Establishment of Two Types of Mammary Epithelial Cell Lines from Chinese Holstein Dairy Cow

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Abstract: Chinese Holstein dairy cow is the main dairy cow in China. With the development of Chinese milk industry, more attention is being placed on the factors that might affect milk quality, especially milk fat and milk protein content. So, the objective of this study was to establish a culture system for two types of bovine mammary epithelial cell line, high fat high protein Mammary Epithelial Cell line (MEC-HH) and low fat low protein Mammary Epithelial Cell Line (MEC-LL) in vitro and elucidate the unique characteristics of these cell lines. Mammary tissues from two lactating dairy cows whose milk were high fat high protein content and low fat low protein content were used as sources of these epithelial cell lines. The tissues were cultured in plastic dishes by the method of tissue cultures. Fibroblast and epithelial cells successively grew and extended from mammary tissues after 6 days from low fat low protein cow but 10 days from high fat high protein cow and pure epithelial cells were obtained by continuous culture and digestion for 3-5 times. The epithelial cells cultured in the 10% FBS maintained a normal growth tendency and had normal cell nucleus. The α-s1 casein and κ-casein expression of epithelial cells were detected by RT-PCR. The positive immunoocytochemistry to cytokeratin 18 suggested the obtained cell line exhibited the specific character of epithelial cells. A TG-kit was used to detect the TG content of the two types of cells, the results showed that the cell lines kept the character of high fat and low fat cows in the 1st and 2nd passages. The obtained mammary epithelial cell lines had normal morphology, growth characteristics, cyto genetic and secretory characteristics. Thus, it might supply a useful tool for studying the gene function about milk fat and protein. The research provides the basis for the future study of lipometabolism mechanism in dairy cow.

Key words: Mammary epithelial, Chinese Holstein dairy cow, cell lines, fat, protein, China

INTRODUCTION

In the bovine species, milk production is well known to correlate with mammary tissue. However, most advanced techniques in optimizing milk production relied on improvements of breeding and husbandry practices (Zakizadeh et al., 2007; Gorgulu, 2011; Martignani et al., 2010). The Chinese Holstein cow is the main dairy cow in China. With the fast development of the Chinese milk industry, more attention is being placed on the mechanisms or factors that might affect milk synthesis and quality, especially fat and protein (Han et al., 2011; Huang et al., 2011).

Collagenase dissociation has been used successfully during isolation and culture of bovine epithelial cells in vitro (Baumrucker et al., 1988; Hryhn et al., 1991). Tissue culture is another method which was successfully used to isolate bovine mammary epithelial cells (Rapp and Melnick, 1964; Rose et al., 2002). Mammary tissue cell or explants have been used over the years as a bioreactor model to understand its physiological function (Pantschenko et al., 2000a). Milk synthesis occurs within clusters of differentiated mammary epithelial cell and is controlled by lactogenic hormones and extracellular matrix interactions (Cifrian et al., 1994). A method for establishment of mammary epithelial cell lines from human has been described (Ceriani et al., 1979). Several MEC lines had been existed including BMBC+H (Schmid et al., 1983), MAC-T (Szyda and Komisarek, 2007), PS-BME (Jack and Mather, 1990), BME-UV (Zavitzion et al., 1996) and L-1 (German and Barash, 2002), CMECs (Hu et al., 2009). However, the level of milk protein synthesis of these MEC lines is either very low or non-existent. These cell lines are not suitable for investigations of mammary gland function.

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The study established two mammary epithelial cell lines from Chinese Holstein cows whose milk contained high fat high protein and low fat low protein with tissue culture method and the MECs were thoroughly characterized via morphology, growth curves, staining analysis of cells nucleus, RT-PCR and immunocytochemistry (Pantschenko et al., 2000b). The comparison about the two cell lines were done by detection of TG contents and the isolated MECs have been used for investigation of gene function about milk fat and protein and lipometabolism mechanism study.

