**Erysipelothrix rhusiopathiae**: bacteriology, epidemiology and clinical manifestations of an occupational pathogen

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**Introduction**

The genus *Erysipelothrix* consists of two named species, *E. rhusiopathiae* and *E. tonsillarum*, and an as yet unnamed third species. All are gram-positive, non-sporing rods. *E. rhusiopathiae* is a pathogen or a commensal or saprophyte of a wide variety of wild and domestic animals, birds and fish. Diseases of economic importance in animals include swine erysipelas, erysipelas of farmed turkeys, chickens and emus, and polyarthritis in sheep and lambs. Man can be infected; the organism is an occupational pathogen, prevalent in those working in association with animals and animal products. Three forms of human disease are recognised, the mildest and most common of these is the skin infection known as erysipeloid.

**Bacteriology**

**History and nomenclature**

Bergey's Manual [2] classifies *E. rhusiopathiae* as a regular non-sporing gram-positive rod. Until recently,
the genus consisted of only the type species, *E. rhusiopathiae*, which was known to demonstrate considerable serological, biochemical and antigenic variation [3]. Genetic analyses have revealed a new species, *E. tonsillarum* [4]. *Erysipelothrix* was originally considered a close relative of the genus *Listeria*, although numerous molecular taxonomic studies have concluded now that the genus is a distinct cluster of organisms, most similar to the streptococci [2, 3].

*E. rhusiopathiae* (from the Greek 'erysipelas' – a disease, 'thrix' – a hair or thread, 'rhusius' – reddish and 'pathus' – disease [5]), literally 'erysipelas thread of red disease' [1], has a long history and is the result of many name changes. Koch [6] first isolated a strain of the genus *Erysipelothrix* in 1876 from a mouse he inoculated with putrefying blood. He described the organism as the 'bacillus of mouse septicaemia' and named it *E. muriseptica*. In 1882, Loeffler isolated a similar organism from cutaneous blood vessels of a pig which had died from swine erysipelas, and was the first to describe fully the infectious agent and the disease it caused in swine [6].

The first description of human disease, later attributed to *Erysipelothrix*, was reported in 1870 in the *British Medical Journal*; further cases were documented in 1873 as erythema serpens [7]. However, it was not until 1884, when Rosenbach isolated an organism similar to Koch's from a patient with localised cutaneous lesions, that *Erysipelothrix* was established as a human pathogen. He coined the term 'erysipeloid' to differentiate between the human streptococcal disease erysipelas and the condition he had observed [6].

Rosenbach [6] distinguished three separate species of the organism, *E. muriseptica*, *E. porci* and *E. erysiploides*, based on their isolation from mouse, pig and man, respectively. It was later realised that these were three nearly identical strains of the same species [3], and they were named *E. insidiosa*, as originally proposed by Trevisan in 1885. This name, and all 36 other documented names for the organism, were rejected in favour of *E. rhusiopathiae* in 1966, a combination which originated in 1918 [8].

Takahashi *et al.* [9] isolated a cluster of avirulent serotype 7 strains which were later found to be genetically distinct from *E. rhusiopathiae* by DNA base composition and DNA–DNA homology studies [4]. These strains formed the basis of a new species, *E. tonsillarum* (from the Latin 'of the tonsils'). Originally, *E. tonsillarum* was considered morphologically and biochemically identical to *E. rhusiopathiae*, but it was shown later that *E. tonsillarum* could ferment sucrose, while *E. rhusiopathiae* could not [10, 11]. *E. tonsillarum* was described as avirulent for pigs, mice and chickens, but pathogenic for dogs [12, 13]. There have been no studies so far into the pathogenicity of this species for man.

**Morphology and growth characteristics**

*E. rhusiopathiae* is a non-motile, non-sporulating, non-acid-fast, slender gram-positive rod, which is easily decolourised [2]. Gram-negative forms are often seen [14], particularly if the culture is old [15]; thus, the organism has been cited occasionally as a gram-negative bacillus [16]. *E. rhusiopathiae* was long described as non-capsulate [2], until recent studies showed the presence of a capsule and suggested a role for it in virulence [17].

