



Coccidioidomycosis: Epidemiology, Fungal Pathogenesis, and Therapeutic Development

Hazael Hernandez¹ · Victor H. Erives¹ · Luis R. Martinez¹

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Abstract

Purpose of Review Coccidioidomycosis can result from the inhalation of infectious spores of *Coccidioides* species (spp.) *immitis* or *posadasii*. Clinical manifestations range from mild flu-like disease to severe disseminated infection that can require life-long therapy. Burden of this mycosis is high in the southwest region of the USA where it is well characterized, and in many areas of Mexico and Latin America where it is inadequately characterized. Here, we provide historical data and current knowledge on *Coccidioides* spp. pathogenesis as well as recent progress in therapeutic and vaccine development against coccidioidomycosis.

Recent Findings The virulence mechanisms of *Coccidioides* spp. are largely unknown; however, production and regulation of a spherule glycoprotein, ammonium production, and melanization have all been proposed as integral factors in *Coccidioides* spp.' pathogenesis. Therapeutic options are limited and not 100% effective, but individualized treatment with triazoles or amphotericin B over the course of pulmonary or disseminated infection can be effective in resolution of coccidioidomycosis. Human immunization has not been achieved but efforts are ongoing.

Summary Advances in therapeutic and vaccine development are imperative for the prevention and treatment of coccidioidomycosis, especially for those individuals at risk either living or traveling to or from endemic areas.

Keywords Antifungals · Coccidioidomycosis · Epidemiology · Vaccine · Valley fever · Virulence

Introduction

Coccidioides species (spp.) are dimorphic fungi, which are present in the environment and can infect and cause disease in humans when arthroconidial spores are inhaled [1]. *Coccidioides* spp. are endemic in the southwest region of the United States (U.S.) and in other semiarid areas in Mexico and Latin America. Disease due to the progression of coccidioidal infection is termed coccidioidomycosis and two species have thus far been identified as etiologies: *C. posadasii* [2] and *C. immitis* [3, 4], the former having been identified more than 100 years after the initial discovery of the fungus [5]. While the two species are genetically different and there may exist

clinically distinct characteristics due to infection that are yet to be identified, no difference in disease progression, diagnosis, or treatment has been established. In fact, pathogenicity of the two species has been proposed to be comparable [6, 7]. Infection with *Coccidioides* spp. commonly remains asymptomatic, but oftentimes presents as pneumonia that can be self-resolving and, in certain cases, life-threatening disseminated mycoses [8]. The public health impact of coccidioidomycosis is significant, with approximately 25,000 reported coccidioidomycosis-associated hospitalizations and over \$2 billion USD in hospital charges in California alone during 2000–2011 [9].

Infection due to *Coccidioides* spp. was first described in Argentina in 1892 by a native doctor who was examining skin lesions on a soldier [5] and shortly after in the U.S. during an examination of a manual laborer from the San Joaquin Valley in California [3] (Fig. 1). After mischaracterization of the etiological agent as a protozoan in 1896 [3], correct taxonomic status of *Coccidioides* as a fungus was established from 1900's animal work involving the inoculation of guinea pigs and rabbits with the coccidioidal arthroconidia which led to coccidioidomycosis and satisfied Koch's postulates [4]. Important insight into the understanding of coccidioidomycosis

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✉ Luis R. Martinez
lmartinez43@utep.edu

¹ Department of Biological Sciences, The Border Biomedical Research Center, The University of Texas at El Paso, 500 W. University Ave., Bioscience Research Building, Room 2.170, El Paso, TX 79968-9991, USA

