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Pathogens of Medical Importance Identified in Hospital-Collected Cockroaches: A Systematic Review

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Abstract: Cockroaches serve as mechanical vectors for medically important pathogens, and their presence in hospitals is a common occurrence. Describe the pathogens and their resistance mechanisms carried by cockroaches collected in hospitals in different parts of the world during the period 2000-2023. This will be achieved through a systematic review of the indexed literature. The studies were identified through a search of the Google Scholar search engine and in electronic databases related to health: LILACS, Scielo Regional, and PubMed. The search strategy was conducted in accordance with the principles for systematic reviews and meta-analysis (PRISMA). The review encompassed 22 studies conducted during the specified period. Blattella germanica and Periplaneta americana are referenced in 15 and 13 of the analyzed studies, respectively. A variety of pathogens, including bacteria, fungi, protozoa, helminths, and rotavirus, were isolated and identified from the external and internal body parts of cockroaches. Bacteria represent the most frequently identified group. The species that appear most frequently in the selected articles are Escherichia coli (11 articles), Staphylococcus aureus (9 articles), Klebsiella pneumoniae (7 articles), and Pseudomonas aeruginosa (6 articles). E. coli and S. aureus bacteria were found to be resistant to antibiotics in 19% and 12% of articles, respectively. Conventional techniques, including seeding in culture media, Gram staining (GS), conventional biochemical tests (CBT), direct parasitological methods, and disc diffusion, have historically been the primary methods for identifying microorganisms and determining antibiotic susceptibility profiles in bacteria. However, there is a paucity of studies that use molecular techniques for bacterial identification and resistance mechanism detection. The identification of pathogens carried by cockroaches collected in hospitals suggests a potential risk of these insects in the transmission of healthcare-associated infections.

Keywords: cockroach; bacteria; hospital; pathogenic parasites; antibiotic resistance

1. Introduction

Cockroaches are ancient insects, with fossil evidence indicating that their lineage can be traced back to the Upper Carboniferous period. [1] Despite the passage of time, modern species have

retained a high degree of morphological and physiological similarity to their ancestors. These insects possess the capacity to adapt to a diverse array of habitats, a trait that has enabled their survival in a multitude of environments. [2-4]

The majority of these insects inhabit tropical and subtropical regions of the globe. [4,5] Approximately 4,600 species have been identified, with over 99% classified as non-domestic. [6] Two notable examples are *Blatella germanica* (Linnaeus, 1767) and *Periplaneta americana* (Linnaeus, 1758) which are among the most common synanthropic cockroaches due to its abundance and cosmopolitan distribution. [5,7-9] The species in question inhabits a variety of environments that provide optimal conditions for its survival, including residential buildings, commercial premises, healthcare facilities, educational institutions, and food-handling establishments.10 These insects are omnivorous, with a preference for foods rich in starch and sugars. [3]

The World Health Organization (WHO) considers cockroaches a significant public health concern due to their role as mechanical vectors, capable of carrying potential pathogens on external and internal parts of their bodies. These pathogens include bacteria, fungi, protozoa, helminths, and viruses. [3,5,11]

The contamination of food, surfaces, and objects of human use with pathogens carried by cockroaches occurs when these insects come into contact with them and regurgitate portions of their partially digested food or deposit their feces on them. [3,12,13] Pathogens from these insects can cause diseases such as bacterial dysentery, giardiasis, amebiasis, and toxoplasmosis, as well as symptoms such as nausea, abdominal pain, vomiting, and diarrhea. [14,15] The presence of cockroaches is a common occurrence in hospitals, which raises concerns about the potential adverse implications for the health of patients and workers. [10,16-18]

In this regard, the World Health Organization (WHO) reports that healthcare-associated infections (HAIs) are responsible for approximately 40,000 deaths annually. It is estimated that 25% of HAIs occur in developing nations and between 5% and 15% in developed countries.[19] The predominant group of pathogens that cause these types of infections are bacteria, many of which are resistant to multiple antibiotics and have been isolated and identified from the bodies of cockroaches. [16,20-23]

The phenomenon of antibiotic resistance (AR) represents a significant global public health concern, with documented associations with elevated mortality rates, prolonged hospitalizations, diminished therapeutic options, increased economic costs, and the potential for hospital-acquired outbreaks. [24] The gravity of this issue is underscored by projections indicating that by 2050, AR is likely to account for 10 million deaths and inflict substantial financial burdens, with a disproportionate impact on low- and middle-income countries.[25]