Based on the colonial appearance of the organism, *Erysipelothrix* morphology is described as smooth (S) or rough (R) [2, 8, 18]. S-form colonies are convex, with a smooth surface and entire edge [2, 15, 19]. R-form colonies are slightly larger with an irregular edge and a flattened, rough surface [14]. All colonies are clear, circular and very small (0.1–0.5 mm diameter after 24 h; 0.5–1.5 mm after 48 h [2, 19]), increasing in size and tending towards a pale blue opacity with further incubation or age [2, 19]. Most strains exhibit a narrow zone of α-haemolysis on blood agar, which can even show slight clearing after 48 h [2]. R-form colonies do not cause haemolysis [18].

Growth in broth was best described by Smith (cited by Jones [2]), who noted that the suspension had a 'faint opalescence … which on shaking was resolved for a moment into delicate rolling clouds'. S forms cause slight turbidity and a powdery deposit, whereas R forms have a tangled hair-like appearance [3].

Cell morphology is closely linked to the colonial characteristics of each form [19]. S forms are slender, straight or slightly curved rods with rounded ends, 0.8–2.5 μm in length and 0.2–0.4 μm in diameter. The rods exist in various formations, often as small chains [2]. The R form exhibits a predominantly filamentous morphology, frequently likened to the mycelial formations of fungi, although branching does not occur. The filaments can be 4 μm to >60 μm in length and can have a beaded appearance with Gram's staining [2]. Long chains of distinct rods can also exist in this form [14, 18].

The origin of R and S forms and their role in disease have received much emphasis in the *Erysipelothrix* literature, possibly because different forms for other pathogens, such as *Streptococcus pneumoniae*, have distinct roles in virulence. For *E. rhusiopathiae* these roles are not definite, and there are conflicting observations on the role of each form. Furthermore, the distinction between S and R forms is not always clear. An intermediate (RS) form, which is the most common conformation [15], has been used to describe colonies sharing the characteristics of both types [8]. Some investigators have suggested that S forms are commonly isolated from acute pig diseases, such as septicaemia, and R forms from more chronic syn-
dromes such as arthritis and endocarditis [8]. However, there are conflicting reports relating to the virulence of R forms. Gorby [20] reported that, in pig disease, R forms are the more virulent type, while Taylor [21] stated that the R form was generally considered less virulent. Similar possible relationships for S and R forms in human disease have not been documented.

Media and conditions of incubation (see below) play a major role in morphology formation; R forms are favoured by incubation at 37°C in acidic pH, while S forms predominate in alkaline conditions (pH 7.6-8.2) with incubation at 33°C [14]. Changing growth conditions has allowed S forms to give rise to R and RS forms, and S forms to originate from R forms [3]. The S form often dissociates to the R form with age [8, 15]. These changes in morphology and cultural characteristics are reported to lead to changes in virulence and antigenic properties [8].

**Growth conditions and requirements**

_E. rhusiopathiae_ is a facultative anaerobe [3]. Newly isolated strains are micro-aerophilic, but laboratory-adapted cultures grow both aerobically and anaerobically, with some strains being favoured by incubation in CO₂ 5% or 10% [2]. The organism can grow at temperatures between 5° and 44°C [2], optimally between 30° and 37°C [18]. Best growth is favoured by an alkaline pH. The optimum pH range has been documented as 7.2-7.6 [2, 5] or 7.4-7.8 [15, 18, 19, 221 and the limits of growth as 6.7-9.2 [23].

Growth is enhanced by the inclusion of serum 5-10%, blood, glucose 0.1-0.5%, protein hydrolysates, or surfactants such as Tween 80 in media [8, 18]. The exact nutritional requirements of the organism are not known [18], but riboflavin, small amounts of oleic acid and several amino acids [24] – particularly tryptophan and arginine [8, 25] – are needed for growth. Higher concentrations of glucose and oleic acid are inhibitory [5].

**Biochemistry**

The genus _Erysipelothrix_ is relatively inactive and gives negative results for catalase, oxidase, methyl red, indole and Voges-Proskauer reactions [15]. Carbohydrate fermentations produce acid without gas, but reaction patterns are variable and depend on the basal medium and indicator used [14, 26]. Andrade's agar with horse serum 10% is the recommended medium for biochemical tests. The majority of strains produce H₂S gas, but again the extent of this production varies with the culture medium. The best reaction is demonstrated on triple sugar iron agar [27]. A more detailed description of the biochemical characteristics of _Erysipelothrix_ can be found in the reviews of Ewald [8], Jones [2] or Reboli and Farrar [3]. Traditionally, biochemical reactions were used to differentiate between _Erysipelothrix_ and morphologically similar bacteria, such as _Listeria_ and _Corynebacterium_ spp. Characteristics used for this purpose included α-haemolysis, lack of motility, lack of catalase production and resistance to neomycin [14]. While traditional biochemical testing may still be of value, particularly in discriminating between _E. rhusiopathiae_ and _E. tonsillarum_, rapid identification can be achieved with an API Coryne System strip (bioMérieux) [28].