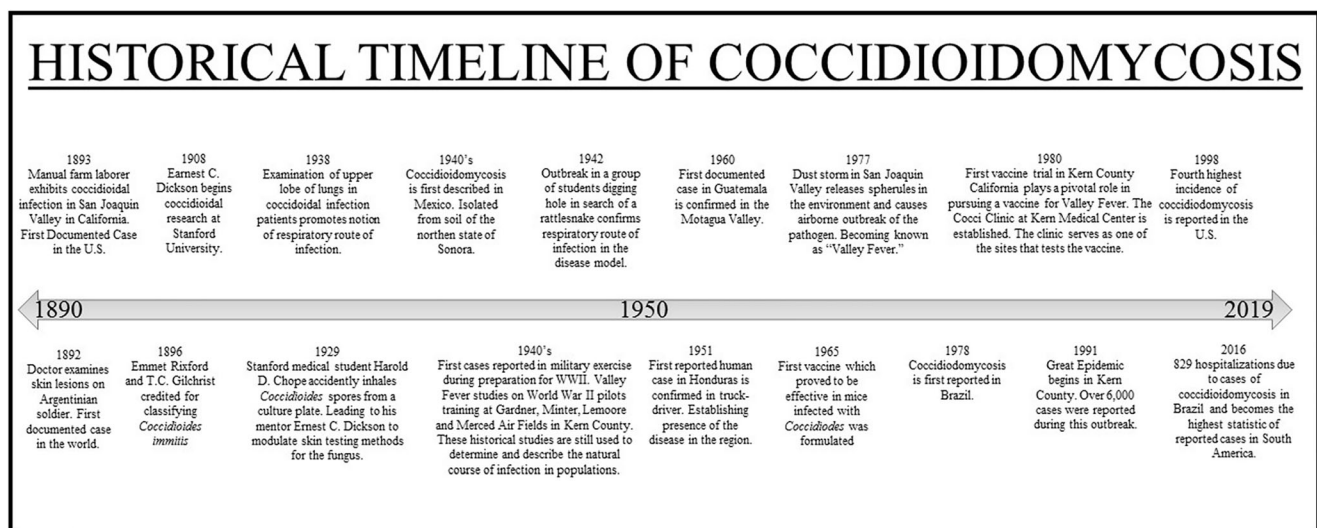


Fig. 1 Historical timeline of coccidioidomycosis

as an environmentally acquired infection developed from an incident in 1929 where a Stanford medical student accidentally inhaled *Coccidioides* spores from a culture plate and developed non-lethal pneumonia [10•]. This case led to a series of seminal findings over the next two decades by Ernest Dickson which, through the use of coccidioidin skin testing methods, robustly established the link between dust exposure, *Coccidioides* infection, and the emerging “valley fever” (“San Joaquin fever” or “desert fever”) in Kern County, California [10•]. Further evidence that supported the proposed respiratory route for infection came from a *Coccidioides* outbreak in a group of students that had dug a hole in search of a rattlesnake in 1942 [11] and from previous descriptions in 1938 of lung upper lobe pneumonia case caused by coccidioidal lung infection [12]. Since then, the relationship of environmental disturbances and the infectious potential of *Coccidioides* spp. has been established and prophylactic and therapeutic solutions are actively being investigated.

In this review, we discuss *Coccidioides* spp. infection and its prevalence, highlighting risk factors and susceptible populations in under-resourced areas. Furthermore, coccidioidal virulence factors along with the current state of vaccination are examined, along with therapeutic strategies in development that hold promise for future treatment. We aim to inform on this neglected medically important fungi while we highlight the necessity for clinician awareness of coccidioidomycosis as a differential diagnosis in endemic areas or for cases involving individuals with recent travel to or from endemic areas.

Coccidioidomycosis: the Desert Dust Disease of the Americas

Coccidioides spp. exclusively inhabit the western hemisphere as thermally dimorphic microorganisms that can exist in

saprotrophic or parasitic growth phases in the environment and in suitable hosts, respectively [13•]. The fungus grows saprotrophically as a mycelium consisting of filamentous hypha networks in soils of semiarid regions such as in the U.S. Southwest or in certain regions of Mexico and Latin America. Maturation of mycelia leads to the formation of single-cell barrel-shaped arthroconidia (spores) that are easily aerosolized by any form of soil disturbance, either natural or anthropogenic. The infectious arthroconidia can then be acquired through respiration and this has been described in many host organisms such as humans and in domesticated animals like dogs, cats, and livestock or alpacas [14, 15]. In non-human primates, exposure to as few as 10–50 arthroconidial spores has been shown to cause disease in 4–6 weeks [16, 17]. Many investigations that challenge animals with arthroconidia use challenge load numbers with a similar order of magnitude. Once inhaled into the warm lungs of a host, arthroconidia initiate the thermal transition to spherules, thus initiating the infectious parasitic phase of the fungus. In vitro, this transition has been demonstrated at 37 °C and 10–14% atmospheric CO₂ in chemically defined media [18] and contact with leukocytes has also been suggested as a stimulant for phase transition in the lungs [19]. The spherules then mature and proceed into the endosporulation stage where they swell into structures with diameters upwards of 100 μm that contain 100–300 single-celled endospores which release upon rupture [13•]. Endospores mature into large second-generation endospore-filled spherules that endosporulate and burst, thereby repeating the fungal parasitic life cycle.

In 60–65% of individuals, acquisition of *Coccidioides* spp. leads to asymptomatic sub-clinical infection. Because acquisition of the fungus happens almost exclusively through the respiratory route, individuals with progression to more severe coccidioidomycosis present with symptoms characteristic of typical pneumonia or influenza such as fever, night sweats,

myalgia, arthralgia, headache, fever, cough, and dyspnea which can develop 1–4 weeks after exposure [20]. Particularly in California, presentation of multiple symptoms is commonly referred to as the “valley fever.” *Coccidioides* spp. are responsible for 17–29% of community acquired pneumonia in endemic regions [21] and many have stressed the importance of its strong consideration as a differential diagnosis in endemic areas [22, 23••]. Disseminated coccidioidomycosis occurs secondary to primary pneumonia in approximately < 5% of individuals infected [24] and coccidioidal meningitis is the most serious form of systemic disease [25]. Disseminated infection is typically treated with administration of amphotericin B or azole antifungals [26]. In addition, dissemination can result in cutaneous coccidioidomycosis secondary to acute infection whose manifestations mimic that of dermatologic diseases [27]. In very rare cases, primary cutaneous coccidioidomycosis can result after acquisition of fungal spores through exposed skin, although only about 30 of these cases have been reported since 1926 [27–29] making this the least common type of *Coccidioides* spp. infection manifestation [28]. For patients with primary cutaneous coccidioidomycosis, prognosis is excellent with proper treatment typically leading to complete resolution [28]. Skin rashes and arthritis can also result [24].

