 10% of cases.⁷⁹ Sex hormone influences, combined with diminished cell-mediated immunity, may explain this predilection for disseminated coccidioidomycosis to rise as pregnancy progresses.^{66,80}

Immunosuppression

Persons that are especially vulnerable to developing severe or disseminated coccidioidomycosis are those with diseases that impair T-cell function. Such conditions include hematologic malignancies, inflammatory arthritis, and diabetes.^{2,63,81–83} In one review of 55 patients with

hematologic malignancies and coccidioidomycosis close to half of all cases occurred in patients with non-Hodgkin lymphoma (NHL) or chronic lymphocytic leukemia (CLL).⁸¹ As compared to other hematologic malignancies, NHL and CLL did not appear to be disproportionately represented since these two hematologic malignancies comprised the majority of the malignancies in their cancer registry. Nonetheless, a striking percentage (20%) of patients did experience extrapulmonary dissemination and 50% of those with disseminated infection died; patients with NHL had the highest rate of dissemination (35%). In patients with diabetes, poor glycemic control (blood glucose levels ≥ 220 mg/dL) may increase the risk of dissemination.⁸³ In addition, medical therapies that impair cellular immunity, such as antineoplastic agents, high dose corticosteroid therapy (equivalent to a prednisone dose of >20 mg/day), or tumor necrosis factor- α antagonists heighten the risk for severe or disseminated coccidioidomycosis.^{84–86} Other immunodeficiency states that confer a significant risk for *Coccidioides* infection are HIV infection and immunosuppression following hematopoietic stem cell or solid organ transplantation.⁸⁷

Transplant

Coccidioidomycosis has been described as a serious infectious complication among organ transplant recipients in endemic areas; infection is believed to most commonly result from reactivation of previously acquired coccidioidomycosis.⁸⁸ Risk factors for post-transplant coccidioidomycosis include a history of coccidioidomycosis, positive serology at the time of transplantation, and the use of antirejection therapy; however, concurrent illnesses such as cytomegalovirus, diabetes, and hepatitis C virus infection have not been shown to be associated with coccidioidomycosis among transplant recipients.^{88–91}

Early retrospective studies of coccidioidomycosis among solid organ transplant recipients in Arizona calculated incidence rates as high as 6.9% among kidney transplant recipients during 1970–1979⁹² and 8.7% among heart transplant recipients during 1979–1984,⁹³ with high rates of dissemination (75%) and overall mortality (63%).⁹² A more recent report from one large Arizona institution found lower rates during 1999–2006 of 1.2% to 2.4% among hematologic and liver transplant recipients respectively, possibly related to their use of targeted antifungal prophylaxis among patients at high risk for post-transplant coccidioidomycosis.⁹⁴ Other reports of small case series indicate that considerable rates ($>30\%$) of disseminated coccidioidomycosis and mortality still occur among transplant recipients.^{89,90}

Several instances of coccidioidomycosis acquired from donated organs have also been documented in persons with no history of exposure to endemic areas, including lung,^{95–98} liver,^{99,100} kidney,^{96,99–101} pancreas,⁹⁶ and heart,⁹⁹ transplant recipients; the majority of these reports describe disseminated and/or fatal cases. The exact risk for coccidioidomycosis transmission through an infected organ is unknown. A prospective study of 568 healthy potential liver or kidney donors in Arizona found that 12 (2.1%) had a positive coccidioidal serology, four of whom proceeded to donate an organ, but none of the recipients developed coccidioidomycosis.¹⁰² Nevertheless, donor-derived coccidioidomycosis is likely to be under-recognized in endemic areas, due to the reasonable assumption that infection in a transplant patient is the result of re-activation of prior infection or acquired from the local environment.^{94,102} Further research is needed to better understand if screening donors or recipients with a serological test can reduce the risk of donor-derived *Coccidioides* transmission.¹⁰²