**Chemical tolerances**

_E. rhusiopathiae_ is a remarkably resistant organism for one that does not form spores [18]. The survival of the organism in the environment is an important factor in the epidemiology of disease. _E. rhusiopathiae_ is also tolerant to numerous chemicals. It can grow in the presence of phenol 0.2% and crystal violet 0.001%, and is said to be one of the organisms most resistant to sodium azide, tolerating 0.1% [23, 29]. Some of these chemical tolerances have been utilised in the development of selective media.

**Antigenic structure**

Watts [30] noted that most strains of _Erysipelothrix_ had two kinds of antigen, a species-specific heat-labile protein antigen and a heat- and acid-stable polysaccharide antigen, which now form the basis for serotyping strains. Dedié (cited by Reboli and Farrar [3]) recognised two major serotypes, A and B. Strains that did not react with these specific antisera were named group N. The Arabic numeral serotyping system of Kucsera [31] superseded the alphabetical scheme, due to variations in previous methods of antigenic extraction. The current standard serotyping method is a double agar-gel precipitation test with type-specific rabbit antisera and antigen prepared by hot aqueous extraction [31, 32].

At present, strains of _Erysipelothrix_ are classified as serotypes 1–26 [33]. A group N still exists for strains that have no type-specific antigen. In swine, 75–80% of isolates are of serotype 1 or 2 (previously group A or B) and the less common serotypes make up the remaining 20% [18]. Some investigators have noted a relationship between serotype and clinical condition in pigs, with serotype 1a most commonly isolated from acute swine illness, and serotype 2a more prevalent in chronic forms of disease [34]. However, other surveys have provided contradictory results, and all clinical conditions can be induced experimentally in susceptible swine with a variety of serotypes [18, 35]. Sneath _et al._ [23] reported that strains from human or pig origin were antigenically similar, but there has been no verification of this since the numerical typing system was established. There is a deficiency in the literature regarding serotypes in human infection, and the epidemiological significance of serotyping in human disease is questionable.
Molecular studies have cast further doubt on the value of serotyping as a reliable taxonomic and epidemiological method of classification. DNA–DNA hybridisation [10], polyacrylamide gel electrophoresis [36] and multilocus enzyme electrophoresis studies [11] classified serotypes 1a, 1b, 2, 4–9, 11, 12, 15–17, 19, 21 and 25 in the species E. rhusiopathiae, and serotypes 3, 7, 10, 14, 20, 22–24 in E. tonsillarum, with differing results for serotypes 13 and 18 [11]. When Erysipelothrix strains were analysed by restriction fragment length polymorphisms (RFLP), both E. tonsillarum and E. rhusiopathiae contained serotype 2 (virulent) and 7 (avirulent) strains. The creation of a third species of Erysipelothrix was suggested to account for a distinct cluster of strains [37].

These findings suggest that there is no direct relationship between serotype and virulence, supporting earlier investigators who noted that, within each serotype, strains of high, low and no virulence existed, and that factors other than serotype were important in the induction of disease in animal models [38, 39]. Further investigation is required to reach a consensus on typing schemes and their role, if any, in studies of pathogenicity.

**Mechanisms of pathogenicity**

Relatively little is known about the pathogenesis of E. rhusiopathiae infections. Strains of E. rhusiopathiae are known to vary considerably in virulence. Despite much investigation, there has been no conclusive evidence of a relationship between virulence and morphology, chemical structure or antigenic structure [18]. Various virulence factors have been suggested, although their relative importance is not yet clear. The presence of a hyaluronidase and a neuraminidase has been recognised [40, 41], but hyaluronidase was detected in strains both virulent and avirulent for pigs. A correlation between the amount of neuraminidase produced and the virulence of strains was noted [41], although later studies demonstrated that avirulent acapsular mutant strains also produced the enzyme [17].