For populations in endemic regions, coccidioidomycosis is a fungal threat that warrants substantial notice yet it is regularly overlooked. Acute pulmonary coccidioidomycosis presents similar symptoms to other respiratory illnesses, resulting in many cases that are likely unnoticed or misdiagnosed [23••]. This is especially true in under-resourced communities or in populations with impaired access to medical care [23••]. Lack of state/nationwide skin or soil testing to determine the endemicity range and prevalence also participates in keeping the fungus’ true threat potential in obscurity. Therefore, a concerted effort should be made by public health officials, community leaders, and clinicians in all endemic regions to increase awareness and preparedness in order to decrease the burden of this insidious disease.

Coccidioidomycosis: Prevalence and the Impact on Public Health

The ability of the arthroconidia to remain as viable infectious spores for years under dry conditions is a significant contributor to *Coccidioides* spp.’ prevalence in endemic areas. Endemic regions all have similar environmental profiles and they have been characterized by their arid or semiarid climate, gentle winters, alkaline soil, and low precipitation [30•]. Environmental or anthropogenic disturbances can then aerosolize the resilient arthroconidia, and human hosts can acquire the spores through inhalation. Two *Coccidioides* spp. with variable geographical ranges cause infection, *C. immitis* and

C. posadasii. *C. immitis* is the predominant species in California and Baja California while *C. posadasii* predominates in the rest of the southwestern states. *C. posadasii* is also responsible for most, if not all recorded cases in Mexico and the rest of Latin America although there are sparse reports of *C. immitis* in locations like Venezuela, Argentina, and Colombia [23••, 31]. However, the possibility of *C. immitis* being endemic to those regions remain uncertain as the possibility of infection due to travel to and from endemic areas could not be excluded [23••]. Genetic analysis suggests that *Coccidioides* was introduced into South America from North America by mammals between 9000 and 140,000 years ago [32].

Dating back to the twentieth century, skin tests for the coccidioidal antigen coccidioidin have been used extensively to determine the endemic geographic ranges of the fungus in the environment [10•, 13•]. Areas surrounding established endemic regions are commonly designated as suspected endemic. In the U.S., many early studies using this method established Arizona and the central valley in California as the most endemic regions and endemicity has now also been established in Utah, Nevada, New Mexico, and Texas where the fungus is becoming an emerging threat. Similar skin testing surveys have been conducted in Latin America to establish the fungus’ geographic range. However, most regions with or without endemic *Coccidioides* spp. lack internal comprehensive investigations and surveys that would establish clear geographical boundaries for the fungus. Thus, the extent to which the fungus is present in the environment of the Americas remains highly speculative. To date, one region with thousands of new yearly cases of autochthonous coccidioidomycosis remains the most thoroughly characterized: the U.S. Southwest.

Coccidioides in the U.S. Southwest

With over a quarter of the U.S. population inhabiting the mainland Southwest, disease from fungal etiologies endemic to the region should not be overlooked, especially *Coccidioides* spp. Morbidity due to coccidioidomycosis in hospitalized patients in the U.S. Southwest is significant, resulting in loss of income and a potentially altered quality of life. In California, one estimate places the average lifetime medical costs associated with treatment at \$57,000 USD per patient across all disease manifestations [33••]. Patients with uncomplicated pneumonia incur ~\$22,000 USD lifetime costs, while those with diffuse/chronic pneumonia incur ~\$132,000 and individuals with disseminated disease including meningitis incur over \$1,000,000 USD [33••]. Moreover, about 75% of symptomatic patients miss work or school [34] for nearly a week when manifestation is uncomplicated pneumonia [33••]. In patients, with disseminated coccidioidomycosis, 10% of individuals have permanent work loss, while the remaining 90% lose an average of 90 work days [33••]. In

California, coccidioidomycosis has emerged as a significant fungal threat in the past few decades with 75% of the state's cases occurring in the San Joaquin Central Valley [35]. From 1998 to 2011, documented cases steadily increased by about 13–16% each passing year, culminating in almost a 1000% increase of reported cases, up from 2271 in 1998 to 22,641 in 2011. The incidence of coccidioidomycosis attenuated to 8232 cases in 2014 but increased since then to 14,364 cases in 2017 [36]. While valley fever gets its name from California, the burden of coccidioidomycosis has historically been highest in Arizona, where approximately two-thirds of all infections were reported up until 2015 [36]. In 2016, only about 51% of cases were reported from Arizona and, in 2017, more cases were reported in California than in Arizona, ending the trend of Arizona having higher incidence for coccidioidomycosis [36]. In total, 14,364 cases were reported in 2017 making it the year with the fourth highest incidence since 1998 [36]. The severe threat that re-emerging coccidioidomycosis poses is being responded to by the National Institutes of Health (NIH). Recently, the NIH posted funding opportunity announcements in support of research activities contributing to the understanding of coccidioidomycosis [37].

Interestingly, natural disasters have been shown to affect the environmental release of fungal spores [38, 39]. Coccidioidomycosis has been associated with a few instances of post-disaster infection. In 1994, a magnitude 6.7 earthquake struck Northridge, California. In conjunction with secondary seismic activity, dust clouds were formed that were strongly implicated in contributing to an increase in aerosolized *C. immitis* spherules in the environment resulting in 203 cases of coccidioidomycosis in Ventura County, CA [40]. A similar airborne outbreak occurred previously in 1977 in the San Joaquin Valley [41] when a dust storm was implicated in 115 valley fever cases including 16 disseminated cases in non-endemic Sacramento county, 18 cases at a U.S. Navy air station in Kings County [42], and 117 more cases in Kern county than the previous year during the months counted [43], as well as many others across other Californian counties [38]. Worryingly, dust storm activity has intensified over the past few decades and the frequency of dust storms has been found to be correlated with valley fever cases [44].

