Acute peritonitis and salpingitis associated with Case Report streptococcal toxic shock syndrome caused by Lancefield group G a-haemolytic Streptococcus dysgalactiae subsp. equisimilis Toshinobu Horii,^{1,2} Sayuri Izumida,³ Kiyotake Takeuchi,⁴ Toyohiro Tada,⁵ Jinko Ishikawa¹ and Koji Tsuboi⁶ ^{1,2}Department of Laboratory Medicine¹ and Division of Infectious Diseases and Control², Correspondence Hamamatsu University School of Medicine, 1-20-1 Handa-yama, Hamamatsu 431-3192, Toshinobu Horii Japan horiihm@hama-med.ac.jp ^{3,4}Department of Clinical Laboratory³ and Department of Obstetrics and Gynecology⁴, Toyokawa City Hospital, 1-19 Koumei-cho, Toyokawa 442-8561, Japan ⁵Department of Pathology, Nagoya City University School of Nursing, Mizuho-ku, Nagoya 467-8601, Japan ⁶Nagoya University School of Health Sciences, 1-1-20 Daiko-minami, Higashi-ku, Nagoya 461-8673, Japan The authors treated a patient for what appears to be the first reported occurrence of acute peritonitis and salpingitis associated with streptococcal toxic shock syndrome (STSS). This was caused by Lancefield group G α -haemolytic Streptococcus dysgalactiae subsp. equisimilis TKCH2004-001. The isolate showed M protein type stc36 and carried the spegg gene. To the best of the authors' knowledge, the present report represents the first case of STSS complicating acute peritonitis and salpingitis caused by Lancefield group G α-haemolytic S. dysgalactiae

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Introduction

Streptococcus dysgalactiae subsp. equisimilis can be classified by a combination of phenotypic and genotypic characterization into various groups: Lancefield group C α -haemolytic, Lancefield group C β -haemolytic, Lancefield group G β -haemolytic and Lancefield group L β -haemolytic S. dysgalactiae subsp. equisimilis (Vieira et al., 1998). Lancefield group C and G S. dysgalactiae subsp. equisimilis (GCS and GGS, respectively) account for 5-8% of human streptococcal infections, including bacteraemia, wound infections, otitis media, purulent pharyngitis and streptococcal toxic shock syndrome (STSS) (Bisno et al., 1996; Hashikawa et al., 2004; Keiser & Campbell, 1992; Lopardo et al., 2005; Natoli et al., 1996; Roth et al., 1999; Turner et al., 1993; Wagner et al., 1996; Woo et al., 2001). Streptococci are a rare cause of peritonitis and salpingitis (Berenguer et al., 1992; Stevens et al., 1989; Vuilleumier & Halkic, 2001).

subsp. equisimilis.

Recently, several cases of STSS caused by GGS have been described (Hashikawa *et al.*, 2004; Woo *et al.*, 2003). Underlying diseases in most patients with GCS- and GGS-related STSS have included diabetes mellitus, malignant neoplasms, cirrhosis and burns (Hashikawa *et al.*, 2004; Lopardo *et al.*, 2005; Wagner *et al.*, 1996; Woo *et al.*, 2003). In the present study, we characterized a clinical isolate of GGS that caused acute peritonitis and salpingitis, with progression to STSS in a patient with a history of endometriosis and resection of an endometriotic ovarian cyst.

Case report

A 43-year-old woman with a history of endometriosis and resection of an endometriotic ovarian cyst was admitted to the Toyokawa City Hospital in July 2004 because of abdominal pain with a sudden mid-menstrual onset. Laboratory results included a white blood cell (WBC) count of 11 600 mm⁻³, a platelet count of 402×10^9 l⁻¹ and a C-reactive protein (CRP) concentration of 6.6 mg l⁻¹. On hospital day 2, shock developed, with hypotension, hypothermia, clouding of consciousness and progressive ascites. In an emergency adnexectomy, the left fallopian tube and ovary were resected. α -Haemolytic *S. dysgalactiae* subsp.