HIV/AIDS

Coccidioidomycosis is also a well-recognized opportunistic infection among HIV-infected individuals, though its incidence and severity in this population appears to have declined in the USA with the broader use of combination antiretroviral therapy (cART) beginning around 1996.^{103,104} A prospective study at an Arizona HIV clinic in 1988 showed a cumulative incidence of active coccidioidomycosis of 25% during 41 months of follow-up, corresponding to an average annual incidence of 7.3%.¹⁰⁵ In contrast, a retrospective review at the same clinic during 2003–2008 found an annual incidence of only 0.9% and a decrease in severity of disease compared to the previous study.⁸⁷ As with other HIV-associated opportunistic infections, CD4 cell counts are the most important risk factors for coccidioidomycosis. Studies conducted both before and after the advent of cART have shown CD4 count to be the only predictor for developing active coccidioidomycosis; factors such as a history of coccidioidomycosis and duration of residence in an endemic area¹⁰⁵ or age, sex, race, ethnicity, plasma HIV RNA level, or receipt of cART were not associated with increased risk for coccidioidomycosis.⁸⁷ Another study matched HIV-infected patients based on CD4 count, age, sex, and county of residence, and found that black race, a history of oropharyngeal or esophageal candidiasis, and not receiving any medication were independent risk factors for developing coccidioidomycosis, while use of protease inhibitors was protective; in patients with a history

of oropharyngeal or esophageal candidiasis, receipt of an azole drug in the 90 days preceding the diagnosis of coccidioidomycosis was associated with a reduced risk for developing coccidioidomycosis.¹⁰⁴

Prisons

Coccidioidomycosis has been described in a number of prisons in endemic areas, with a large degree of associated morbidity. In California, a large burden of disease occurs in state and federal correctional institutions in the San Joaquin Valley,¹⁰⁶ and recently, incidence has been increasing in some prisons, with rates as high as 7% during 2006–2010.¹⁰⁷ The reasons for high rates among prisoners are not entirely clear, but may be due, at least in part, to large numbers of immune-naïve persons who have been relocated from other, less-endemic areas. Public health officials have instituted aggressive measures to reduce the morbidity and mortality of this disease, including widespread education of prisoners and staff and exclusion of immunosuppressed individuals from prisons located in highly endemic areas. Still, the annual cost of coccidioidomycosis care and treatment in California prisons is estimated to exceed \$23 million.¹⁰⁷ Further research into additional methods aimed at reducing the numbers of infected prisoners is needed. One promising strategy may be to risk stratify prisoners via the spherulin skin test (Spherusol, Allered Laboratories, Inc. San Diego, CA, USA, is FDA-approved but not yet commercially available) prior to entry in a prison in a highly endemic region.

Recent epidemiologic trends

The incidence of coccidioidomycosis has risen substantially during the past two decades (Figure 3). In Arizona, this increase first became apparent during the early 1990s, when the annual incidence more than doubled from 7/100,000 persons in 1990 to 15/100,000 in 1995.¹⁰⁸ Coccidioidomycosis became a mandatory reportable disease in Arizona in 1997; the incidence rose from 21/100,000 in 1997 to 37/100,000 in 1999 and continued to rise to 91/100,000 in 2006.⁶¹ Incidence decreased slightly during 2007–2008,⁶⁴ but then increased dramatically during 2009 (154/100,000) to 2011 (248/100,000).¹⁰⁹ Reasons for the observed increase since 2009 are not known, but may be due in part to a major commercial laboratory changing its reporting practices to conform with the other laboratories in Arizona, by reporting positive enzyme immunoassay (EIA) results as cases without confirmation by immunodiffusion.⁶⁴ An enhanced surveillance study suggests that the use of laboratory-only criteria for case reporting, including a single positive EIA test, accurately reflects the burden of disease in Arizona.¹¹⁰ Other reports show that EIA testing has a high false-positivity rate¹¹¹ and more recent evidence has suggested the performance characteristics of EIA testing varies among different patient groups.¹¹²

Diagnostic practices in coccidioidomycosis vary between institutions with some facilities employing highly sensitive but nonspecific testing, such as the aforementioned EIA, while others perform more labor intensive methodologies such as complement fixation and