Just as dust storms in California left an increase of coccidioidomycosis cases in its wake, it has been suggested that incidence of infection in Arizona can be predicted by other natural processes such as seasonal precipitation [45]. Human disturbance of soil through industry has also been suggested to possibly increase *Coccidioides* infectious potential [46]. Similarly, coccidioidomycosis cases have been reported after military exercises [10•] and construction projects [47]. Climate change has also been proposed as a driver of fungal proliferation in the environment and expansion of the endemic range of *Coccidioides* spp. [48] and the influence of climate change on valley fever is being increasingly analyzed [49]. In

order to develop an appropriate public health system response, anthropogenic disturbances of soil caused by construction or climate change that affect the incidence of fungal disease like valley fever should continue to be investigated thoroughly.

In all southwestern states, there are areas purported to be highly endemic, established endemic, or both. However, many surrounding areas to the endemic states are also suspected to be endemic and should not be ruled completely safe, especially for those individuals with higher risk for infection. In fact, recently, the fungus has been identified in soil in Oregon [50] and cases have been reported as far north as Washington State in the Northwest of the U.S. [51]. Moreover, valley fever has been diagnosed in workers at Dinosaur National Monument [52] which spans the northeastern reaches of Utah and extends into Colorado. This and other cases may indicate that the endemic range for this fungus is increasing. It is suspected that pathogenic coccidioidal spores are found in the soils of many arid areas throughout North America, and their spread to new areas should be considered in the future. The overall increase of coccidioidomycosis in the southwest is troubling, and while the reason(s) have not been fully elucidated, *Coccidioides* spp. are emerging as a significant fungal threat to public health.

Coccidioides in Latin America

Coccidioidomycosis is of particular importance in Latin America due the large amount of individuals involved in agriculture and thus potentially at higher risk of exposure to aerosolized fungi [23]. In addition, significant segments of the population are medically underserved and resources for detection and/or treatment of coccidioidomycosis can be scarce or unavailable leading to high burden of disease. Likewise, clinical and epidemiologic data can be sparse [23••, 31], which can result in underrepresentation of coccidioidal burden and potential unpreparedness for future infections. *C. posadasii* remains the most prevalent coccidioidal species in Mexico and the rest of Latin America [23••, 53]. In Mexico, coccidioidomycosis was first described in the mid-1940s [54, 55] and later isolated from soil in the northern state of Sonora [56]. Skin surveys during this time established endemicity in areas with soil profiles similar to soil in the U.S. Southwest [23••]. Therefore, much of the Mexican research focus has been on northern states with deserts, which lie directly across the border from endemic regions in the U.S. such as the states of Baja California, Sonora, Chihuahua, Coahuila, and Nuevo Leon [57–59]. In Latin America, Mexico has the highest incidence of coccidioidomycosis where it is known to be as prevalent as in endemic regions in the U.S. [60]. The current burden of coccidioidomycosis in Mexico is unknown; however, up until its dismissal as a reportable disease in 1994, an average of 1500 cases were reported yearly [23••]. For example, in a study involving

668 individuals from rural and urban communities in Coahuila, skin tests revealed that 93% of individuals were positive to coccidioidin [58, 61].

In Central America, *Coccidioides* spp. are also characteristically endemic in arid and semiarid regions of Honduras and Guatemala [2, 23••]. Two early skin studies in Guatemala and Panama in 1945 and 1950, respectively, indicated less than 1% positive reactivity to coccidioidin in tested individuals [62, 63]. In 1951, coccidioidomycosis in a truck driver from Honduras was described, becoming the first such case in the region and thereby establishing its presence in Honduras [64]. In Guatemala, the first human case was reported in 1960 [65, 66] and the fungus' endemicity was established in the Motagua Valley in western Guatemala based on evidence from an investigation on 6 individuals with the disease. This observation was supported by further findings from skin tests performed on ~ 10,000 individuals from the valley in the late 1960s [67].

In South America, coccidioidomycosis is also not reportable and studies are limited making the true incidence uncertain. Reported cases and skin test surveys that have been conducted in Colombia, Venezuela, Brazil, Paraguay, Bolivia, and Argentina are reviewed in detail [23••] with infection rates ranging from 2 to 46%. Five cases have been reported in Colombia from 1958 to 2015 and 114 in Venezuela from 1948 to 2004 [23••]. In Brazil, coccidioidomycosis was first reported in 1978 and studies have been conducted on incidence in specific states since then [31, 68]. In 2016, the nation's burden of coccidioidal disease was carefully elucidated and 829 hospitalizations due to coccidioidomycosis were described [69], making Brazil the South American country with the highest reported number of cases. However, this statistic might be misleading and indicative of underreported coccidioidomycosis in other South American countries rather than a greater disease burden in Brazil. Mycology is an extensively studied field in Brazil, and the nation is host to many excellent fungal investigators and clinicians, and this might account for the greater number of reports coming from the country. Ultimately, just like in the U.S., *Coccidioides* spp. are likely to be found in many areas that are yet to be described in scientific literature and it is probable that they might also be expanding into neighboring regions. A concerted effort by health officials in nations with reported cases might lead to a better understanding of coccidioidomycosis' impact on Latin America.

