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Streptococcus pyogenes infection in a free-living European hedgehog (*Erinaceus europaeus*)

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Abstract

Streptococcus pyogenes, a common pathogen of humans, was isolated from the carcass of a free-living European hedgehog (*Erinaceus europaeus*) found in northern England in June 2014. The animal had abscessation of the deep right cervical lymph node, mesenteric lymph nodes and liver. The *S. pyogenes* strain isolated from the lesions, peritoneal and pleural cavities was characterised as *emm* 28, which can be associated with invasive disease in humans. This is the first known report of *S. pyogenes* in a hedgehog and in any free-living wild animal that has been confirmed by gene sequencing. As close associations between wild hedgehogs and people in England are common, we hypothesise that this case might have resulted from anthroponotic infection.

Streptococcus pyogenes (Lancefield group A streptococcus; [GAS]) is a Gram-positive facultative anaerobic coccus which is the main cause of bacterial pharyngitis in humans as well as causing severe invasive disease, such as streptococcal toxic shock syndrome (Cunningham 2000). The rate of GAS infections reported in 2006 from England, Wales and Northern Ireland was 3.3 per 100,000 population, with the highest levels in the young and the elderly (Lamagni et al., 2008).

The European hedgehog (*Erinaceus europaeus)* has a range that extends across Europe and is widely distributed in Great Britain despite recent dramatic population declines (Harris et al., 1995; Wembridge, 2011). Relatively little is known about the infectious diseases of hedgehogs, but there have been reports of potentially zoonotic infections in this species including *Salmonella* spp., *Borrelia* spp., *Cryptosporidium* sp. and mecC-positive *Staphylococcus aureus* (Meredith and Milne, 2009; Riley and Chomel, 2005; Skuballa, et al., 2007; Widen et al., 1996).

A wild hedgehog was found dead in June 2014 within a suburban garden in Northumberland, England. Wild hedgehogs had been regularly observed at the site but were not provided with supplementary food, although this could have occurred in neighbouring properties. The garden was frequented by five apparently healthy domestic cats. No other domestic or livestock species were reported in the immediate vicinity. No previous sign of ill-health was seen in any wild hedgehog prior to finding the carcass of the

submitted animal. A post-mortem examination was conducted consisting of systematic external and internal inspection of all body systems, combined with microbiological and histopathological examinations, as described below.

The hedgehog was an adult male and was in poor body condition with severe, diffuse periodontal disease and an oronasal fistula connecting the empty socket of the upper right first incisor with the right nasal cavity.

Several tissues, including the liver, the right deep cervical lymph node and multiple mesenteric lymph nodes, contained multiple abscesses containing yellow turbid fluid. Aseptically collected tissue samples were plated onto each of the following media: (a) Columbia blood agar supplemented with 5% horse blood (CBA) (Oxoid Ltd., Basingstoke, UK) incubated under aerobic, anaerobic conditions and in air supplemented with CO_2; (b) Columbia agar with chocolate horse blood (Oxoid Ltd.) incubated in air supplemented with CO_2; (c) Xylose-Lysine Deoxycholate (XLD) agar (Oxoid Ltd.) incubated under aerobic conditions. All culture media were incubated at 37 °C and reviewed at day one, two and five. Samples from the right deep cervical lymph node, hepatic abscess, peritoneal cavity and pleural cavity yielded a pure, confluent growth of small (< 1 mm diameter), translucent, entire, butyrous, beta-hemolytic colonies on the CBA plate after 48 hours of anaerobic incubation. The isolate, a Gram-positive coccus, was both oxidase-and catalyse-negative and was identified as a beta-hemolytic *Streptococcus* sp. Lancefield group A using a Prolex TM Lancefield latex agglutination test (Pro-Lab Diagnostics Ltd., Wirral, UK). The isolate was further identified as *S. pyogenes* using API 20 Strep identification strip (bioMérieux UK Ltd.) (API 20 Streptococcus profile 0160514). Using the CDC *emm* sequencing protocol (http://www.cdc.gov/streplab/protocol-*emm*-type.html), this was characterized as an *emm* 28 strain. No other significant microbial isolates were obtained.

