



UNIVERSITÀ DEGLI STUDI DI TORINO

AperTO - Archivio Istituzionale Open Access dell'Università di Torino

The application of a multicolor ZOO-FISH on secondary bovine oocytes showed its potential use for aneuploidy detection.

This is the author's manuscript	
Original Citation:	
Availability:	
This version is available http://hdl.handle.net/2318/150235	since
Published version:	
DOI:10.1007/s10577-014-9435-7	
Terms of use:	
Open Access	
Anyone can freely access the full text of works made available as "Open Access". Works made available under a Creative Commons license can be used according to the terms and conditions of said license. Use of all other works requires consent of the right holder (author or publisher) if not exempted from copyright protection by the applicable law.	

(Article begins on next page)



UNIVERSITÀ DEGLI STUDI DI TORINO

The final publication is available at Springer via http://dx.doi.org/(10.1007/s10577-014-9435-7)

The application of a multicolor ZOO-FISH on secondary bovine oocytes showed its potential use for an euploidy detection

<u>A. Pauciullo¹</u>, A. Perucatti¹, A. Iannuzzi¹, D. Incarnato¹, V. Genualdo¹, L. Pucciarelli¹, M. Rubessa², D. Di Berardino², L. Iannuzzi¹ (alfredo.pauciullo@cnr.it)

¹National Research Council (CNR), ISPAAM, Laboratory of Animal Cytogenetics and Gene Mapping, Naples, Italy; ²University of Naples Federico II, Department of Agriculture, Portici (NA), Italy

The female gametes are more susceptible to chromosome segregation errors during meiosis I division and therefore they are the major contributors to the embryo aneuploidies. The evaluation of aneuploidies in bovine oocytes is useful for monitoring the reproductive health of this species and FISH is the main method employed for this purpose. To date only 2-3 chromosomes were simultaneously investigated for aneuploidy detection in cattle oocytes. In this work we propose a multi-color ZOO-FISH by the simultaneous detection of 6 specific chromosome painting probes on secondary oocytes matured in vitro. Standard procedures were employed for 24h in vitro oocytes maturation, whereas specific autosomal probes were prepared by microdissection and DOP-PCR using river buffalo mitosis (2n=50). Probes were labelled with spectrum-green and -orange in a second DOP-PCR. Three sequential rounds of FISH were achieved for the same slides. Each round was realized using two probes simultaneously hybridized on MII oocytes with the corresponding first polar bodies (I pb). Slides were counterstained with DAPI in antifade. Digital images were captured in gray-scale and pseudocolored by the software. Six specific probes, painting 3 out of 5 sub-metacentric river buffalo chromosomes (BBU 1p, 1q, 3p, 3q, 4p and 4q) were sequentially hybridized on BTA secondary oocytes with the corresponding (I pb). The different colors of the probes allowed the identification of 6 cattle chromosomes (BTA 1, 5, 8, 19, 27 and 28) both on MII and polar bodies evidencing no abnormalities for the investigated cells, but confirming their potential use for aneuploidy detection in bovine oocytes. This result opens further opportunity of investigation for clinical cytogenetic applications also in the other species with difficult CGH karyotype.

This study was supported by CISIA-VARIGEAV project, National Research Council (CNR) of Italy and by RARECA project, PSR-214 of the Campania Region, Italy.