Risk Factors for the Acquisition of Coccidioidomycosis

Since *Coccidioides* spp. and their intimate association with the environment began to be characterized in the twentieth century, much knowledge has been acquired about related risks for

infection with the fungus due to environmental exposure [30•]. Unfortunately, significant insight has been gathered directly from outbreaks in individuals exposed to dust in endemic regions. For example, it is now known that there are very high rates of exposure and coccidioidomycosis among incarcerated individuals in prisons, especially in California [70, 71]. While officials have made an effort to address the issue, one study found that health costs due to *Coccidioides* infection in California prisons were approximately \$23 million USD from 2006 to 2010 [72]. Individuals participating in outdoor activities such as hunting, sports, or construction are also at higher risk for infection [47, 73–75]. Furthermore, many outbreaks have arisen as a result of military exercises with some of the first cases reported in the 1940s during military exercises when troops were preparing for overseas battle in World War II [10•, 76].

The increase in globalization and fast travel is also an important consideration as patients might present with coccidioidomycosis in a non-endemic area after travel to an endemic area [77–79]. In fact, the incidence of reported cases in the U.S. outside of recognized endemic zones has increased in the past few decades [77, 80–83]. In Canada, a high number of reported cases in Ontario have even prompted reviews that investigate the impact of fungi on the city [84]. Outside of the western hemisphere, coccidioidomycosis has been observed in recent travelers of endemic zones returning home to Europe [85–87], Australia [88–90], and Asia [91–94]. The threat of coccidioidomycosis is thus a very important consideration for traveling individuals with a propensity for coccidioidal infection, such as immunocompromised individuals. Recent travelers to endemic areas that acquire coccidioidomycosis with more serious manifestations require medical assistance in their areas of residence and it is up to health care providers to be alert for coccidioidomycosis in the differential diagnosis, particularly in the setting of a pneumonia or other invasive infection non-responsive to antibacterial agents.

Gender is a determinant of risk to acquire coccidioidomycosis with men having higher rates of infection and dissemination [30•]. Although coccidioidomycosis occurs in all age groups, adults aged 40 and older have demonstrated a higher incidence for the disease [30•]. Moreover, an increased rate of dissemination has been described in non-Caucasian groups such as Hispanics, African Americans, and especially Filipinos [43, 76, 95]. Furthermore, American Indians and Alaskan Natives experience high coccidioidomycosis-associated hospitalization rates and high morbidity [96]. Medical status of individuals also plays a role, with pregnancy, diabetes, and cardiopulmonary disease being associated with infection and at least pregnancy with disease severity. Immunocompromised individuals are recognized to be at greater risk for an opportunistic infection by *Coccidioides*. In patients with HIV/AIDS, diminished CD4⁺ T cell counts is the greatest risk determinant, as there is an inverse

correlation between the risk for disseminated coccidioidomycosis and total CD4⁺ counts [97, 98]. Organ transplant recipients [30, 99, 100] and patients on immunosuppression with high-dose corticosteroid [30] or receiving a TNF- α antagonist [101] are also at increased risk for coccidioidomycosis.

Coccidioides spp. Virulence Factors

Coccidioidal arthroconidia that are aerosolized through environmental or anthropogenic means initiate infection after inhalation by a host. Once in the host respiratory system, arthroconidia undergo the thermally induced switch to the pathogenic-“parasitic” phase of the fungus’ life cycle. Fungal spherules begin isotropic growth and hundreds of endospores begin to differentiate within the large spherule. As coccidioidal spherules mature in these initial stages of pulmonary infection, a membranous layer termed the spherule outer wall (SOW) composed of polysaccharides, lipids, and proteins is shed from the spherule [102]. Isolated SOW from cultures of *Coccidioides* demonstrate high immunoreactivity to anti-coccidioidal antibodies isolated from patients [103]. SOW glycoprotein (SOWgp) has been established as the immunodominant molecule within SOW fractions responsible for this in vitro reactivity and for eliciting a human immune response to coccidioidal infection [104, 105]. A defining characteristic of SOWgp is that it contains proline/aspartate-rich motifs that tandemly repeat 3–6 times, depending on the clinical isolate [106]. Concerted evolution of these tandem sequences has been proposed as a mechanism that could potentially enhance the fungi’s ability to evade host immune responses [107]. Also, proline-rich regions (PRRs) have been proposed as key motifs in proteins that participate in the adhesion of many microorganisms to host tissues [108, 109]. This has been demonstrated in fungi containing glycoproteins with PRRs such as in *Colletotrichum lindemuthianum* [105, 108] and *Candida albicans* [109] and in the bacterium *Streptococcus mutans* [110] which contains a cell surface adhesin with a PRR.