Abbreviations: ARDS, acute respiratory distress syndrome; GCS, group C *Streptococcus dysgalactiae* subsp. *equisimilis*; GGS, group G *Streptococcus dysgalactiae* subsp. *equisimilis*; STSS, streptococcal toxic shock syndrome.

equisimilis TKCH2004-001 was isolated from ascitic fluid. After initiation of treatment with a carbapenem, panipenem (0.5 g intravenously twice daily; Sankyo), three sets of blood samples were collected for aerobic and anaerobic cultures; no isolate was recovered. Pathologic examination of the resection specimen disclosed a ruptured endometriotic cyst of the left ovary. Pus consisting of coccoid organisms, inflammation, and fibrin were present both in the ruptured ovary and the left tube (Fig. 1). The pathologic diagnosis was acute inflammation of the left tube and ovary with exudative serositis and infections with cocci. On hospital day 3, the patient developed acute respiratory distress syndrome (ARDS) and disseminated intravascular coagulation, dying of septic shock on day 5.

S. dysgalactiae subsp. equisimilis TKCH2004-001 was identified on the basis of phenotypic analysis (API 20 Strep, bioMèrieux Vitek). Lancefield serologic testing (Seroiden Strepto kit; Eiken Chemical) assigned the isolate to group G. The isolate was stored at -70 °C in heart infusion broth (Nissui Pharmaceutical) containing 2% Supplement F (Nissui Pharmaceutical) and 20% (v/v) glycerol. Before further testing, the bacteria were inoculated on trypticase soy agar II with 5% sheep blood (TSA II, Nippon Becton Dickinson) and incubated at 37 °C overnight. Phylogenetic analysis was performed by 16S rRNA gene (rDNA) sequencing, as described elsewhere (Hashikawa *et al.*, 2004). A 1272 bp sequence was amplified by PCR, corresponding to positions 49–1380 of the 16S rDNA sequences of clinical

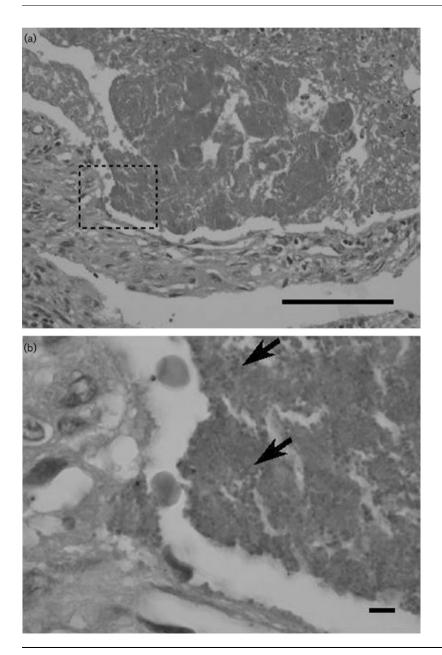


Fig. 1. Histopathologic appearance of cocci (b, arrow) in a meshlike fibrin network in the lumen of the fallopian tube. Panel (b) is a higher-power view (\times 1700) of the area delineated by a broken square in (a) (\times 425). Haematoxylin and eosin staining. Bar, 100 µm. isolates of *S. dysgalactiae* subsp. *equisimilis*, including Hiroshima, Fukuoka, A-22-84, A-ka and A-sa strains (Hashikawa *et al.*, 2004). *Smal*-restricted fragment patterns of chromosomal DNA from the present *S. dysgalactiae* subsp. *equisimilis* TKCH2004-001 and Hiroshima, Fukuoka, A-22-84, A-ka and A-sa were compared by PFGE. The restriction patterns of *S. dysgalactiae* subsp. *equisimilis* TKCH2004-001 and A-22-84 were indistinguishable from each other (data not shown).