The bacteria that most frequently cause infections in hospitalized patients include *S. aureus*, *Streptococcus* sp., *Acinetobacter* sp., *S. coagulase negative*, *P. aeruginosa*, *E. coli*, *Proteus mirabilis*, and *K. pneumoniae*. Some of these bacteria represent a global threat due to their increasing resistance to available antibiotics and their role in transmitting such resistance between bacteria of the same and different species. [26,27]

Over the past decade, there has been a notable increase in research related to the pathogens carried by cockroaches collected in various environments, including hospitals [6]. However; there has been a paucity of studies that have synthesized this information with the aim of facilitating access to it and enhancing the understanding and comprehension of personnel working in such health institutions. Accordingly, the objective of the present study was to describe the pathogens carried by cockroaches collected in hospitals in different parts of the world during the period 2000-2023 and their resistance mechanisms by means of a systematic review of the indexed literature.

2. Materials and Methods

2.1. Design

A systematic review of the literature was conducted to identify the pathogens and resistance mechanisms carried by cockroaches collected from hospitals in various parts of the world.

2.2. Methodology used in the search for relevant studies

The search strategy was conducted in accordance with the guidelines set forth in the PRISMA 2020 statement and through manual examination [28].

- The following keywords were established: cockroach, bacteria, hospital, pathogenic parasites, and antibiotic resistance. In order to optimize the retrieval of relevant information, the bibliographic citations of the articles examined were selected.
- The literature search was conducted using Google Scholar and electronic databases pertinent to the field of health, namely LILACS, the SciELO portal, and PubMed.
- 3. The articles were identified based on their title, language, date of publication, and the time interval between January 2000 and April 2023 (the extreme years of the interval were included).
- Reports that met at least one of the exclusion criteria proposed by the authors of this study were excluded.
- 5. The selected articles were found to meet all the inclusion criteria established by the authors of this work.

2.3. Inclusion criteria for studies.

- Reports of isolation and identification of pathogenic organisms of medical importance from external or internal parts of cockroaches.
- Articles published in the period from January 1, 2000 to April 30, 2023. For the establishment of
 this period of time, we took into account what was described by Guzman et al.[6] who express
 that since 2000 there has been an increase in publications in Pubmed related to the topic of
 cockroaches and the pathogens identified from the body of these insects.
- Studies carried out with cockroaches collected in hospital environments, although the research
 also describes the capture of these insects in other places such as homes, hotels, markets, schools,
 or restaurants.
- Scientific reports from anywhere in the world that are published in the form of a thesis, scientific article or letter to the editor in English, Spanish or Portuguese.

2.4. Exclusion criteria for studies.

- Articles in which the scientific name of the cockroach species collected is absent.
- Exclusively experimental research studies conducted in a laboratory setting on pathogens isolated from cockroaches collected in hospital settings.
- Publications that constitute review articles.

2.4. Statistical Analysis

Microsoft Office Excel 2007 version was used to present the results in tables and figures.

3. Results

3.1. Search for information

Figure 1 shows the flowchart of the selection process of the reports that make up the review based on the updated PRISMA guide [28]. Initially, using the established keywords, their combinations, date of publication and languages, 109 records on the subject were identified. Thirty-four articles were excluded because they were considered duplicates. In addition, 22 were eliminated because they were dissertations, conferences and book data.

The 53 reports resulting from the screening of the records were examined based on the inclusion and exclusion criteria previously established by the authors. Finally, 22 articles were selected for review. Most of the articles were published in Pubmed 65/109 (59.6%).

Figure 1. Flowchart of the report selection process based on the PRISMA guide.

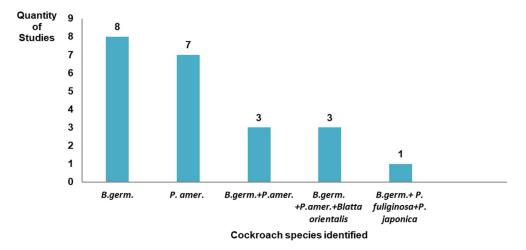
3.2 Distribution of articles by continent and country.

(n=22)

Table 1 shows that 95.4% of the research comprising the present review was conducted in developing nations [29]. Of these, 45.4% were published in countries on the Asian continent, while 39% and 23% were in African and American nations, respectively. Iran, Brazil and Ethiopia accounted for 59% of the studies involved in this review.