In vitro adherence assays performed to investigate recombinant SOWgp’s role as putative adhesion have demonstrated the glycoproteins’ ability to bind to mammalian extracellular matrix (ECM) components such as laminin, fibronectin, and collagen [105]. During inhalation and infection, these types of ligands might mediate adherence of the fungus to host respiratory tissues. In fact, in a SOWgp knockout strain of *C. immitis*, spherules display a 30–50% reduction in their ability to bind to fibronectin and laminin. Furthermore, in an experiment evaluating the effect of SOWgp gene deletion on mice survival, the mutant without SOWgp showed decreased virulence. Animals challenged intranasally with *C. immitis* with the SOWgp gene died after 21 days while mice infected with the *C. immitis* SOWgp knockout had significantly greater

survival, with 58% of animals surviving beyond 40 days post-infection. SOWgp is an established antigen suggesting that the loss of virulence in the SOWgp *C. immitis* mutant might be attributable to the fungus’ decreased ability to bind to host tissue or, alternatively, due to the loss of SOWgp as an antigen, which may influence a strong immune response by the host resulting in collateral tissue damage.

Expression of SOWgp has been shown to be cyclic: prevalent during early spherule development but significantly decreasing during endosporulation [105]. During pre-endosporulation, the maturing spherule releases copious amounts of antigenic SOWgp, making the fungus vulnerable to opsonization and phagocytosis by leukocytes. When the spherule bursts, spherule contents and endospores that are the appropriate size to be phagocytized are released, initiating an intense host inflammatory response. Studies have shown that the fungus combats this with the production of a metalloproteinase (Mep1) [111] that digests SOWgp, effectively enhancing the endospores’ ability to evade opsonization and subsequent phagocytosis [112]. In mice immunized with SOWgp, high antibody titers were observed and survival after challenge with *mep1* knockout arthroconidia significantly increased compared to immunized mice challenged with wild-type arthroconidia. Moreover, in vitro exposure of spherules (devoid of SOWgp after stripping by the action of purified Mep1) to alveolar macrophages resulted in enhanced phagocytosis and killing of the fungus.

Like in the medically relevant *Cryptococcus* spp., coccidioidal release of ammonia during infection is also implicated in the virulence of this fungus [113, 114]. *Coccidioides* spp. respond to pH changes in their environment. For example, when the fungus is grown in acidic cultures in either the saprobic or parasitic phase, the fungus responds by secretion of NH₃/NH₄⁺, resulting in an increase in the pH of the culture. The enzyme responsible for production of a large portion of cellular ammonia is urease [115, 116], which catalyzes the hydrolysis of urea into two ammonia molecules that neutralize acid by spontaneously protonating to form ammonium at physiological pH [117]. Ureases have been established as determinants of pathogenesis in other microorganisms [118] such as in *Helicobacter pylori*, which is able to use ureases to acclimatize to the acidic conditions of the gastric milieu [119]. These bacterial ureases have been implicated in exacerbated tissue damage by accumulation of ammonium in infected tissues [113]. A similar mechanism of damage has been proposed for *Coccidioides* spp. and studies have provided evidence corroborating this hypothesis [113, 120]. In one investigation, 55% of mice challenged with a *C. posadasii* strain lacking the urea gene survived beyond 50 days post challenge while a wild-type strain challenge resulted in 100% mortality by day 18 [113]. The ureidoglycolate hydrolase enzymatic pathway has been proposed as another source of coccidioidal ammonia [113]. In this study, a

challenge in mice performed with a mutant strain in urea and ureidoglycolate hydrolase showed 90% survival after 30 days.

Coccidioides spp. also produce melanin-like pigments [121] similar to those observed in other medically relevant fungi such as *Cryptococcus neoformans*, *Aspergillus* spp., *Exophiala dermatitidis*, *Sporothrix schenckii*, *Histoplasma capsulatum*, *Paracoccidioides brasiliensis*, *Scedosporium prolificans*, and *Blastomyces dermatitidis* [122–126]. Melanization is a key contributor towards fungal virulence and environmental persistence with extensive studies demonstrating that this dark pigment on the cell wall protects fungi against extreme temperature, UV light, toxic metals, antimicrobial peptides, antifungals, and nitrogen/oxygen oxidants [127]. While the impact of melanization has been described in other fungi, further studies of *Coccidioides* spp. need to be conducted to further elucidate the extent to which melanization enhances coccidioidal environmental persistence and pathogenicity.

SOWgp production, SOWgp dysregulation by the action of Mep1, urease activity, and melanin production are four key means by which this fungus can adversely impact/damage infected hosts. Three of these virulence mechanisms and their concerted contribution to the overall virulence of *Coccidioides* spp. have been extensively reviewed [106•]. With the completion of *Coccidioides* spp. sequencing projects and the advent of newer genetic manipulation techniques, there surely is promise that novel virulence mechanisms of this pathogen will be elucidated aiding in the development of prophylactic and therapeutic strategies to combat coccidioidomycosis.

