Detection of *Treponema pallidum* subsp. *pallidum* from Skin Lesions, Serum, and Cerebrospinal Fluid in an Infant with Congenital Syphilis after Clindamycin Treatment of the Mother during Pregnancy

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We report here a case of congenital syphilis in a newborn after clindamycin treatment in pregnancy. Using PCR detection of tmpC (TP0319) and DNA sequencing of the genes TP0136 and TP0548, DNA sequences identical to *Treponema pallidum* subsp. *pallidum* strain SS14 were detected in the infant’s skin lesions, serum, and cerebrospinal fluid.

CASE REPORT

A 2,200-g male infant was born by emergency cesarean section for placenta previa after a 35-week gestation to a 20-year-old secundigravida mother; the amniotic fluid was turbid.

The infant’s mother was diagnosed with secondary syphilis in the 15th week of pregnancy. She had a macular rash, and her serological findings were as follows: a rapid plasma reagin (RPR) test result of 1:64, microhemagglutination assay for antibodies to *Treponema pallidum* (MHA-TP) positive, fluorescent treponemal antibody-absorption (FTA-ABS) test positive, enzyme-linked immunosorbent assay (ELISA) immunoglobulin G (IgG) positive, and ELISA IgM positive. Her husband had no clinical signs of the disease at that time, but his serology was positive: RPR of 1:32, MHA-TP positive, FTA-ABS positive, ELISA IgG positive, and IgM negative. The mother was given clindamycin therapy due to a history of allergy to penicillin, which was not tested by a penicillin skin allergy test. There was a significant decrease in the RPR titer at the time of delivery. The serological findings were as follows: RPR of 1:16, MHA-TP positive, ELISA IgG positive, and ELISA IgM slightly positive.

The infant developed respiratory distress shortly after birth. He required mechanical ventilation and was therefore admitted to the neonatal intensive care unit (NICU) with a presumptive diagnosis of congenital syphilis. Upon admission to the NICU, examination revealed a liver palpable at 3 cm below the right costal margin in the midclavicular line, and a spleen palpable 2 cm below the left costal margin. Abdominal gray-scale ultrasound showed no sonographic abnormalities of the hepatic parenchyma or the biliary structures. Long-bone radiographs demonstrated osteochondritis of both upper and lower extremities, a finding consistent with a diagnosis of congenital syphilis. Laboratory examination revealed a white blood cell count of $6.1 \times 10^9$ cells/liter (with 28% neutrophils), a hemoglobin level of 100 g/liter, and a platelet count of $11 \times 10^9$ cells/liter. The C-reactive protein level was 138.1 mg/liter.

The infant’s skin changes were the most significant with regard to the diagnosis of congenital syphilis. There were numerous vesicles with nonhemorrhagic content on his palms, fingers, soles, and crura. The vesicles on the infant’s fingers and soles were stripped open, baring shallow erosions (Fig. 1). There were suppurative margins in the areas adjacent to the vesicles and shallow erosions and a discrete macular rash without desquamation on the patient’s chest. No nasal discharge was presented.

Therapy with aqueous crystalline penicillin G (100,000 U/kg/day) was initiated, and serological diagnostics and PCR of samples from skin lesions, cerebrospinal fluid (CSF), and serum were performed. Serological diagnostics included RPR, MHA-TP, FTA-ABS, ELISA IgG and IgM, and Western blot IgG and IgM.

Serological findings were as follows: RPR of 1:16, MHA-TP positive, FTA-ABS positive, ELISA IgM positive, ELISA IgG positive, Western blot IgM positive (antibodies to TpN15, TpN17, and TpN47), and IgG positive. CSF serological analysis was interpretable due to contamination with blood and the small sample of CSF taken.

A nested PCR protocol amplifying the tmpC gene (TP0319, which encodes a putative membrane lipoprotein) was used to detect *T. pallidum* subsp. *pallidum* in the clinical samples (3). Molecular analysis revealed *T. pallidum* subsp. *pallidum* in samples from the infant’s skin lesions, serum, and CSF. Sequencing of the genes TP0136 and TP0548 in the PCR-positive samples revealed DNA sequences identical to *T. pallidum* subsp. *pallidum* strain SS14.

On postnatal day 8, mechanical ventilation was finished. On day 14, sepsis developed, and a diagnosis of hepatic insufficiency and ascites was made. A peripheral venipuncture blood...
DNA sequencing of the genes TP0136 and TP0548 identified in the samples using PCR detection of tmpC (TP0319) and DNA sequencing of the genes TP0136 and TP0548.

culture yielded Enterobacter cloacae producing extended-spectrum β-lactamase. This culture showed sensitivity to chloramphenicol, ciprofloxacin, colistin, amikacin, and meropenem. The infant was started on ciprofloxacin therapy. On day 18, the patient’s status rapidly worsened due to hepatic insufficiency, which ended in extreme hyperbilirubinemia (610 μmol/liter), and the infant died.