3.3. Cockroach species collected and identified in the studies analyzed.

Figure 2 shows that five cockroach species were identified in the 22 articles included in the review. [17,30-50] B. germanica and P. americana were the most common, reported in 15 (68%) and 13 (59%) of the studies described, respectively. [17,30-50] In seven studies (32%) more than one cockroach species was captured in the same hospital environment. [17,36,37,41,46,48,50]



B. germ: Blattella germanica; P: Periplaneta; P. amer: Periplaneta americana

3.4. Microorganisms isolated and identified from cockroaches collected in hospitals.

Table 1 shows the microorganisms isolated and identified from the bodies of cockroaches collected in hospitals. Bacteria represented the predominant group, being reported in 89% of the articles included in the review, followed by fungi and helminths with 13% and protozoa with 9%.

Bacteria of potential medical importance appearing in the greatest number of articles were: *E. coli* (11 articles), *S. aureus* (9 articles), *Enterobacter aerogenes* and *K. pneumoniae* (7 articles), *P. aeruginosa* and *Citrobacter freundii* (6 articles) and *Enterobacter cloacae*, *Klebsiella oxytoca* and *Proteus vulgaris* (5 articles). *Enterococcus faecalis*, *Morganella morganii*, *Proteus mirabilis*, *Staphylococcus epidermidis*, *Streptococcus pneumoniae*, *Streptococcus pyogenes* and *Streptococcus agalactiae* appear less frequently (Table 1).

Regarding fungi, two articles reported the identification of the species *Aspergillus niger* from the external part of cockroaches, while *Candida glabrata*, *Candida krusei* and *Aspergillus fumigans* were isolated in a different study from those analyzed in the review. The remaining members of this group were not classified at the species level but at the genus level, such as *Mucor* spp, *Penicillium* spp and *Rhizopus* spp (Table 1).

Helminths harmful to human health were identified: *Enterobius vermicularis* (eggs, larvae and adults) and *Ancylostoma duodenale* (larvae) in two and one of the studies analyzed, respectively. In addition, the presence of organisms belonging to the genera *Ascaris* (adults) and *Taenia* were detected (Table 1).

Regarding protozoa, the presence of *Lophomonas blattarum* and *Entamoeba coli* was identified in two and one of the studies, respectively. In other studies, organisms such as *Blastocystis* spp., *Cyclospora* spp., and *Cystoisospora* spp. were reported (Table 1).

Isolation and identification of medically important microorganisms from the external and internal parts of cockroaches was reported in 12 (55%) of the 22 articles reviewed. In all studies, pathogens were detected in both body parts of these vectors. [17,32,33,34,39,40,42,43-45,47,50]

One article (4.5%) reported the identification of viruses (rotavirus) carried by cockroaches on the external and internal parts of their bodies. [40]

3.5. Antibiotic resistance and mechanisms of resistance in bacteria identified in cockroaches.

Antibiotic resistance in bacteria isolated and identified from cockroaches collected in hospitals was addressed in 64% (14/22) of the studies included in this review (Table 1).

Figure 3 shows that gram-negative bacteria were identified in the highest percentage of the articles analyzed where AR is reported. In this regard, *E. coli* stands out with 8 studies (57%), followed by *K. pneumoniae* and *P. aeruginosa*, which are described in 4 (29%) and *E. cloacae* in 3 (21%). In the case of gram-positive bacteria, *S. aureus* appears in 5 studies (36%).

AR in bacteria was frequently observed in antibiotic groups such as penicillins, cephalosporins, aminoglycosides, amphenicols, tetracyclines and quinolones. [17,30-33,39,40,45,46]

Regarding antibiotic resistance mechanisms, the BlaZ gene coding for beta-lactamase enzymes and the production of extended-spectrum beta-lactamases were detected in one and two studies, respectively. [17,45,50] These enzymes are responsible for the resistance mechanism characterized by inactivation of the antibiotic. [51] On the other hand, the presence of genes coding for Penicillin-binding proteins involved in the resistance mechanism called target site modification [51], appears in one investigation. [48] Two articles reported the presence of tet(K) and tet(L) genes encoding proteins involved in the mechanism of antibiotic efflux through the energy-dependent pump. In addition, these studies were able to identify the tet(M) and tet(O) genes involved in the mechanism of target site alteration that protects the ribosome from the action of the antibiotic. [17,48,52] Finally, the mechanism of resistance (target site modification) to colistin mediated by the mcr-1 gene was detected in E. coli isolates. [50,53]

Table 1 shows that in 17 of the 18 studies (94%) bacteria were identified by seeding in culture media (general and differential), Gram staining (GS) and Conventional Biochemical Tests (CBT). In addition, five of the 17 articles referred above (29.4%) used other techniques such as Analytical Profile Index (API), Serological Tests and PCR. In one of the 18 investigations, only the culture media seeding method and MALDI-TOF technology were used.