Therapeutic Approaches to Fight Coccidioidomycosis

As with other fungi, there is currently no therapy that results in resolution of coccidioidomycosis in all cases. Patients with coccidioidomycosis that proceeds to a more severe form are typically burdened with high morbidity and often mortality. Antifungal therapy is expensive and hospital costs can quickly accumulate especially if intensive care is needed. Therefore, it is imperative to find the most effective treatment with minimal acquired damage to tissues and organs. The majority of patients that become infected with *Coccidioides* spp. remain asymptomatic or infection only leads to a mild manifestation. Thus, primary coccidioidal pneumonia in immunocompetent individuals will usually resolve without medical intervention or administration of antifungal therapy [26]. In fact, studies of antifungal administration during early pulmonary infection show no evidence in support of such treatments [26, 128, 129]. In addition, no randomized clinical studies exist that justify early therapy in healthy hosts [26, 130]. In fact, practice guidelines have suggested that treatment of early primary coccidioidomycosis should be highly individualized [26, 131]. For example, in immunocompetent patients, azole therapy might not be

necessary; however, periodic reassessment might be appropriate to ensure resolution of mycosis or identify complications. In patients with an overt pulmonary manifestation, biopsy or radiological examination might be performed to rule out malignancy. In patients with hydrocephalus due to disseminated meningeal mycoses, surgical implementation of a shunt may be required [26]. However, these medical devices are at extremely high risk for infection and clogging, which can exacerbate morbidity. In pregnant mothers, alternative therapies might be pursued due to the possible teratogenicity of azoles. Nevertheless, recent findings have determined that this might only be the case in the first trimester of pregnancy and, even then, low doses during first trimester might be an option [132]. Thus, administration of azoles might be appropriate during second and third trimesters, if so determined by a clinician [26, 131]. Initiation of antifungal therapy has proven to be effective in many cases, but there is no guarantee that infection will not return after discontinuation. Thus, antifungal therapy is often indefinite and lifelong, especially for more advanced coccidioidomycosis [26, 131, 133].

The pharmacology and treatment of coccidioidomycosis has been excellently reviewed [134••]. Like in the treatment of other fungal infections, triazole antifungal drugs (fluconazole, itraconazole, voriconazole, and posaconazole) and amphotericin B are the agents administered to counter coccidioidal infections [134••]. Triazoles are largely fungistatic and they exert their effect through inhibition of the ergosterol synthesis pathway in fungi which leads to destabilization of membrane-associated enzymes and eventual inhibition of fungal growth. Fluconazole is the most commonly administered antifungal drug for treatment of coccidioidomycosis as it is relatively inexpensive and available in intravenous or oral preparations [134••]. The drug has excellent oral bioavailability and it is unaltered by food or gastric conditions. Furthermore, fluconazole is not protein bound and distributes widely into most body tissues and fluids such as the central nervous system (CNS) [135, 136]. This is especially useful for the treatment of disseminated coccidioidomycosis in the CNS. In vitro, fluconazole exhibits higher minimal inhibitory concentrations (MICs) in *Coccidioides* spp. compared to other azoles [137]; however, this has not been rigorously corroborated in patients [134••]. Itraconazole is also commonly administered [134••] and some evidence has shown that its use is superior in the treatment of some disseminated forms of coccidioidomycosis [138]. Reduced relapse rates after discontinuation of therapy with itraconazole compared to fluconazole have also been observed. Itraconazole is available in capsule and oral solution formulations, as well as intravenously in some countries outside of the U.S., and the different forms of the drug exhibit differing bioavailability and dispersal in the body. After the development and introduction of fluconazole and itraconazole in the 1990s, there remained a need for second-generation triazoles to account for fluconazole's limited spectrum of antifungal activity and itraconazole's absorption limitations [139]. This led to the

development of voriconazole and its approval by the U.S. Food and Drug Administration (FDA) in 2002. Voriconazole is a synthetic derivative of fluconazole which contains a fluorinated pyrimidine and an α -methyl group and exhibits expanded activity. There is limited experience with the use of voriconazole in the treatment of coccidioidomycosis and the drug is used in patients who are intolerant or refractory to the other azoles [134••] and has been effective in such patients [140]. Voriconazole has also been used to successfully treat coccidioidal meningitis [140, 141]. Although voriconazole exhibits excellent bioavailability and distribution properties, it is also toxic [142, 143]. Posaconazole is another azole with limited clinical experience for coccidioidomycosis. It is available intravenously or by tablet. In animal models, the drug has been found to be the most active azole [144, 145] and the drug has also demonstrated superior sterilization in tissues in comparison to itraconazole. Large clinical trials with posaconazole have not been conducted, and thus far, no clinical benefit over other triazoles has been established [134••].

Amphotericin B, a polyene, is usually reserved for severe cases of coccidioidomycosis [26]. Amphotericin B binds irreversibly to ergosterol, resulting in disruption of fungal membrane integrity and ultimately cell death. Before the development of the triazoles, amphotericin B was the primary antifungal agent used in the treatment of coccidioidomycosis. With the advent of triazole treatment, amphotericin B is usually reserved as the final option in treatment, or for widely disseminated coccidioidomycosis [146] because the drug exhibits high toxicity due to off target binding to cholesterol in human cells. The drug has proven efficacy in some patients with disseminated coccidioidomycosis to the CNS when applied intravenously but it shows low penetration into the CNS.

All of the therapeutic options available for the treatment of coccidioidomycosis have positive and negative aspects regarding cost, bioavailability, tolerability, penetration, toxicity, and pharmacokinetic/pharmacodynamics characteristics. Therefore, as [134••] points out, there is an existing need for therapeutics that improve on adverse characteristics of the currently available clinical options [134••]. Currently, there are several novel agents in preclinical development such as nanoparticle and cochleate formulations of amphotericin B [147], itraconazole formulations with enhanced absorption SUBA-itraconazole [148], glucan synthase inhibitors [149], and chitinase inhibitor nikkomycin Z. Ultimately, the search is ongoing for agents to battle *Coccidioides* spp., and it is necessary due to the fungus' resurgence and infectious potential.