Adhesion to porcine kidney cells in vitro was greater for virulent strains [42]; however, the role of this factor in disease has not been investigated further. Further work on adhesion has been carried out [43], but this was in relation to arthritis in swine as a model for rheumatoid arthritis in man. Recently, the presence of a labile capsule was reported and acapsular mutants were constructed by transposon mutagenesis. In contrast to the parental strain, the mutants failed to resist phagocytosis by murine polymorphonuclear leucocytes and could not survive within murine macrophages, suggesting that the capsule was an important virulence factor [17, 44]. The mutant has been used in the development of a diagnostic PCR [45] (see below) and vaccine studies [46]. Further investigation of the pathogenesis of E. rhusiopathiae infection is required.

**Epidemiology**

E. rhusiopathiae and infections caused by this organism occur world-wide [6, 14]. Infections of man and animals have been documented from Africa, Australia, several countries in the Americas, Japan, China and throughout Europe [8]. Human disease can originate from an animal or environmental source.

**Animal disease**

Swine erysipelas caused by E. rhusiopathiae is the disease of greatest prevalence and economic importance [18]. There are three clinical forms: a severe acute septicaemic form of sudden mortality; a milder, subacute urticarial form characterised by purple diamond-shaped lesions on the skin and a chronic form with endocarditis or arthritis [14]. Swine erysipelas is economically detrimental to the pig industries of North America, Europe, Asia and Australia [18].

As well as affecting swine, E. rhusiopathiae causes infections in a wide variety of domestic and wild mammals (including marine mammals), domestic, game and wild birds, and man [3, 8, 19, 22]. Polyarthritis of sheep and lambs, and erysipelas in calves, ducks and domestic turkeys are also economically significant diseases caused by E. rhusiopathiae [6, 19]. In Australia, the organism is an emerging problem in farmed emus [47].

Domestic swine are believed to be the most important animal reservoir of E. rhusiopathiae. The organism is shed by diseased animals in faeces, urine, saliva and nasal secretions, which can contaminate food, water, soil and bedding, leading to indirect transmission of the organism [18]. Furthermore, an average of 20–40% of healthy swine, and in some herds up to 98%, harbour Erysipelothrix in the lymphoid tissue of the alimentary tract, particularly in the tonsils [18, 48, 49]. One study demonstrated that both virulent (serotypes 2, 6, 11, 12 and 16) and avirulent (serotype 7) serotypes were found on the tonsils [9]. Another showed that the faeces of apparently healthy animals contained virulent organisms [50]. The maintenance of E. rhusiopathiae in nature appears to result from asymptomatic carriage in animals and subsequent dissemination of the organism to the environment [6].

Mice are susceptible to infection [6], but other rodents seem to be affected only occasionally [51]. These animals can harbour the organism and are important reservoirs in some environments, such as meat packing plants [6]. Insects have been reported to carry E. rhusiopathiae, and are occasional vectors [6, 19, 52];
however, this is not a known route of infection for man.

Environmental reservoirs

The environment appears to be secondary in importance to animal reservoirs as a source of *E. rhusiopathiae*. However, in some circumstances, such as in marine environments, the organism may survive long enough to create a significant hazard to man [6].

*E. rhusiopathiae* is a saprophyte associated with some groups of animals, particularly marine fish, molluscs and crustaceans [6, 53, 54]. Freshwater fish and some species of bird are also hosts [14]. The organism survives and grows on the exterior mucoid slime of fish, without causing disease in the fish themselves [6]. It is likely that *E. rhusiopathiae* survives by a similar mechanism on the exterior slime layer of other marine creatures. The slime on fish appears to be an important source of infection for man. In early reports, fish caught under ‘aseptic conditions’ did not harbour *Erysipelothrix* [55], so investigators concluded that the organism was transmitted via the slime from other fish [54, 55]. Boxes used for transport of fish seemed to play a vital role in the transmission of *E. rhusiopathiae*, and many human cases resulted from contact with these objects [5, 54, 55].