Samples from brain, heart, kidney, liver, lung, spleen, trachea, small and large intestine, the right cervical lymph node were fixed in 10% neutral buffered formalin, embedded in paraffin, sectioned and stained with haematoxylin and eosin, Gram's, Ziehl–Neelsen and Periodic acid–Schiff stains using routine methods. Histopathological examination of the liver showed diffuse congestion, hemorrhagic foci, loss of architecture and multiple focal-to-coalescing areas of acute hepatocellular necrosis with co-localization of Gram-positive cocci. Microscopic examination of the right deep cervical lymph node showed this to contain a mixed infiltrate of degenerative neutrophils and lymphocytes and multiple focal-to-coalescing areas of acute necrosis. No abnormalities were detected on examination of the remaining tissues. No additional organisms were detected on histological examination.

We identified disseminated *S. pyogenes* infection as the cause of death of a wild hedgehog in Britain. The animal had periodontal disease, which is commonly seen in wild hedgehogs (Robinson and Routh, 1999) and a tooth root abscess. These lesions might have been the site of entry for the bacterium, from where it spread to other tissues. To the best of our knowledge, this is the first report of *S. pyogenes* infection in a free-living wild animal that has been confirmed by *emm* gene sequencing and the first report of the *emm* 28 strain in a non-human species.

Prior to the availability of molecular techniques in the 1970s, identification of *S. pyogenes* was performed through the assessment of biochemical characteristics combined with Lancefield grouping. This is considered an inaccurate sole means of identification and is now conducted in conjunction with gene sequencing of the M-protein antigen, known as *emm* typing (Facklam, 2002). Infection with *S. pyogenes* has been reported in wild rodents in the USA (Bell et al., 1958) in which it was associated with disseminated abscessation but as this study pre-dated moleculartyping, these results are unconfirmed. *Streptococcus pyogenes* infection has previously been reported in domestic animals including dogs and cats in Europe and the USA (Greene, 2011; Weese and Fulford, 2011) and in cattle in

Denmark (Henningsen and Ernst, 1937) and Canada (Pullin, 1947). Clinical disease in dogs and cats has not been reported except for one equivocal case of canine conjunctivitis (Sprot et al., 2012). Reports of infection in farmed cattle have been associated with mastitis (Pullin, 1947; Henningsen and Ernst, 1937). Disease associated with the bacterium has also been recorded in captive wild animals that exist in close contact with humans, including a laboratory rhesus monkey (*Macaca mulatta*) with streptococcal toxic shock syndrome (García et al., 2006) and farmed white-tailed deer (*Odocoileus virginianus*) with pneumonia (Whitlock, 1939), both reported in the USA.

Until the current study, *S. pyogenes emm* 28 had not been reported from any non-human species. In humans, *emm* 28 strain can be associated with both superficial and invasive disease and accounts for a third of the cases of puerperal sepsis in Europe (Luca-Harari et al., 2009). In the UK, reports of human cases are sporadic with no reported seasonal or spatial pattern of infection (Public Health England, *unpublished data*).

In humans, *S. pyogenes* is transmitted via aerosol-inhalation, skin and mucous membrane exposure to nasal, pharyngeal or wound secretions and, rarely, via contaminated food sources (Boston University Research Compliance, 2012). Whilst we cannot exclude that the hedgehog contracted *S. pyogenes* infection from another (non-human) animal, such as a domestic pet or a mastitic cow, the rarity of described non-human infections indicates that this is unlikely. We postulate that infection in this case occurred due to anthroponotic transmission.

Direct human-hedgehog contact frequently occurs in wildlife rehabilitation centers and is a known route of exposure for people to zoonotic diseases (Riley and Chomel, 2005). Indirect contact with hedgehogs may occur within gardens and although fomite transmission is not thought to be a common source of *S. pyogenes* infection in humans, transmission via this route remains a possibility.

The frequency of interactions between humans, domestic animals and wildlife is increased by the anthropogenic loss of wildlife habitat which augments the risk of these groups being exposed to novel pathogens (Daszak et al., 2000; Epstein and Price, 2009). Many studies have focused on zoonotic diseases originating from wildlife but relatively few have concerned anthroponoses (otherwise known as reverse zoonoses), where human pathogens adversely affect captive or wild animals (Messenger et al., 2014). In addition to being a threat to wildlife health, there is the possibility that anthroponoses may spillback to humans (Henningsen and Ernst, 1937). This report of GAS infection in a hedgehog may be an isolated case, but further research on the pathogens affecting wildlife is required to better understand any disease threats to conservation or public health.

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