State of Developing a Coccidioidal Vaccination

Similar to other fungal etiologies, thus far, there is currently no vaccine for prophylactic immunization against *Coccidioides* spp. However, patients that recover from coccidioidomycosis

typically acquire lifelong immunity, indicating that development of a vaccine that can achieve similar results is feasible. At the moment, we do not have a comprehensive understanding of coccidioidomycosis' pathogenesis, but great efforts have and are being made to identify strategies that can be used in the development of an effective vaccine. In the 1960s, the first vaccine which proved to be effective in mice infected with *Coccidioides* was formulated [150]. Mice were immunized with formalin-killed spherules (FKS) and this was later shown to reduce the severity of disease in a primate model. The vaccine eventually moved to double-blind phase III clinical trials, but its progress was stopped after the killed spherule vaccine did not demonstrate significant reduction of incidence or severity in the vaccinated group in humans and it induced an over reactive immune response [151]. In 2007 and 2009, another live-attenuated vaccine began to be investigated [152] when heat-killed *Saccharomyces cerevisiae* was reported to protect against aspergillosis and candidiasis, thus prompting investigation on the effect of the vaccine on coccidioidomycosis. It was ultimately determined that heat-killed *S. cerevisiae* conferred less protection than formalin-killed spherule vaccination and subcutaneous immunization with a fraction derived from mechanically derived spherules [153].

With the advent of more contemporary techniques and a more advanced understanding of immunology and *Coccidioides* spp.' genome, identification of potential antigens that can be recombinantly produced and used for immunization have also become the focus of investigations. Since then, a large number of candidate antigens have been discovered [154] and their efficacy after combination with adjuvants are also being evaluated. Furthermore, two live-attenuated vaccines have been isolated and evaluated in murine models. Both attenuated mutant strains used for the vaccine formulation were created by knocking out chitin-related genes that disable the microorganisms' ability to undergo transformation from its saprobic form to the parasitic or its ability to endospore within the lungs. To test whether the mutants lacked virulence, mice were infected with an extremely high dose of arthroconidia (5000 spores) that is orders of magnitude higher than that necessary for death. All the mice survived, enabling researchers to investigate their immunization potential [155]. For instance, mice immunized with 7500 arthroconidia had 100% survival after 75 days with a challenge with a lethal dose of non-attenuated arthroconidia. While these results are promising, severe hurdles with the FDA have to be overcome before these strains can be used in trials with humans. Further studies focusing on the immune responses generated by live-attenuated vaccination and safety issues associated with delivering live-attenuated coccidioidal strains particularly in immunocompromised individuals are necessary [156]. However, these findings provide a proof-of-concept study that opens a novel area of research and a potential therapeutic strategy to prevent and reduce the devastating

consequences of coccidioidomycosis, particularly in individuals living in endemic regions.

Conclusion

Coccidioidomycosis is an emerging threat to human health on two continents in the western hemisphere. While the burden of this disease might at times show patterns of attenuation, current infection trends indicate that the prevalence of coccidioidomycosis is increasing in magnitude and that the causative agent is evolving and expanding outside of historically endemic regions. In the southwest region of the U.S., coccidioidomycosis is an overt threat, as it affects local populations at epidemic levels. Many of the U.S. most populous cities are in the Southwest, which means that tens of millions of individuals that already reside there are potentially at risk. Moreover, the southwest population is increasing rapidly [157] and 6 out of the 10 fastest growing cities in the U.S. reside in this region. As many individuals migrate to the U.S. Southwest, there is coccidioid risk to individuals without previous exposure to *Coccidioides* spp. Due to the increasing population density, there is also an associated increase in agricultural, commercial, and industrial projects being carried out, which only serves to exacerbate the propagation of *Coccidioides* spp. and thus their burden of disease. In Latin America, the threat of coccidioidomycosis is more insidious in nature as it is likely to be a far more significant threat than scientific or clinical records indicate. Socio-economic conditions and the current status of health care access in Latin American regions present significant challenges in the diagnosis and management of coccidioidomycosis.

In the past few decades and due to global climate change, evidence indicates that there has been an increase in natural disasters. Climatic phenomena such as dust storms, hurricanes, mudslides, unaccompanied high winds, and tornados are all drivers of aerosolization. An increase in events such as these should raise an increase in concern, especially since fungal outbreaks, particularly coccidioidomycosis, following a natural disaster have been observed in areas affected by such events. Although the threat of coccidioidomycosis is considerable, significant advances have been made in endeavors with a mission to ameliorate its burden. Investigations into *Coccidioides* spp. virulence and pathogenesis are rigorous and ongoing, and now there is special emphasis in drug and vaccine development for prevention and treatment of this neglected mycosis. Continuing support of these scientific endeavors and an increased effort for public education on the topic will surely result in decreased disease burden and an eventual resolution to coccidioidomycosis.

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Compliance with Ethical Standards

Conflict of Interest The authors declare that they have no conflict of interest.

Human and Animal Rights and Informed Consent This article does not contain any studies with human or animal subjects performed by any of the authors.

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