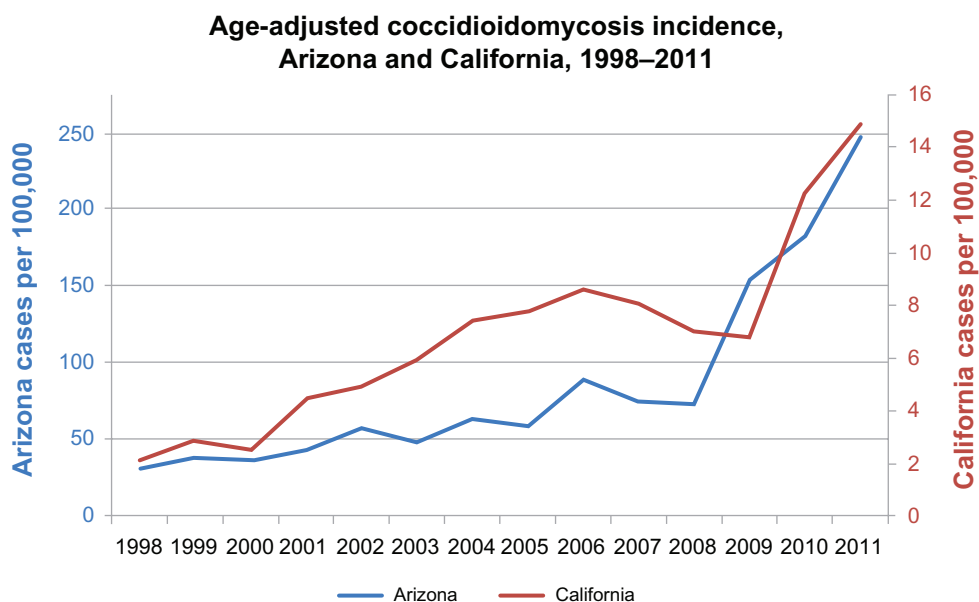


Figure 3 Incidence of coccidioidomycosis in Arizona and California, 1998–2011.

immunodiffusion testing that are available only at reference laboratories.¹¹³ Recently coccidioidal antigen testing has become commercially available, however this test has thus far proven useful only in cases of widely disseminated infection¹¹⁴ and has a poor sensitivity in veterinary samples.¹¹⁵

Although it is likely that differences in testing methodology have contributed in part to a reported increase in Arizona, a similar incidence trend has occurred in California. Average statewide annual incidence increased from 2.5/100,000 during 1995–2000 to 8/100,000 in 2006,⁶⁵ and after a slight decline during 2007–2009, increased during 2010 and 2011 (14/100,000).¹⁰⁹ During this time, the only major change to California's surveillance was a transition to a laboratory-based reporting system in 2010. Coccidioidomycosis incidence also rose substantially during 1998–2011 in other endemic areas (where reporting practices have remained relatively consistent), including New Mexico, Nevada, and Utah,¹¹⁶ perhaps indicating improved awareness of the disease, changes in testing practices, increased travel or relocation to endemic areas, and growth of the “at-risk” immunosuppressed population.¹¹⁷ Other hypotheses include increased spore dispersal due to environmental changes (ie, temperature, moisture) or human activity such as construction.¹¹⁸

Analyses of hospitalization data further suggest that the recent increase reflects an increase in actual disease. In Arizona, the number of persons discharged with a primary or secondary diagnosis of coccidioidomycosis increased from 69 in 1998 to 598 in 2001, with the highest hospitalization rate (29/100,000) occurring among persons ≥ 65 years.¹¹⁹ An analysis of national pediatric coccidioidomycosis hospitalizations showed that rates were stable during 2003–2005 but increased significantly during 2005–2006.¹²⁰ In California, coccidioidomycosis-related hospitalization rates remained fairly stable during 1997–2002, with an average annual rate of 3.7/100,000.¹²¹ A review of California hospital discharge data from 2001–2008, however, showed an overall increase in coccidioidomycosis-related hospitalizations during 2001–2006, peaking at 6/100,000 in 2006 before decreasing in 2007 and 2008.⁶⁴ Overall, data on coccidioidomycosis-related hospitalizations in recent years are limited. Given that one study estimated that over 40% of reported Arizona coccidioidomycosis case-patients require hospitalization,¹¹⁰ this topic deserves further study.