DNA from skin lesions, blood serum, and CSF was isolated by using a QIAamp DNA minikit (QIAGEN) according to the manufacturer’s instructions. PCR detection of T. pallidum subsp. pallidum was performed as described previously (3). Briefly, the primers tTP0319F (5’-CTGCCTCATCCTCTCTCTA-3’) and tTP0319R (5’-ACCAGACTTCAGCCTATC-3’) were used to produce the first amplicon (773 bp) of nested PCR protocol detecting tmpC gene (TP0319). During the second step, 451-bp PCR product was amplified by using the following primers: iTP0319F (5’-GAAGGTGGTGACTTGAGAAG-3) and iTP0319R (5’-CAACACACCTCCGGCCAT-3’). Each PCR (25 μl) contained 0.125 μl of 10 mM concentrations of each deoxynucleoside triphosphates, 2.5 μl of 10× PCR buffer, 0.25 μl of each primer (100 pmol/μl), variable volumes of sterile distilled water (12 to 21 μl), and the examined DNA isolate (1 to 10 μl). To this reaction, 0.05 μl of Taq polymerase (5,000 U/ml; New England Biolabs) was added. PCRs were amplified as follows: 94°C for 1 min, followed by 30 cycles of 94°C for 30 s, 58°C for 30 s, and 72°C for 1 min, followed in turn by 72°C for 10 min. The second step of the PCR was performed identically with the following exceptions: 1 μl of the PCR containing the 773 bp amplicon was used as a template, and 40 amplification cycles were applied. Final amplicons were analyzed by using 2% agarose gel. Chromosomal DNA of T. pallidum subsp. pallidum comprising TP0136 and TP0548 genes was PCR amplified by using the primers TP0136F (5’-AGTGCTCTTCCTCGTGGTTC-3’) and TP0136R (5’-CACGTGGGTGTCACATT3’), resulting in a 1,207-bp PCR product, and the primers TP0548F (5’-GCCTCTCCATATGATCGTGCTAT-3’) and TP0548R (5’-GAGCCACTTCAGCCCTACTG-3’), resulting in a 1,066-bp amplicon. PCRs were set up as described above, and cycling conditions were as follows: 94°C for 1 min, followed by 30 cycles of 94°C for 30 s, 55°C for 30 s, and 72°C for 2 min, and finally by 72°C for 10 min. The resulting PCR products were purified using QIAquick PCR purification kit (QIAGEN) and subjected to dideoxy terminator sequencing using amplification primers and additional internal primers. Sequence analysis was performed in DNASTAR software (Lasergene).

The number of treponemal DNA copies (inferred from the maximal dilution of PCR-positive samples) was identical in serum and CSF and came to approximately 10^5 DNA molecules per 1 ml of undiluted serum and CSF samples.

Treponemal DNA in congenital syphilis can be detected in a number of clinical samples including serum, whole blood, amniotic fluid, paraffin-embedded placental tissue, bullous targe-

Skin lesions either in the form of a typical vesiculobullous eruption (especially over the palms of the hands and the soles of the feet) or a maculopapular skin rash over the body are common presentations of early congenital syphilis and have been described in several studies (5, 6, 10, 11). Detection of treponemal DNA in whole blood and serum in patients with congenital syphilis appears to be more reliable than in adults (7, 9). This is probably due to the relatively higher concentration of treponemes in the blood of these patients. In our case, the number of treponemal DNA copies emphasizes the fact that in early congenital syphilis the concentration of treponemes in serum and CSF is similar and notably higher than in adults (3). These findings are consistent with the study by Michelow et al. in which central nervous system involvement in infants with congenital syphilis was best predicted by IgM immunoblotting of serum or a PCR assay of serum or blood (7).

Parenteral penicillin G is the only therapy with documented efficacy for syphilis during pregnancy. In experiment on rabbits, single intramuscular doses of clindamycin (15 or 40 mg/kg) did not decrease treponemal counts significantly, but single injections of penicillin (10,000 U/kg) reduced treponemal counts by more than 250-fold. Multiple intramuscular injections of clindamycin reduced counts by five- to sevenfold, whereas multiple doses of penicillin decreased treponeme...
counts by greater than 300-fold. Despite partially crossing the placenta, clindamycin is far less effective than penicillin in treating syphilitic lesions (2, 8). Pregnant women with syphilis who report a penicillin allergy should be desensitized and treated with penicillin. Skin testing may be useful in pregnant women to establish whether a penicillin allergy exists (1).

In conclusion, this report documents the importance of treating syphilis during pregnancy with penicillin and endorses the use of molecular techniques to identify \textit{T. p. pallidum} in clinical samples to diagnose congenital early syphilis.

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