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edited by:
André de Almeida
Fabrizio Ceciliani
David Eckersall
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Romana Turk



Colostrum protein uptake in neonatal lambs examined by descriptive LC-MS/MS

Lorenzo E. Hernández-Castellano^{1,2*}, Anastasio Argüello¹, André M. Almeida^{3,4}, Noemí Castro¹ and Emøke Bendixen⁵

¹Department of Animal Science, Universidad de Las Palmas de Gran Canaria, Arucas, Spain

²Veterinary Physiology, Vetsuisse Faculty, University of Bern, Bern, Switzerland;

lorenzo.hernandez@vetsuisse.unibe.ch

³Centro Interdisciplinar de Investigação em Sanidade Animal, FMV-UL; Instituto de Investigação Científica Tropical, Lisboa, Portugal

⁴Instituto de Biologia Experimental e Tecnológica and Instituto de Tecnologia Química e Biológica da UNL, Oeiras, Portugal

⁵Department of Molecular Biology and Genetics, Aarhus University, Aarhus, Denmark

Introduction

Colostrum is the first source of nutrition in neonatal ruminants, supplying not only nutrients, but having also a fundamental biological function, promoting immunoglobulin transfer from the dam to the newborn (Hernández-Castellano et al., 2014a). Newborn ruminants are hypo gammaglobulinemic at birth, as the complexity of the ruminant placenta does not allow a sufficient transfer of immunoglobulins (Ig's) from the dam to the foetus (Lérias et al., 2014). Therefore, newborn ruminants depend entirely on passive immunity transfer from the mother to the neonate, through the suckling of colostrum (Hernández-Castellano et al., 2014b). The aim of this study is to describe the proteomes of sheep colostrum and lamb blood plasma after suckling, using SDS-PAGE for protein separation and in-gel digestion, followed by LC-MS/MS of resulting tryptic peptides for protein identification.

Material and methods

Sample collection and preparation

The study was based on individual analysis of plasma samples from 4 single partum lambs (Canarian dairy breed), studied as two experimental groups, as well as two samples from the standard pool of colostrum used for lamb feeding. The experiment took place at the experimental farm of the Veterinary Faculty of the Universidad de Las Palmas de Gran Canaria (Spain). During the experimental period (from birth up to 14 h after birth), the colostrum group (C group) received colostrum feeding at 2 h after birth. The non-colostrum group (NC group) was not fed at 2 h after birth. Blood samples were collected directly before feeding at 2 and 14 h after birth from the jugular vein in 2.5 ml tubes with K-EDTA. Blood was centrifuged at 2,190 g for 5 min at 4 °C and the obtained plasma was frozen at -80 °C until further analysis. Plasma and colostrum samples were analysed at Aarhus University (Aarhus, Denmark). A total of 200 µl from each sample were homogenized with 1