For the identification of fungi, methods based on seeding in culture media and examination of macroscopic and microscopic characteristics of the growths were used. Protozoa and helminths were classified using direct parasitologic methods, and rotaviruses were identified using ELISA technology (Table 1).

3.7. Methods Used to Detect Antibiotic Resistance and its Mechanisms in Bacteria Identified in Cockroaches

As shown in Table 1, the predominant method used to detect antimicrobial resistance was the use of disk diffusion (Kirby Bauer). This method was used in 12 of the 14 studies in which AR was investigated. On the other hand, PCR was used in three of the 12 articles (25%) and the Etest method in one of the 14 studies (7.1%). It is worth noting that one of the studies reviewed did not report the method used to detect AR (Table 1).

The detection of genes and enzymes that indicate the antibiotic resistance mechanism present in the bacteria analyzed was performed in five of the 22 studies (22.7%). [17,37,45,48,50] The PCR technique was used in three of them, [17,37,48] in one the Double Disc Synergy Test (DDST) [45] and in another article RT-PCR and DDST. [50]

Table 1. Microorganisms isolated and identified from cockroaches collected from hospitals in different parts of the world, 2000 - 2023.

different parts of the world, 2000 - 2023.			
Author, year, reference number	Country	Microorganisms isolated and identified from cockroaches	Methods used in the identification of microorganisms, determination of antimicrobial susceptibility profile and antibiotic resistance mechanisms.
Prado <i>et al.</i> , 2002 (30)	Brazil	Bacteria: E. coli, K. pneumoniae, E. cloacae, S. marcescens, H. alvei, E. gegorviae, Serratia spp., K. oxytoca, P. vulgaris, Morganella morganii	IM (Bacteria): Seeding in culture media (general and differential), Gram stain (TG) and conventional biochemical tests (CBT). PSA (Bacteria): Disc Diffusion Method (Kirby Bauer).
Marinésia <i>et al.</i> , 2006 (31)	Brazil	Bacteria: S. coagulase-negative, E. aerogenes, S. marcescens, H. alvei, E. cloacae, E. gergoviae, Serratia spp. Fungi: Yeast and Filamentous fungi	IM (Bacteria): Seeding in culture media (general and differential) and PBC.
Elgderi <i>et al.,</i> 2006 (32)	Libya	Bacteria: E. coli, K. pneumoniae, K. oxytoca, K. ornithinolytica, E. cloacae, E. aerogenes, Pantoea sp., C. freundii, C. braakii, C. youngae, C. amalonaticu, S. marcescens, S. liquefaciens, P. mirabilis, P. vulgaris, M. morganii, H. alvei, Buttiauxlla agrestis, Aeromonas hydrophila, Aeromonas	