**MATERIALS AND METHODS**

**Samples selection and collection of mammary tissue:**

Milk fat and protein content from 11 Chinese Holstein cows which were grown in the same circumstance were analyzed by milk composition analyses machine for 9 days and monitored 2 times per day (Table 1 and 2). The mammary tissue from the No. 33 cows whose milk fat and protein content were highest (fat content was 4.30, protein content was 3.91) and the No.16 cow whose milk fat and protein content were lowest (fat content was 2.41, protein content was 2.80) were selected as samples were butchered in slaughter house for their mammary tissue.

**Isolation of mammary tissue and culture of mammary epithelial cells:** Two types of bovine mammary tissue were obtained from a high-fat high-protein and a low-fat low-protein lactating Chinese Holstein dairy cows. Fresh tissues were placed in an incubation box at 37°C in PBS (Tiangen, Beijing, China) solution with 5% (v/v) penicillin and streptomycin (Tiangen) and immediately transported to the laboratory. Mammary tissues were then washed with PBS solution for several times until the solution was lipid and without milk. Tissues were then cut into about 1 mm² rubble and washed again with PBS solution until tissue was clean. The smaller pieces of tissue were transferred onto cell culture dishes (Falcon, Franklin Lakes, NJ, USA). The basal media was prepared in advance: DMEM/F12 ( Gibco Invitrogen, Carlsbad, California, USA) with 10% (v/v) fetal bovine serum (GIBCO) (Ethier, 1986) and 1% (v/v) penicillin and streptomycin (Tiangen, Beijing, China) and 1% (v/v) epithelial growth factor (OIBCO) which could promote the growth of the epithelial cells. About 2-3 drops of basal media were added ensuring that the tissue was not dry. Then, the cell culture dishes with tissue were incubated at 37°C, 5% CO₂ in a CO₂ incubator (Themo, Marietta, Ohio, USA). After 6 h, 5 mL basal media were added to every dish ensuring that the tissues would not float and separate from the bottom of the culture dish. The basal

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* Milk fat content in No. 16 cow was 2.41% which was the lowest in all the 11 cows, milk fat content in No. 33 cow was 4.30% which was the highest in all the 11 cows, p = 0.000

**Table 2: Milk protein content in 11 Chinese Holstein cows (%)**

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* Milk protein content in No. 16 cow was 2.80% which was the lowest in all the 11 cows, milk protein content in No. 33 cow was 3.91% which was the highest in all the 11 cows, p = 0.000
media was replaced with fresh media every 48 h until culture dish was full of cells. Cells were detached with 0.25% trypsin-0.02% EDTA (GIBCO) and transferred to new culture dishes which were used to remove fibroblasts. Subsequently, the pure mammary epithelial cells were isolated after 3-5 passages.

**Characteristic of mammary epithelial cells by growth curves draw:** About $6 \times 10^5$ cells/well were seeded in 96 well flat-bottom culture plates (Falcon) containing basal media. The number of cells were counted by counting plate under an inverted microscope after every 24 h until 9 days. Before counting the cells, researchers digested all the cells in a well and added 200 μL PBS. Then, researchers dropped 10 μL the mixed PBS solution in the counting plate. Within 9 days, the basal media were replaced with fresh media every 48 h without digestion. The growth curves were drew on the basis of the earlier data.

**Cells nucleus staining analysis by Giemsa stain solution:** The purified cells which were cultured in cover glass slides (Tiangen, Beijing, China) were examined via staining analysis of cells nucleus. The cells were rinsed with PBS solution three times and were suspended with formalin (Tiangen) solution. The slides were then stained with Giemsa stain solution (Tiangen) 1 mL for 15 min, washed with distilled water and dried at room temperature. Cells nucleus were visualized and detected with microscope.