In contrast to the dramatic recent increase in incidence, coccidioidomycosis-associated mortality has remained largely unchanged. Based on death certificate data, the mortality rate in Arizona actually decreased slightly

from 0.9/100,000 in 1996 to 0.5/100,000 in 2005.⁶¹ A review of US death certificates identified a total of 3089 coccidioidomycosis-associated deaths during 1990–2008, but found that mortality rates have remained fairly stable since 1997, with an overall age-adjusted mortality rate of 0.59 per 1 million person-years during the study period.¹²² Possible reasons for the stable mortality rates despite the overall increase in incidence include treatment advancements or a disproportionate increase in primary pulmonary coccidioidomycosis diagnoses compared to the more severe forms of disease.⁶¹ Although mortality rates have remained stable, the number of documented coccidioidomycosis-related deaths is likely underestimated.¹²²

Conclusion

The public health burden of coccidioidomycosis is substantial and has been increasing in recent years. Current prevention messages focus on common-sense methods to reduce exposure to soil or dust where *Coccidioides* is common in the environment, such as wearing a dust mask, wetting soil before participating in soil-disturbing activities, or limiting these types of activities altogether, particularly among people at risk for severe disease. However, it is important to point out that few to no data exist that demonstrate the effectiveness of any of these measures. Wide-scale measures to reduce airborne dispersal of *Coccidioides* such as watering construction sites, paving roads, planting grass, or other vegetation have also been proposed, but these methods are not likely to be overly effective because airborne conidia can travel for miles.

Determination of prior *Coccidioides* exposure to evaluate the risk for disease represents a promising public health strategy. Measurement of coccidioidal cellular immunity using skin tests has been a valuable clinical and epidemiologic tool since the 1940s; however, *Coccidioides* skin test reagents have not been commercially available in the USA for over a decade.¹²³ The return of widespread access to coccidioidal skin tests in endemic areas could help identify and thus better manage specific groups of people at risk for infection.

Because there are no proven methods to prevent coccidioidomycosis, additional research into strategies to reduce the associated morbidity is required. Continued efforts to promote awareness among the public may help to reduce delays in diagnosis and treatment, as evidence suggests that persons with coccidioidomycosis who knew about the disease before seeking healthcare were more likely to request testing and be diagnosed sooner than those who were

unfamiliar with the disease.¹¹⁰ Similarly, increased awareness among healthcare providers about coccidioidomycosis diagnosis and treatment is needed,¹²⁴ especially because the symptoms are often indistinguishable from those of other community-acquired respiratory infections. Admittedly, it is not definitively known whether earlier diagnosis and treatment can lead to improved outcomes, but other benefits of diagnosis, such as reduced anxiety or unnecessary medical treatment or procedures, make recognition of coccidioidomycosis essential.

Further research into the optimal antifungal treatment regimen for coccidioidomycosis is also warranted, particularly with regard to the role of antifungal treatment for primary pulmonary disease.^{2,125} Currently, guidelines from the Infectious Diseases Society of America recommend treatment of primary pulmonary disease in persons who are at risk for developing severe or disseminated disease.² However, the role of antifungal medications in this clinical syndrome is controversial. Some experts recommend treatment of all persons with symptomatic respiratory disease, while others prefer to observe these patients closely. More research is needed to determine if existing or newly-developed antifungal agents can reduce the severity or duration of disease. Finally, efforts to create a preventive vaccine are ongoing;¹²⁶ and if developed, a vaccine could prove to be a cost-effective strategy to reduce the burden of disease among some at risk populations.¹²⁷

Disclosure

The authors have no conflicts of interest to declare.

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