		caviae, P. aeruginosa, Acinetobacter	
Tachbele <i>et al.</i> , 2006 (33)	Ethiopia	sp., Streptococcus.sp. Bacteria: Shigella flexneri, E. coli O15717, S. aureus, Bacillus cereus.	IM (Bacteria): Seeding in culture media (general and differential), TG, PBC and Serological Tests. PSA (Bacteria): Disc Diffusion Method (Kirby Bauer).
Salehzadeh <i>et al.,</i> 2007 (34)	Iran	Bacterias: E. coli, Haemophilus spp., S. hemolítico group A y B. Fungi: Candida spp., Mucor spp., Aspergillus niger, Rhizopus spp., Penicillium spp., Aspergillus fumigans Helminths: Enterobius vermicularis, Ascaris spp.	IM (Bacteria): Seeding in culture media (general and differential), TG y PBC. PSA (Bacteria): Seeding in culture media (general and differential) (Kirby Bauer). IM (Fungi): Seeding in Sabouraud agar medium and observation of macro and microscopic morphological characteristics of the sample. IM (Helminths): Direct parasitologic methods (saline and Lugol) and observation of the sample under the light microscope.
Aparecida <i>et al.,</i> 2008 (35)	Brazil	Bacteria: Salmonella spp., E. coli, C. freundii, H. alvei, S. aureus, E. aerogenes, Serratia spp.	IM (Bacteria): Seeding in culture media (general and differential), TG y PBC. PSA (Bacteria): the method used is not shown.
Karimi <i>et al.,</i> 2009 (36)	Iran	Bacteria: E. coli, K. pneumoniae, K. oxytoca, E. cloaceae, E. aerogenes, P. mirabilis, P. vulgaris, M. morganii, C. freundii. C. diversus, Edwardsiellae trada, S. aureus, S. epidermidis, S. saprophyticus, S. group D, P. eruginosa	IM (Bacteria): Seeding in culture media (general and differential), TG y PBC.
Saitou <i>et al.</i> , 2009 (37)	Japan	Bacteria: P. aeruginosa	IM (Bacteria): Seeding in culture media (general and differential), PBC y API. PSA (Bacteria): Using the Etest. MRA: PCR.
Risco et al., 2010 (38)	Cuba	Bacteria: Acinetobacter calcoaceticus, Alcaligenes faecalis, C. diversus, C. freundii, E. aerogenes, E. agglomerans, E. cloacae, Enterococcus sp., E. coli, K. oxytoca, K. pneumoniae, P. mirabilis, P. vulgaris, P. stuartii, P. aeruginosa, S. marcescens, S. aureus, S. epidermidis. Fungi: Aspergillus spp., Mucor spp., Rizopus spp.	IM (Bacteria): Seeding in culture media (general and differential), TG y PBC. IM (Fungi): Seeding in Sabouraud agar
Tilahun <i>et al.,</i> 2012 (39)	Ethiopia	Bacteria: K. oxytoca, K. pneumoniae, E. cloacae, C. diversus, P. aeruginosa, Providencia rettgeri, K. ozaenae, E. aerogenes, S. aureus, E. coli, Shigella flexneri, Enterococcus faecalis.	IM (Bacteria): Seeding in culture media (general and differential), TG y PBC. PSA (Bacteria): Disc Diffusion Method (Kirby Bauer).
Tetteh-Quarcoo et al., 2013 (40)	Ghana	Bact eria: K. pneumoniae, E. coli, P. vulgaris, C. ferundii, E. cloacae,	IM (Bacteria): Seeding in culture media (general and differential), TG y PBC.

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		P. aeruginosa, Enterococcus faecalis, K. oxytoca. Helminths: Ancylostoma duodenale, Taenia spp. Virus: Rotavirus	lugol and observation under a light
Motevali <i>et al.,</i> 2014 (41)	Iran	Fungi: Candida spp., Rhodotorula spp., Aspergillus spp., Fusarium spp., Penicillium spp. Geotrichum spp., Alternaria spp., Cladosporium spp., Trichoderma spp., Mucor spp., Chrysosporium spp.	IM (Fungi): Seeding in Sabouraud's dextrose agar with chloramphenicol, tube germination test (yeast) and observation of macro and microscopic morphological characteristics of the sample.
Menasria <i>et al.,</i> 2014 (42)	Argelia	Bacteria: S. aureus, C. erundii, E. cloacae, E. aerogenes, K. pneumoniae, S. marcescens.	IM (Bacteria): PBC y API.
Suresh <i>et al.,</i> 2015 (43)	India	Bacteria: Salmonella B, Salmonella D, Salmonella E, Shigella B, E. coli, S. aureus.	IM (Bacteria): Seeding in culture media (general and differential), TG, PBC y Pruebas Serológicas. PSA (Bacteria): Disc Diffusion Method (Kirby Bauer).
Cazorla <i>et al.,</i> 2015 (44)	Venezuela	Protozoans: Entamoeba blattae, Nycthoterus ovalis, Leptomonas spp., Cyclospora spp., Entamoeba coli, Cystoisospora spp., Lophomonas blattarum, Lophomonas striata Helminths: Enterobius vermicularis, Thelastoma spp., Hammerschmidtiella spp.	,
Ikechukwu et al., 2017 (45)	Nigeria	Bacteria: E. coli, Salmonella spp., Shigella spp.	IM (Bacteria): Seeding in culture media (general and differential), TG y PBC. PSA (Bacteria): Disc Diffusion Method (Kirby Bauer). MRA (Bacteria): DDST
Abdolmalek et al., 2017 (17)	Iran	Bacteria: S. aureus	IM (Bacterias): Seeding in culture media (general and differential), TG y PBC. PSA (Bacterias): Disc Diffusion Method (Kirby Bauer) and PCR. MRA (Bacterias): PCR
Nazari <i>et al.,</i> 2020 (46)	Iran	Bacteria: E. coli, S. coagulasa- negativa, Proteus spp., Enterococcus spp., Micrococcus spp., Pseudomona spp., Serratia spp.,Streptococcus.B, Streptococcus A, S. aureus.	(general and differential) 1(v PB(
Khodabandeh <i>et</i> <i>al.</i> , 2020 (47)	Iran	Fungi: Aspergillus niger, Rhizopus spp., Penicillium spp., Mucor spp., Candida glabrata, Candida krusei	IM (Protozoans): Seeding in different culture media; observation of macro and microscopic morphological characteristics of the sample.
Chehelgerdi <i>et</i> al., 2021 (48)	Iran	Bacteria: S. pneumoniae, S. pyogenes, S. agalactiae	IM (Bacterias): Seeding in culture media (general and differential), TG, PBC y PCR. PSA (Bacteria): Disc Diffusion Method (Kirby Bauer) and PCR. MRA (Bacteria): PCR