Once in the environment, *E. rhusiopathiae* can survive for long periods although it does not form spores [18]. The organism is ubiquitous, and can be found wherever nitrogenous matter decomposes, retaining virulence and viability for months in putrid material [56]. Survival in swine faeces for 1–5 months, depending on seasonal conditions [19], and in soil for up to 5 years from the time of the last diseased pig [57], has been demonstrated. However, this latter report did not consider the possibility of asymptomatic shedding. *E. rhusiopathiae* has been recovered from sewage effluent from abattoirs, streams, drains and fertilizer [3, 15], surviving in drinking water for 4–5 days and sewage for 10–14 days [5]. It was long thought that the organism could live in soil indefinitely, and early reports suggested that the source of infection was soil [5]. However, studies by Wood [58] did not support this widely held belief; *E. rhusiopathiae* survived for a maximum of only 35 days, depending on temperature and soil condition. Despite this limited endurance, the organism does survive long enough in soil to be a potential source of infection to animals and man.

*E. rhusiopathiae* persists in animal tissues for long periods, despite chilling, freezing, or curing [6]. The organism is resistant to pickling, smoking and salting [14]. *Erysipelothrix* can also survive in decaying tissue, and will remain viable in a carcass for 12 days in direct sunlight, for 4 months in putrefied flesh, for 9 months in a buried carcass and at least 10 months in refrigerated tissue [19]. *E. rhusiopathiae* has been isolated from fresh fish, pork and chicken for human consumption [59, 60]. The widespread distribution of *E. rhusiopathiae* can be attributed to the ability of the organism to survive for long periods in the environment, and the fact that the organism can colonise or infect a wide variety of animals [6].

Human infection

Risk of humans infection is based on the opportunity for exposure, and factors such as age, sex, race and socio-economic status relate only to this opportunity [3, 61]. Individuals involved in occupations or recreations with contact with animals, animal products or animal wastes are at greatest risk. Thus, *E. rhusiopathiae* infection is said to be occupationally related [56]. It follows that those in occupations with most frequent animal contact, such as butchers, abattoir workers, veterinarians, farmers, fishermen, fish-handlers and housewives are the most commonly infected [5, 6, 19, 62]. However, cases have been documented from a very wide variety of occupations (see [1] for a complete list). The common names for human infection reflect this occupational mode of acquisition. These include seal finger, whale finger, blubber finger, fish hand, fish poisoning, fish handler’s disease and pork finger [3, 6, 62, 63].

Human infection can occur from contact with infected animals, their secretions, wastes or products, or organic matter contaminated by any of these [6]. Infection is initiated either by an injury to the skin with infective material or when a previous injury is contaminated. There have been a few documented instances of penetration of the skin by the bacteria [61], and of infection by ingestion of contaminated food products [62]. There have been no reports of person-to-person transmission [3]. Modes of infection tend to be very occupation-specific, and transmission is generally by vehicles. These include contaminated objects causing wounds, such as knives, needles, dissecting instruments, fish teeth and spines, fish hooks, bone splinters, and crab, lobster and crayfish claws [6]. If a wound is already present, infection can result from contact with any of a very wide variety of contaminated objects [6, 62].

Infections in both man and animals appear to have a seasonal incidence, with most cases occurring in the summer months [6]. While it has been suggested that the biological activity of *E. rhusiopathiae* is related to temperature [55], others think that it is likely to be due to increased contact between people and sources of infection during these months [6].

Clinical manifestations in man

*E. rhusiopathiae* can cause three forms of human disease which closely resemble disease in swine. These
are erysipeloid (a localised cutaneous form), a generalised cutaneous form and a septic form often associated with endocarditis [14]. It is possible that the incidence of human infection is declining. However, infection is possibly under-diagnosed, because of the resemblance it bears to other infections, difficulties in isolation and identification of the causative organism, and the rapid response to empiric antimicrobial therapy [64].

**Erysipeloid**

Erysipeloid is the most common form of human infection [14]. It is an acute localised cutaneous infection, described as a local cellulitis [3]. Erysipeloid usually occurs on the hand or fingers, reflecting the occupational nature of acquisition of the disease [62]; however, lesions have been described on many areas of the body [6].

The incubation period is usually <4 days, but can be up to 7 days after exposure [65]. The infection consists of a distinctive, well-demarcated, slightly elevated violaceous lesion [19]. The peripheral edge spreads slowly as the centre fades [66], and while vesicles are occasionally present, there is no suppuration or pitting [14, 56]. There is associated local swelling, and an intense itching or a severe burning or throbbing pain [6], which is inconsistent with the mild look of the lesion [1]. Systemic symptoms can occur; 10% of cases experience fever and joint ache, and lymphadenitis and lymphadenopathy appear in 33% of patients [67]. Arthritis can manifest in an adjacent joint. The disease is self-limiting and usually resolves in 3–4 weeks without therapy [3], although relapses may occur if untreated [6].