MICs were determined by an Etest method (AB Biodisk) in accordance with the manufacturer's instructions. MICs of β -lactams, macrolides and fluoroquinolones for S. dysgalactiae subsp. equisimilis TKCH2004-001 were as follows: $0.016 \ \mu g \ ml^{-1}$ for penicillin G, $0.125 \ \mu g \ ml^{-1}$ for oxacillin, $0.19 \ \mu g \ ml^{-1}$ for ceftazidime, $0.016 \ \mu g \ ml^{-1}$ for cefpirome, $0.032 \ \mu g \ ml^{-1}$ for cefepime, $<0.016 \ \mu g \ ml^{-1}$ for imipenem, $0.064 \ \mu g \ ml^{-1}$ for clarithromycin, $0.5 \ \mu g \ ml^{-1}$ for azithromycin, $0.064 \ \mu g \ ml^{-1}$ for clindamycin and $0.38 \ \mu g \ ml^{-1}$ for levofloxacin. The M protein type was determined as stc36 according to an M protein gene (emm) typing protocol by the sequencing of the emm gene (http://www.cdc.gov/ncidod/biotech/strep/strepindex.html). Detection of superantigenic genes was performed by PCR; primers for the speA, speB, speC, spegg and speF genes were designed as described elsewhere (Hashikawa et al., 2004). The spegg gene, but none of the other superantigenic genes, was detected in S. dysgalactiae subsp. equisimilis TKCH2004-001.

Discussion

The presently reported patient developed acute peritonitis and salpingitis caused by GGS, with progression to septic shock, ARDS and multiple organ failure. This is compatible with the well-recognized syndrome of STSS (Alouf & Müller-Alouf, 2003; The Working Group on Severe Streptococcal Infections, 1993). S. dysgalactiae subsp. equisimilis TKCH2004-001 was isolated from ascitic fluid, while three sets of blood cultures following initiation of panipenem administration failed to recover an organism. Underlying diseases included endometriosis with previous resection of an ovarian cyst, an unusual setting for GCS- and GGSrelated STSS (Hashikawa et al., 2004; Wagner et al., 1996; Woo et al., 2003). A routine cervical smear examination 1 month previously may have been a risk factor for infection. S. dysgalactiae subsp. equisimilis TKCH2004-001 showed susceptibility to antimicrobials, including β -lactams, but therapy with panipenem failed to reverse STSS. Streptococci are a rare cause of peritonitis and salpingitis (Berenguer et al., 1992; Stevens et al., 1989; Vuilleumier & Halkic, 2001). Few previous cases of peritonitis involving STSS caused by Streptococcus pyogenes have been reported in English (Stevens et al., 1989; Vuilleumier & Halkic, 2001). Thus, to the best of our knowledge, the present report represents the first case of STSS complicating acute peritonitis and salpingitis caused by GGS. The case supports the view that GGS can cause overwhelming septic states.

Both α -haemolytic and haemolysin-deficient isolates of S. dysgalactiae subsp. equisimilis have been identified phenotypically by the API 20 Strep system as well as genotypically by 16S rDNA sequencing (Dierksen & Tagg, 2000; Woo et al., 2003). Our α-haemolytic isolate, TKCH2004-001, was identified by both methods as S. dysgalactiae subsp. equisimilis. An earlier study proposed that streptococcal superantigens may be responsible for development of STSS caused by GCS and GGS (Hashikawa et al., 2004). In the present case, S. dysgalactiae subsp. equisimilis TKCH2004-001 carried the spegg gene, which is a homologue of the S. pvogenes superantigenic gene speG, although involvement of Spegg in the pathogenesis of GGS-related STSS remains unclear. It would be advisable to determine if the superantigenic genes are actually expressed, since superantigenic genes such as speB are found in a number of S. pyogenes isolates, although the degree of expression of the genes varies from strain to strain (Nagamune et al., 2005). M protein, known to be a major virulence factor of S. pyogenes, inhibits activation of the alternative complement pathway and impedes phagocytosis by polymorphonuclear leukocytes (Campo et al., 1995). Although an earlier study found the M proteins of GCS and GGS isolated from STSS patients to be highly variable (Hashikawa et al., 2004), the M protein type identified in our TKCH2004-001 isolate was stc36, which had also been identified in GGS A-22-84, isolated as a causative pathogen from an STSS patient at a location in Japan far from Toyokawa City (Hashikawa et al., 2004). In the present study, the PFGE genotypes of S. dysgalactiae subsp. equisimilis TKCH2004-001 and A-22-84 were indistinguishable from one another. Our results suggest that it is important to monitor genetically clonal dissemination involving certain GGS among STSS patients.

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