Farzad <i>et al.</i> , 2021 (49)	Iran	Protozoans: Gregarina sp.,	IM (Protozoans): Direct parasitologic
		Lophomonas blattarum. Entamoeba	methods: samples paired in saline, Lugol
		sp. Blastocystis sp. Nyctotherus sp.	stained, then light microscoped.
Landolsi <i>et al.,</i>		Postorio V manuscrico F coli F	IM (Bacteria): Seeding in different culture
			media and mass spectrometry (MALDI-TOF).
2022	Tunisia	Bacteria: K. pneumoniae, E. coli, E. cloacae, C. sedlaki	PSA (Bacteria): Disc Diffusion Method (Kirby
(50)		стоисие, С. ѕешикі	Bauer) and PCR.
			MRA (Bacteria): DDST and Real Time PCR

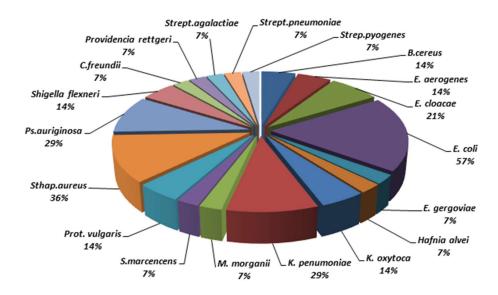


Figure 3. Distribution of bacterial species with antibiotic resistance in the articles analyzed.

4. Discussion

A systematic review of pathogens of medical importance isolated from cockroaches collected in hospitals in different parts of the world over a 24-year period was performed. A literature search strategy similar to that used by other authors in reviews of isolated fly, cockroach, and beetle pathogens was established for the selection of studies. [54-56] This allowed a thorough process of identification, screening, and selection of articles based on the inclusion and exclusion criteria proposed by these authors.

Ninety-five percent of the studies in the review come from developing countries [29] in the Americas, Africa, and Asia. [17,30-36,38-50]. Similar findings have been reported by different authors in their respective review articles. [55-57] In the countries where these studies were conducted, there are socioeconomic factors that negatively affect the health conditions found in second and third level hospital facilities. [17,30,38, 39,40,58] The presence of cockroaches in hospitals, parasites in their integument and gastrointestinal tract is an indicator of poor sanitation in health facilities. [1,44] On the other hand, there are reports of cockroaches collected in hospitals in Japan, France, and Poland [37,59,60] showing that this phenomenon is independent of the geographical location and economic situation of nations, which becomes an epidemiological alert for developed countries.

B. germanica and *P. americana* were the predominant cockroach species in the studies included in this review. [17, 30-50] Both are recognized as the most abundant of this group of insects worldwide. [7-9] Trade and the ability of both species to adapt to a wide range of conditions have played a key role in their distribution. [1, 2,3-5,11,61] *Blatta orientalis, Periplaneta fuliginosa,* and *Periplaneta japonica* are other species considered to be of medical importance with limited worldwide distribution reported in the articles reviewed. [1,2,36,37,48,50]

In seven of the 22 articles analyzed (32%), several species of cockroaches coexisting in the same hospital facility and carrying pathogens of medical importance were captured. [17,36,37,41,46,48,50] This fact could increase the likelihood of the spread and mechanical transmission of these organisms in the hospital environment. The high infestation of *P. americana* in the drainage system of hospitals, as well as its size, 3 to 4 times larger than *B. germanica*, could facilitate this phenomenon. [17] On the other hand, Karimi *et al.* [36] collected 305 cockroaches in three hospitals in Iran belonging to the species *P. americana* (65.6%), *B. germanica* (12.1%) and *Blatta. orientalis* (22.3%). The first and second mentioned species were caught in corridors, laundry rooms, basements, food stores, and facilities. However, the third species predominates in places where the temperature is colder in relation to the collection sites of the other two species. The difference in habitat is another aspect that favors the spread of pathogens by cockroaches.

