

Pili genes pattern in Group B streptococci from newborn infections and pregnant women in Europe (DEVANI Project)

Monica Imperi¹, Daniela Rinaudo², Roberta Creti¹, Marco Pataracchia¹, Roberto Rosini², Annalisa Nuccitelli², Paula Kriz³, Mogens Kilian⁴, Markus Hufnagel⁵, Androulla Efstratiou⁶, Manuel de la Rosa⁷, Pierrette Melin⁸, Antoaneta Detcheva⁹, Lucilla Baldassarri¹ and Domenico Maione² on behalf of the DEVANI study group

¹ Istituto Superiore di Sanità, Rome - Italy

² Novartis Vaccines & Diagnostics, Siena - Italy

³ National Institute of Public Health, Prague, Czech Republic

⁴ Department of Medical Microbiology and Immunology, Aarhus University, Denmark

⁵ University Medical Center Freiburg - Germany

⁶ Health Protection Agency, Colindale - UK

⁷ Hospital Universitario Virgen de las Nieves, Granada - Spain

⁸ Centre Hospitalier Universitaire de Liege - Belgium

⁹ National Center of Infectious and Parasitic Diseases, Sofia, Bulgaria

Objectives

Evaluation of the presence and expression of genes coding for pili in a collection of group B streptococci (GBS) isolated from newborn infection and pregnant women in the course of the DEVANI (Design of a Vaccine Against Neonatal Infection) project.

Methods

GBS isolates from pregnant women (PW) and cases of newborn infection (NI) were collected in 8 European countries (Belgium, Bulgaria, Czech Republic, Denmark, Germany, Italy, Spain, United Kingdom) during 2009/10 under the auspices of DEVANI. Total no. of strains examined was 1078 and 192 from PW and NI, respectively. Isolates were screened by multiplex PCR and FACS analysis to evaluate respectively gene presence and surface-exposure of pili.

Results

The most common gene patterns found were PI-2a alone, PI 1+2a and PI 1+2b, while the PI-2b gene alone was very rare. The most prominent result was that a majority of isolates from NI carried the PI-1+2b gene pattern, while the most common pattern among PW was PI-1+2a. Most of analyzed strains express at least one pilus on their surface.

Conclusions

All isolates contained at least one gene coding for pili. When present pili 2a and 2b were highly surface exposed.