**RT-PCR for α-s1 casein and κ-casein expression in mammary epithelial cells:** Total RNA from mammary epithelial cells were isolated with RNA extraction kit (Tiangen). The expression of α-s1 casein and κ-casein were examined by RT-PCR. The integrity and concentration of the RNA were verified by analyzing 5 μL of each sample on a 1% agarose gel and ultraviolet spectrophotometer (Alpha). Reverse transcription system: 5 μL RNA RT buffer, 4 μL: dNTP mixture (10 mM), 3 μL: Oligo-dT, 1 μL: AMV Reverse transcriptionase (5 U μL$^{-1}$), 1 μL: RNase inhibitor (40U μL$^{-1}$), 0.5 μL; Total RNA, 6 μL, RNase free H$_2$O, 4.5 μL. Reverse transcription procedure: 42°C, 50 min; 95°C, 10 min, 0°C and 5 min. Primers of α-s1 casein and κ-casein were designed with primer 5.0 (Table 2) and synthesized by Shanghai Sangen Biological Engineering Technology Corporation.

**Examination of Cytokeratin 18 by immunofluorescence in mammary epithelial cell:** Cytokeratin 18 was examined by seeding $1 \times 10^5$ cells/well in 6 well flat-bottom culture plates (Falcon). CK18 expression was examined after 24 h. First, the cells were washed by PBS solution and fixed 15 min with ice-cold acetone-formaldehyde (Tiangen) (8:2). Cells were then washed 5 min with PBST for three times. Nonspecific reactivity was blocked with goat serum (Takara, Dalian, China) for 1 h at room temperature and washed 5 min with PBST for 3 times. First antisera (BOSTER, Beijing, China) anti-cytokeratin 18 was diluted 1/100 in PBS solution and incubated for 1 h at room temperature. The cells were washed 3×5 min with PBST. Second, antibody FITC-conjugated monoclonal anti-rabbit IgG (BOSTER) was diluted 1/100 in PBS and incubated in the dark for 30 min. Cells were washed with PBS for three times and visualized with a fluorescence microscope.

**Comparison of the TG content in MEC-HH and MEC-LL for the difference of the two types of cell lines:** To examine cell lines whether kept the characters of high fat and low fat cows. A TG-kit (BHKT, Beijing, China) was used for detecting TG content in both MEC-HH and MEC-LL. The $1 \times 10^5$ cells/well were seeded in a 6 well flat-bottom culture plates, DMEM/F12 without serum were added into the wells. After 48 h, researchers collected the cells then adjusted the concentration to ensure the cells number was at equal. The cells were broke by a transonic machine. The other operation followed the manufacturer’s protocols. To decrease the influence of same culture condition, the cells in passage 2 and 5 were chose.

**Statistical analysis:** All paired comparisons were subject to a two tailed Student t-test with p<0.05 considered statistically significant.

**RESULTS AND DISCUSSION**

**Morphology of MEC-HH and MEC-LL:** For the mammary tissue from the low fat low protein cow, within 6 days but for the mammary tissue from the high fat high protein cow, within 10 days, most cells began to grow outward from the attached organoid or cell clumps. Elongated cells and large polygonal cells spread and grew from the organoid and formed sheets of cells. The cells formed preconfluent monolayers by day 10 for MEC-LL and day 14 for MEC-HH. At this time, two kinds of cells located in the plates. After about 2 weeks, primary epithelial cells became the dominant cell types. Generally, it took more time to digest epithelial cells than fibroblasts with trypsin and EDTA solution. Subsequently, the pure mammary epithelial cells were isolated after 3-5 passages (Fig. 1). We can see, most of the isolated cells that extended from the tissue had a cobblestone like shape, clear boundary and were connected tightly.
Fig. 1: Purified Chinese Holstein cow high fat high protein mammary epithelial cells and low fat low protein mammary epithelial cells. Tissue culture method was employed to culture cells; a, b) symbiosis of fibroblast and epithelial cells separated from cows with high fat high protein milk and low fat low protein milk, respectively; c, d) purified epithelial cells grew from mammary tissue with high fat high protein and low fat low protein, respectively and e, f) epithelial cells separated from cows with high fat high protein milk and low fat low protein milk showed honeycombing