Cockroaches host and carry a variety of pathogens on the external and internal parts of their bodies, such as bacteria, infective forms of helminths, fungi, protozoa, and rotavirus, as reported by several authors. [55,57] The nocturnal habits of these vectors, their reproductive capacity, the ease with which they enter synanthropic places (homes, restaurants, hospitals, etc.), the ingestion of a wide variety of foods, including waste, and their free movement over a wide range of surfaces such as floors, tables, pipes, ceilings, cracks, and corridors condition the contact with ubiquitous microorganisms. [1,11,23,58] In addition, many pathogens can persist for months on dry inanimate surfaces, depending on the optimal conditions of temperature, humidity and amount of inoculum, which facilitates the contact of cockroaches with them and consequently favors their mechanical vector capacity. [1,3,11,37,44,61,62] In this sense, different authors in their research do not find statistically significant differences between the distribution of potentially pathogenic bacteria identified on the outside and inside of the body of cockroaches captured in hospitals. [17,33,39,40,43]

In fact, bacteria were the most frequently identified group in the studies included in the review. Several authors acknowledge the predominance of these microorganisms among the pathogens isolated from cockroaches. [1,5,11] This finding is not surprising given the high prevalence of these infectious agents in healthcare-associated infections. [19,20] However, Elgderi *et al.* [32] captured specimens of *P. americana* from hospitals and homes in Tripoli, Libya, show no evidence that insects from hospitals are more likely to carry bacteria than those from homes. This suggests that the presence of these pathogens in these insects is primarily related to the sanitary conditions of the environments they inhabit.

Bacteria commonly identified in the review articles were: *E. coli, S. aureus, E. aerogenes, K. pneumoniae, P. aeruginosa, C. freundii, E. cloacae, K. oxytoca,* and *P. vulgaris,* all of which are known to cause human infections, particularly in the hospital setting. [63-67] *K. pneumoniae* and *S. aureus* cause nosocomial pneumonia; *P. aeruginosa* causes surgical wound and burn infections; *E. coli* causes urinary tract infections; and both *E. aerogenes* and *E. cloacae* are opportunistic microorganisms that commonly cause infections in patients admitted to health care facilities. [54,63] These pathogens have been isolated from cockroaches collected from various parts of the hospital, such as the operating room, intensive care unit, and neonatal unit. [30,31,39]

Several review articles suggest that cockroaches may play an important role in hospital-acquired parasitosis. Carzola *et al.* [44] identified enteroparasites of medical importance in *P. americana* cockroaches captured in Venezuela. Among the parasites identified were *E. vermicularis* and trophozoites of the protozoan *Lophomona blattarum* isolated from patients with pulmonary disease in Iran, China, Spain, India, and Peru. [49, 68]

The studies analyzed in this research show the presence of fungi of medical importance carried by cockroaches collected in hospitals. Mycoses in hospital environments are common over the world, mainly caused by *Aspergillus* spp. and *Candida* spp. [57] In the same line, Motevali *et al.* [41] demonstrated the presence of different species of fungi *A. flavus*, *A. niger*, *A. fumigatus*, *C. albicans* and *C. glabrata* in *B. germanica* and *P. americana* collected from three hospitals in Iran.

Viruses (rotaviruses) were identified in only one study in this review. This finding is the first report of identification of these infectious agents in cockroaches collected from hospitals in Ghana. [40] Rotavirus causes severe and fatal diarrhea in young patients worldwide and accounts for half of

all hospitalizations for this condition in children under 5 years of age in developed countries. It is also responsible for approximately 25% of all hospital-acquired viral infections, particularly in immunocompromised children. [57] However, the scarce detection of viruses in the articles included in this review may be due to the fact that the identification of pathogens in cockroaches tends to focus on bacteria, helminths, fungi and protozoa due to the availability of resources and ease of diagnosis. [1,11,40,58]

AR in bacteria is responsible for 70,000 patient deaths annually. [25] This phenomenon is reported in 14 articles in the current study, and mainly in bacteria of the ESKAPE group (*Enterobacter* spp., *E. coli, K. pneumoniae, S. aureus* and *P. aeruginosa*). [69] Moreover, other bacteria that are potentially harmful to humans have also been described: *H. alvei, P. vulgaris, S. marcescens, S. pneumoniae, S. agalactiae*, and *S. pyogenes*. These factors combine to create a global health risk associated with AR.