**Diffuse cutaneous form**

This form is more generalised than erysipeloid, and includes the rare cases in which lesions progress from the initial site to other locations on the body, or in which there is development of lesions remote from the site of inoculation [68]. The lesions are similar to those of the localised form, but bullous lesions can also occur [69]. Systemic symptoms are more frequent and include fever, malaise, joint and muscle pain and severe headaches [6], and polyarthritis in rare instances. The clinical course is more protracted and recurrences are more frequent than with erysipeloid [56]. Very few cases have been documented; only 1 of 100 cases reported by Klauder [68] was generalised and none of the 500 cases reported by Nelson [67] was of this form.

**Septicaemia and endocarditis**

A more serious manifestation of *E. rhusiopathiae* infection is septicaemia, to which endocarditis has almost always been linked. In 49 cases of systemic infection in 15 years, 90% were associated with endocarditis [20]. Although septicaemia and endocarditis are relatively uncommon, there does appear to be an increase in incidence [14, 20], which could either reflect increased exposure or improved diagnosis.

*E. rhusiopathiae* endocarditis has a mortality of 38% and presents as an acute or subacute form, the latter being more frequent. In their summary of cases of endocarditis, Gorby and Peacock [20] discussed predisposing factors, and comparisons with other forms of endocarditis were made. *E. rhusiopathiae* endocarditis had an increased male to female ratio, possibly the result of occupational exposure, and mortality was almost double the rate of endocarditis of other aetiologies. The majority of patients had normal native heart valves and were immunocompetent. A history of alcohol abuse, believed to be a risk factor for the development of this complication, was noted in 33% of patients. Only 36% reported a previous erysipeloid lesion, and 89% of patients were in occupations involving contact with animals.

Recent reports [16, 70] have demonstrated that *Erysipelothrix* bacteraemia without endocarditis is more common than was thought previously, occurring mainly in immunocompromised patients. This increased rate was linked to a more thorough identification of blood culture isolates from these patients.

**Miscellaneous infections**

There are reports of other infections associated with *Erysipelothrix*. These have included chronic arthritis [69], cerebral infection, [71, 72] and osseous necrosis [68]. Recent case reports have focused on novel presentations and complications of *Erysipelothrix* infection. These have included erysipeloid with co-existing orf [64], persistent bacteraemia in a hospitalised patient [73], bacteraemia in an HIV-positive patient [74], endocarditis with acute renal failure [75], septicaemia and lupus nephritis [76], and septicemia in a neonate [77]. In reports of systemic infection, the typical predisposing factors have been involved: either immunocompromised patients with atypical infection without cardiac involvement or immunocompetent patients with endocarditis. Renal involvement and alcoholism were factors noted in this second group.

*E. rhusiopathiae* has been implicated recently in a syndrome known as ‘crayfish poisoning’, which affects lobster fishermen in Western Australia and bears a clinical resemblance to erysipeloid [78]. A possible association between the organism and infections in fishermen was noted in 1947 by Sheard and Dicks [79], but proper identification was hampered by field conditions. It was not until 1996 that this infection was investigated further. The presence of other potential pathogens did not allow a direct causal relationship between *E. rhusiopathiae* and ‘crayfish
poisoning’ to be established; however, the results were suggestive of this [78].

**Treatment and prevention**

Susceptibility data are still limited [3], despite recent reports on the subject [9, 80, 81]. *Erysipelothrix* is highly susceptible to penicillin, cephalosporins and clindamycin [20, 23, 80]. Most strains are resistant to aminoglycosides, trimethoprim-sulphamethoxazole, polymyxins, sulphonamides, streptomycin, novobiocin and vancomycin. The organism is variably susceptible to chloramphenicol, tetracyclines and erythromycin [3].