![Cell Growth Curve](image)

Fig. 2: MEC-HH and MEC-LL growth curve. The growth curve conformed to the rule of S sigmoid curve suggested the cells maintained a favorable growth performance

The result of growth curve of the two types of mammary epithelial cells: During the 1st 3 days of the latent phase, the growth rate was slow but during the following 3 days of growth there was an increase in cell number followed by a steady phase for the last several days of culture (Fig. 2). Thus, the growth curve conformed to the rule of S sigmoid curve for proliferating cells with a lag-phase, exponential phase and steady phase. This suggested that the isolated bovine mammary epithelial cells, no matter MEC-HH or MEC-LL, all maintained a favorable growth performance.

The result of staining analysis of cells nucleus: The result from Giemsa staining showed that the cells displayed polygon, the cells nucleus are round or elliptic.

![Giemsa Staining](image)

Fig. 3: The results of Giemsa staining in mammary epithelial cell lines. There were 2-4 nucleoli per cell suggest that cells we isolated were in a active state. a) The result of high fat high protein epithelial cells in Giemsa staining. b) The result of low fat low protein epithelial cells in Giemsa staining

There are 2-4 nucleoli per cell (Fig. 3). The result indicated that the cells we isolated were in an active state and could be used for the followed studies.
Fig. 4: Chinese Holstein cow mammary epithelial cells α-s1 casein and κ-casein protein identification results. RT-PCR was employed for detecting α-s1 casein and κ-casein in MEC-HH and MEC-LL. M for D2000 standard for the molecular weight Marker, 1 lane for the α-s1 casein of MEC-HH, 5 lanes for the κ-casein of MEC-HH, lane 3, 7 for the negative control fibroblasts for α-s1 casein, lane 2 for the α-s1 casein of MEC-LL, lane 6 for the κ-casein of MEC-LL, lane 4, 8 for the negative control fibroblasts for κ-casein.

The RT-PCR result of α-s1 casein and κ-casein expression: Reverse transcription-PCR amplification of mRNA from bovine mammary gland cultures demonstrated the cells we isolated expressed α-s1 casein and κ-casein (Fig. 4). The α-s1 casein is 1736 bp and κ-casein is 780 bp. The RT-PCR result indicated that the mammary epithelial cells we isolated had normal secretion function.

Cytoskeleton 18 expression in mammary epithelial cell lines: Although the cell lines we established appeared to have epithelial cell morphology, we further investigated homogeneity at the sixth passage and also at the resuscitated cells by examining the protein expression of cytoskeleton. Cytoskeleton 18, a specific protein for epithelial was examined by immunofluorescence. Most isolated epithelial cells were positively-stained and the fibroblast cells were negatively-stained as a control (Fig. 5). The results indicated that the cells we isolated were pure epithelial cells without other type of cells. The positive staining for cytokeratin-18 was a powerful result to prove the specific epithelium character of mammary epithelial cells.

TG content in MEC-HH and MEC-LL in passage 2 and 5: We detected TG content in MEC-HH and MEC-LL in passage 2 and 5. The statistical significance of differences in mean values was assessed using SPSS 13.0 Statistic Software. In passage 2, \( p = 0.009 < 0.05 \) was considered statistically significant but in passage 5, \( p = 0.844 > 0.05 \) was considered not significant. Thus, the results indicated that TG content was different between MEC-HH and MEC-LL in passage 2 but in passage 5 TG content in the two types of cells were nearly equal (Table 3 and 4).

