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▶ To cite this version:

Laurence Finot, Catherine Hue-Beauvais, Etienne Aujean, Fabienne Le Provost, Eric Chanat, et al.. Delineating the MaSC/progenitors committed to the development in the bovine mammary gland at puberty. Mammary Gland Biology Gordon Conference, May 2018, Barga, Italy. 2018. hal-02791174

HAL Id: hal-02791174 https://hal.inrae.fr/hal-02791174

Submitted on 5 Jun 2020

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Delineating the MaSC/progenitors committed to the development of the bovine mammary gland at puberty



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RESULTS

Xenotransplantation of Holstein heifers mammary explant in mouse mammary gland resulted in the development of an epithelium *in vivo*

We observed the development of epithelial outgrowths in 6 out of 8 transplanted mice (figure 1a), highlighting the presence of MaSC in the bovine mammary explant.



BACKGROUND

During puberty and gestation, Mammary Stem Cells (MaSC) and their progeny drive the development of the mammary gland. In this study, we investigated the epithelial MaSC/progenitor populations within the bovine mammary tissue at puberty, a key physiological stage during which MaSCs expand and differentiate into mature cells.

MATERIEL AND METHODS

Explants (0,2 mm²) were sampled from the mammary gland

Figure 1a : mammary gland whole mount

Immunofluorescence analysis of outgrowths **revealed** bovinespecific epithelial structures containing basal and luminal cells, expressing keratin (KRT)14 and KRT7, respectively, in a dense stromal tissue producing collagen (figure 1b).

Figure 1b : mammary gland IHC 🖨 🚺

Distinct epithelial populations were present during the developmental phase of the mammary ductal-alveolar architecture at puberty.

Flow cytometry highlighted 4 cell populations belonging to the epithelial lineage expressing CD49f and CD24 or not (figure 2a). The mean percentage of the cell population is indicated in red (n=3).



Xenotransplantation assay

Bovine mammary gland explants (0,05 mm²) were transplanted in 3-weeks nude mice (Balb/c AnNRj-Foxn1^{nu/nu}). Mouse mammary glands were collected two months later and processed for whole mount (Carmine Alun staining) and **immunofluorescence** (IHC) **analysis**.

Phenotyping assays

- Bovine mammary gland explants were dissociated to single **cells** using an enzymatic (collagenase/hyaluronidase/trypsin) dissociation protocol.
- Single cells were stained with anti-CD49_f and anti-CD24 antibodies and co-expression of these markers was assessed by flow cytometry.
- The expression of CD10 (a basal lineage marker) and the activity of ALDH1 (a MaSC/progenitor marker) were assessed on the CD49_f / CD24 populations **by flow cytometry**.

Within these 4 epithelial populations, only the CD49_f^{high}CD24^{pos} and the CD49^{*high*}CD24^{*neg*} cells were able to form mammosphere *in vitro* (figure 2b)



The cell populations sorted on the basis of CD49_f and CD24 staining were cultured with Matrigel during 7 days to monitor the formation of **mammospheres**



Most of the MaSC cells (CD49^{*high*}CD24^{*pos*} population) exhibited an ALDH1 activity (70%), a characteristic of the luminal restricted MaSC/progenitor. ALDH1 negative MaSC (30%) would represent a quiescent MaSC pool

	% of cells	% of cells with
Populations	expressing CD10	ALDH1 activity
CD49 ^f low CD24 ^{neg}	7%	87%
CD49 ^f low CD24 ^{pos}	75%	78%
CD49 _f high CD24 neg	93%	0.3%
CD49 _f high CD24 pos	92%	68%

In-depth phenotyping highlighted that MaSC express basal marks and suggested that two subpopulations would mingle in the MaSC pool (in Finot et *al*, 2018; submitted)

Luminal Progenitor Ductal Progenitor CD49^{low} CD24⁻ CD10⁻

Proposed epithelial cell lineage scheme in the bovine mammary gland at puberty

PERSPECTIVES

We aim at providing new insights into the fate of bovine MaSC and progenitors (both in their proportion and their molecular signature) at key physiological stages of development, including lactation and dried off. In an agronomic context as dairy, understanding the fundamentals of the mammary epithelial development and turnover is of importance to improve animal robustness through the enhancement of lactation efficiency.