Erysipeloid can be treated effectively with oral penicillin [3]. Although infection is usually self-limiting, relapses and progression to more serious forms are possible. Oral penicillin will resolve a case of erysipeloid in around 48 h, while intravenous penicillin is recommended for more serious *E. rhusiopathiae* infections [14]. While the mortality rate for endocarditis has been reduced from 100% in the pre-antibiotic era, there is still a 38% fatality rate despite available treatment [20]. This rate could be partly explained by the use of vancomycin, to which the organism is resistant, in empiric therapy for endocarditis [81]. Therefore, early diagnosis of all forms of *E. rhusiopathiae* infection is essential [14]. In those individuals allergic to penicillin, cephalosporins have been described as the most appropriate alternative, as clindamycin and erythromycin are only bacteriostatic towards *E. rhusiopathiae* [3].

Before the advent of penicillin therapy, some alleviation of symptoms could be achieved with hyperimmune serum. However, the resulting serum sickness was often more severe than an episode of erysipeloid, and this treatment was confirmed to be of little value for cutaneous infections [14, 82]. Commercial vaccines in the form of bacterins, lysates or live attenuated strains of *E. rhusiopathiae* serotype 2 offer protection to pigs and turkeys [25]. Vaccination is not a viable option in man, because clinical erysipeloid appears to convey little or no immunity [55, 66], as evidenced by relapse and/or re-infection.

Since Heilman and Herrel [52] first reported the success of penicillin therapy for *Erysipelothrix* infections, there has been no recorded resistance of the organism to this antibiotic. One experiment involving serial passage of 75 strains for 8 months did not produce resistance [14]. Plasmids were not found in *E. rhusiopathiae* in early investigations [80, 81], but later studies were able to detect them [83]. Plasmids appear to play no critical role in *E. rhusiopathiae* resistance. Antibiotics contained in animal feed have been reported to influence the resistance of some strains of *E. rhusiopathiae*, although the mechanism remains uncertain [9].

Containment and control of *E. rhusiopathiae* are the most effective means of preventing the spread of infection in man and animals. An awareness of the infection is essential for preventing the spread of the organism throughout a work environment. *E. rhusiopathiae* can be killed by commonly available disinfectants [19]. However, many investigators have noted that structurally complex equipment is difficult to clean, and because the organism is able to survive in organic matter, disinfecting without cleaning is useless [85, 86]. If disinfection is impractical, other control measures become even more significant.

Control of reservoir populations of *E. rhusiopathiae* is impractical or impossible, because of the widespread distribution of the organism, the large variety of animal hosts and its ability to persist in the environment. However, the possibility of human infection can be reduced by awareness, safe work practices and sensible precautions.

**Isolation and laboratory identification**

It has been reported that medical practitioners who see cases of human erysipelas regularly find the lesion and other symptoms are ‘so typical that a biopsy and subsequent isolation is neither necessary nor justifiable’ [87]. As a result, the majority of identification protocols have been developed with swine erysipelas in mind. However, if human cases are declining, doctors could be less likely to recognise the infection, and, therefore, isolation techniques and methods for identification of *E. rhusiopathiae* may gain new importance.

**Cultural methods**

Traditional cultural methods for *E. rhusiopathiae* isolation involve the use of selective and enrichment media. Identification is based on Gram’s stain, cultural morphology, motility, haemolytic characteristics and biochemical properties, particularly H$_2$S production [3]. For bacteraemia or endocarditis, a blood sample cultured in standard blood culture media is sufficient for isolation [19], as *E. rhusiopathiae* is not particu-
larly fastidious [3]. The organism is more difficult to isolate from cases of erysipeloid, because it is said to live deep in the skin [2]. A biopsy at the advancing edge of the lesion and extending the entire thickness of the dermis is required. Aspirates of the lesion or associated bullae and vesicles are usually less rewarding [19]. Swabbing does not usually detect the pathogen [65].

Biopsies and aspirates are incubated in an infusion broth of glucose 1%, in air or CO2 5–10% and subcultured to blood agar every 24 h. If a sample is likely to be heavily contaminated, such as from soil, faeces, or animal tissue, selective measures are required [2]. A number of selective and enrichment broths have been described. The most commonly used is Erysipelothrix selective broth (ESB), a liquid medium containing serum, tryptose, neomycin, vancomycin and kanamycin [88]. Packer’s medium (SACV) makes use of the organism’s tolerance of sodium azide and crystal violet [29], and is frequently used for subculture after growth in ESB [18]. Modified blood azide (MBA) is similar to SACV, but does not include crystal violet [89]. Bohm’s medium contains azide, kanamycin, phenol and water blue [8].