To decrease the influence of milk quality in different raised conditions, we chose 11 lactating Chinese Holstein cows grew in the same circumstance. To ensure the data is accurate, we persistently monitored milk quality of the 11 cows for 9 days with milk composition analysis machine and we calculated the average milk fat content and milk protein content then two cows whose milk fat and milk protein were highest and lowest was chose and butchered. The cells isolated from the two cows kept their characteristic about milk fat and protein within 2 passages but in the 5th passage, the two types of cells had almost same TG contents. We considered the possible reason was that cells absorbed nutrition from the same medium in vitro and after cultured a long time, the two types of cells tended to assimilation.

Table 3: The primer design and the length of PCR amplification fragments

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<td>κ-casein</td>
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The previous studies suggested that DMEM and DMEM/F12 were more suitable than F12 and RPMI1640 (Shamay and Gertler 1986). Evaluation of the cell population doubling time suggested that 10% FBS was
the optimal concentration in growth media (Park et al., 1979; Akers et al., 1986). Some results suggested that the concentration of Fetal Bovine Serum (FBS) in media had a strong effect on the proliferation of bovine mammary epithelial cells (Mackenzie et al., 1982). When compared with 0% FBS treatment, bovine mammary epithelial cells grown in 5-10% serum underwent a 3-4 fold increase in cell number during the 12 days of culture (Mackenzie et al., 1982). The results suggested that DMEM/F12 were suitable for the growth of mammary epithelial cells and a frozen solution with 90% FBS and 10% DMSO was optimal for cell preservation. Though at the first, the cells isolated from high fat high protein cow grew slower than cells isolated from low fat low protein cow (the possible reason was that the cells growth rate was influenced by lipometabolism), the grow rate of the two types of cells tend to be equal after 2 passages and there is no significant different between the two types of cell lines in morphology, growth characteristics, cytogetetic and secretory characteristics.

The growth curves showed that the mammary epithelial cells, no matter MEC-HH or MEC-LL maintained a favorable growth performance. During the 1st 3 days, the cells grew slow and stayed in a latent phase. But during the following 3 days of growth there was an increase in cell number, during the last several days, the growth rate was steady. Thus, the growth curve conformed to the rule of S sigmoid curve for proliferating cells (Barcellos-Hoff et al., 1989). Therefore, the two types of mammary epithelial cells we isolated had a normal proliferate capability and this response was similar to the results of Cifrian (Baumrucker et al., 1988).

Cytokeratin 18 is one of the most common members of the intermediate filament gene family and generally exists together with its filament partner keratin 8. It is expressed in single layer epithelial tissues of the body and is specific for epithelial cells. Immunocytochemical staining of goat mammary tissue showed that cytokeratin 18 was found in both epithelial and myoepithelial cells of goat mammary gland (Hurley et al., 1989). Bovine myoepithelial cells were negative to anti-cytokeratin 18 monoclonal antibody. In the study, the obtained bovine mammary epithelial cells which had stained positive for anti-cytokeratin 18, exhibited the same phenomenon as reported previously. The positive staining for cytokeratin-18 was a powerful result to prove the specific epithelium character of MECs.

Epithelial differentiation is also characterized by expression of milk proteins such as α-s1-casein and κ-casein, the production of milk fat rich in triglycerides, sources of energy and essential fatty acids (Gertler et al., 1982). Casein secretion is the hallmark of the bovine mammary epithelial cells (Beswick and Kennelly, 1998). The results from the study showed that the cells isolated had normal secretion function.

**CONCLUSION**

The study demonstrated that the establishment of these two bovine mammary epithelial cell lines from Chinese Holstein cattle exhibited normal function. And the isolated mammary epithelial cells have been used for investigation of gene function about milk fat and protein and lipometabolism mechanism study.

**ACKNOWLEDGEMENTS**

This research was supported by the National R. and D. Project of Transgenic Organizations of Ministry of Science and Technology of China (2008ZX08007-001, 2009ZX08009-1 676 and 2009ZX08007-005B) and National Natural Science Foundation of China (No. 31000991). Researchers thank the anonymous reviewers for their time and energy in reviewing this study. Chunyan Lu and Runjun Yang contributed equally to this research.

**REFERENCES**