Bacterial resistance in cockroach species to a wide range of antibiotics such as penicillin, tetracycline, gentamicin, ceftaroline, aztreonam, chloramphenicol, cefepime, ceftazidime, erythromycin, ampicillin, and amickacin is highlighted. [17,30-33,38,39,44,45] Prado et al. [30] isolated 15 species of enterobacteria from *P. americana* captured in a public hospital in Brazil. Antibiotic susceptibility testing revealed resistance to most of the antimicrobials tested in the study. Abdolmalekiet et al. [17] provided the first report of phenotypic and genotypic evaluation of antibiotic resistance in Methicillin-Resistant *Staphylococcus aureus* (MRSA) isolates from external and internal parts of *P. americana* and *B. germanica* collected in hospitals. The isolates also showed a high prevalence of resistance to the antibiotic penicillin, ceftaroline, tetracycline, gentamicin and trimethoprim-sulfamethoxazole. In the neonatal ward of a hospital in the capital of Ethiopia, 400 cockroaches of the species *B. germanica* were collected. Of these, *K. pneumoniae*, *K. oxytoca*, *Providencia rettgeri*, *C. diversus*, *Citrobacter spp.*, *E. cloacae*, *P. aeruginosa*, *E. coli*, *E. aeroginosa*, *S. aureus*, and *Acinetobacter* spp. were identified, and multidrug resistance was observed in all of them. [45] The above indicates that cockroaches may be involved in the spread of AR among bacteria. [70]

A bacterial strain may develop multiple mechanisms of resistance to one or more antibiotics, and an antibiotic may be inactivated by different mechanisms in different bacterial species. Therefore, knowledge of these mechanisms is necessary as a fundamental step for the effective treatment of diseases with antimicrobial drugs. [69] In this review, the detection of AR mechanisms in bacteria was found in five of the 18 studies dealing with antibiotic resistance [17,37,45,48,50].

In Nigeria, using phenotypic methods, they found a mechanism of inactivation of beta-lactam drugs by the expression of extended-spectrum beta-lactamases (ESBL). [45] Chehelgerdi *et al.* [48] described for the first time antibiotic resistance in Iran by phenotypic and genotypic pathways in *Streptococcus* spp. isolated from cockroaches collected from hospital environments. These authors identified genes involved in the mechanisms of resistance to penicillins (pbp), tetracyclines (*tet K, tet M, tet O,* and *tet L*), macrolides (erm and mef), and streptogramins A and B (rplV) using PCR technique. In Tunisia, the production of BLEE (blaCTX-M1) and the presence of *mcr-1* genes responsible for the colistin resistance mechanism in the *E. coli* ST648 lineage were described for the first time. [50]

In terms of the methods used to identify microorganisms and determine antibiotic susceptibility profiles in bacteria, conventional technologies such as seeding in culture media, Gram staining, conventional biochemical tests, and direct parasitological methods predominated. A possible explanation for this phenomenon is that these are the techniques used in developing countries, which were the basic scenarios in which the research was carried out. On the other hand, advanced technologies that provide better diagnostic quality and faster results (API [32,37,42], PCR [17,48,50] and MALDI-TOF [50]) were not available to most of the laboratories in the studies analyzed in this review. This fact affects the quality of the results, since in many cases the microorganism can only be identified at the genus level, which prevents us from knowing whether they are of medical importance. In addition, there are probably other pathogens that cannot be identified due to technological limitations.

On the other hand, in the detection of enzymes and genes that indicate the presence of antibiotic resistance mechanisms, it was observed that four articles included in this review employ the PCR technique, three of which correspond to developing countries [17,48,50] carried out in the period 2017-2021. However, in Japan, this technology was used for the same purpose in 2009. [37] It is undeniable that these studies allow the identification of new pharmacological targets and the design of specific antibiotics to provide more precise treatments to combat infections caused by bacteria. [71]

The current research has the following limitation: in many studies, the nomenclature of the identified microorganisms was described down to the genus, which did not allow us to know the species and, consequently, its medical importance. For this reason, the analysis of the results emphasizes pathogens identified down to the species level.

5. Conclusions

Cockroaches in hospitals all over the world might carry potentially pathogenic organisms, which are recognized as etiological agents of human diseases belonging to the groups of bacteria, fungi, helminths, and protozoa. This reinforced the notion that the presence of these insects in healthcare facilities represents a potential risk due to their involvement in healthcare associated infections and as reservoirs of multidrug-resistant bacteria. Given the scarcity of publications on the subject addressed in this review, it is necessary to increase and deepen the study of pathogens of medical importance in cockroaches, a topic of importance for global health.

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