Incubation of liquid media at 37°C for 18–24 h is usually sufficient for growth. Colonies are visible after 24 h on most solid media, although incubation for 48 h is recommended for ESB and 72 h for SACV. An alternative for biopsy specimens is to refrigerate them at 4–5°C for 4–5 weeks in a liquid enrichment media and then subculture to SACV [15]. Each medium has advantages, but none is ideal. ESB is still regarded as the best selective medium, despite a report showing that some strains grow poorly due to kanamycin susceptibility [90]. MBA requires less incubation time than SACV, but is not as selective and is not suitable for heavily contaminated samples [89]. Bohm’s medium does not seem to have been widely used, although the reasons for this are unclear.

Mouse protection test

This test is traditionally regarded as the best confirmatory test of E. rhusiopathiae identity, because most strains of the organism are highly virulent for laboratory mice. One group of animals is inoculated subcutaneously with a 24-h broth culture and equine hyperimmune E. rhusiopathiae antiserum, and the second group receives broth culture but not antiserum. If the organism is E. rhusiopathiae the second group, but not the first, will die within 5–6 days [2]. However, strains must be pathogenic for mice for this method to be useful. While this test is useful in studies of swine erysipelas, the reliability of this method for human pathogenic strains remains unknown.

Fluorescent antibody test

Direct and indirect assays have been used to confirm the identity of E. rhusiopathiae in tissues [91, 92], in broth [93] and in human infection [94]. However, Harrington [93] noted that this method was not as sensitive as cultural methods, and as a result it has not been used widely.

API Coryne system

The API Coryne system (bioMérieux) is a commercial strip system for the identification of coryneform bacteria. Soto [28] compared conventional biochemical reactions with the commercial system for corynebacteria and related genera, including Erysipelothrix. The system had few misidentifications and all four strains of E. rhusiopathiae tested were correctly identified by use of the strip. The investigators concluded that the commercial strip was a good alternative to traditional biochemical methods, permitting reliable and rapid identification of coryneform bacteria.

PCR

Two PCR methods are available for the detection of Erysipelothrix species [45, 95]. While they were both developed for swine erysipelas, a PCR method was employed for detection of organisms in human specimens in a recent study in Australia [78]. Makino et al. [95] based their primers on a region of the 16S rRNA gene, specific to Erysipelothrix but shared by both E. rhusiopathiae and E. tonsillarum. Shimoji et al. [45] made use of the avirulent transposon mutant created during capsule studies to develop an E. rhusiopathiae-specific PCR. The primers were designed from sequences presumed to be associated with the virulence of E. rhusiopathiae. This assay would be particularly useful for monitoring of swine disease. However, the value of this test in a human clinical situation is uncertain, due to the lack of information regarding the pathogenicity of E. tonsillarum for humans. The time saved by using the PCR is the greatest advantage it has over all other methods for detecting Erysipelothrix.

Erysipelothrix PCR is very sensitive; Makino et al. [95] detected <20 bacteria in a mouse spleen, although the limit detected by Shimoji [45] was only 1000 per reaction mixture, despite a broth enrichment procedure and the use of a DNA extraction kit. Other advantages include the ability to detect an organism in a contaminated sample, as the primers recognise only the specific sequences. With the Makino PCR, the organism does not have to be alive for detection. However, a major drawback is that organism viability cannot be assessed and other procedures would be required for this task. The use of broth enrichment as described by Shimoji [45] could be used to overcome this problem; a modification of this technique was also used in the Australian study [78].
PCR is a powerful tool for the detection of Erysipelothrix in all kinds of samples and is able to overcome many problems inherent in other diagnostic methods. However, it is likely that a combination of culture and molecular techniques will be used for accurate diagnosis in the future.

Conclusions

Although uncommon, it is likely that human infections with E. rhusiopathiae are under-diagnosed. The organism’s slow growth and small colony size mean that it may be overlooked in the routine diagnostic laboratory or overgrown with secondary pathogens such as Staphylococcus aureus and Streptococcus pyogenes. A high index of suspicion and the application of modern molecular techniques will no doubt improve this situation. As a pathogen of animals, particularly swine, Erysipelothrix is of great economic importance and good animal husbandry practice is essential to reduce impact. However, the organism’s resilience and ability to survive are important in both human and veterinary medicine. Most human infections result from occupational exposure and this possibility can be reduced through awareness and safe work practices.

References

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