





Differences between cloned and noncloned cows during the first lactation

F. Montazer-Torbati, M. Boutinaud F. Jaffrezic, D. Lalöe C. Richard, H. Jammes, J.P. Renard

E. Devinoy

eve.devinoy@jouy.inra.fr

INRA UMR1313, GABI, Jouy en Josas, France







Introduction

- The high variability in bovine milk production depends both on genetic and environmental factors
- Whereas genomic selection has contributed to largely improve milk production, mechanisms underlying the influence of the environment are mostly unknown
 - They can be studied using cloned cows (obtained through nuclear transfer), presumed to have identical genomes









Experimental procedures

Prim'Holstein cows: Cloned cows (n=9) and non cloned cows (n=9)

• Produced at similar period of the year



Raised in similar conditions







Reproduction

- All cows were inseminated at the age of 1.5 year
 - Eight cloned and 7 non-cloned cows delivered
- Body condition score at calving was lower (ΔBSC= -0.6) for cloned cows









Milk production during 200 DIM

Milk yield was highly variable among animals but similar between the 2 groups
Milk protein and fat contents were less variable in cloned cows

AUGUST - 4 SEPTEMBER 2015

WARSAW. POLAND





Milk composition around 67 DIM

• Lower protein, fat and lactose contents were observed in cloned cows



Mammary biopsy analyses

- Around 67 DIM, biopsies were collected from a rear quarter of the udder
- Five biopsies in each group contained more than 60% of mammary epithelial tissue



Cloned cow



Non-cloned cow

• They were from a sub-selection of cows which had a milk production representative of that observed among cows in each group.







RNA and milk synthesis

Levels of transcripts involved in most casein, fat or lactose syntheses were highly variable and in the average lower in cloned cows but not different between cloned and non-cloned cows, (except CSN1S2)









Milk somatic cell counts (SCC) in cloned cows

- Higher milk SCC, as an average (p=0.1)
- Increased frequency of high SCC (p=0.09)

• Higher apoptotic rate (TUNEL)



During lactation, a slow renewal of mammary cells is observed (0,3% per 24h) Mammary specific DNA methylation profiles are likely to be maintained by DNMT1



DNA methylation- DNMT1 mRNA

 DNMT1 RNA levels were higher (p=0.05) in cloned cows *DNMT1* RNA levels were related to the apoptotic rates (TUNEL)









Conclusions

- Lactation and milk production in cloned cows exhibit slight differences as compared to those of non-cloned cows
 - Higher somatic cell counts, higher apoptotic rates in relation with modifications of global DNA methylation profiles may explain such differences









Thank you for your attention ! Acknowledgements

Collaborators:

- F. Montazer-Torbati, post doc
- M. Boutinaud, INRA UMR 1348, PEGASE
- N. Brun, INRA UMR 1313, GABI
- F. Jaffrezic, D. Lalöe , INRA UMR 1313, GABI
- H. Jammes, J.P. Renard, INRA UMR 1198, BDR
- M. Nguyen, post doc
- C. Richard, V. Hallé, INRA UMR 1198
- A. Neveu, Valérie Gelin, INRA UCEA 1298

Fundings:



- French « Agence Nationale pour la Recherche »: Project « Epigrani »
- APIS-GENE
- INRA, department PHASE



