













DGAT1 K232a polymorphism greatly affects mammary gland activity of milk component synthesis

Faucon, F.,

Rebours, E., Robert-Granie, C., Bernard, L., Menard, O., Miranda, G., Dhorne-Pollet, S., Bevilacqua, C., Hurtaud, C., Larroque, H., Gallard, Y., Leroux, C., Martin, P.

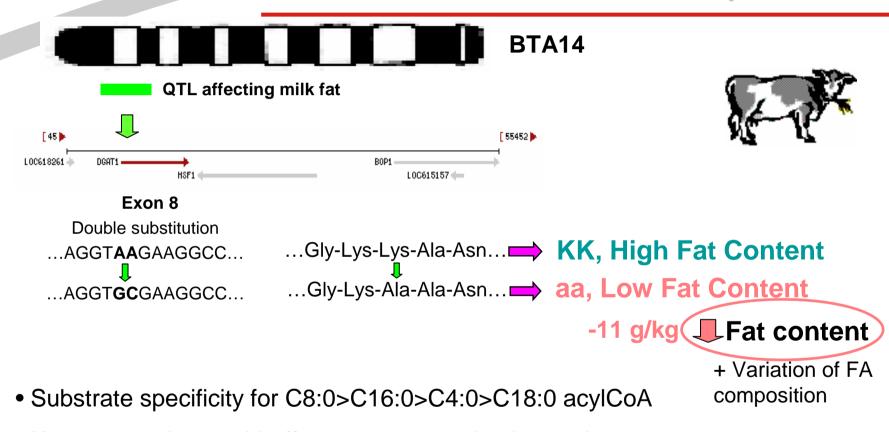
INRA : Unité Génomique et Physiologie de la Lactation Equipe Génomique Expressionnelle et Lait

felicie.faucon@inst-elevage.asso.fr





DGAT1 K232a mutation and consequences



• K232a mutation could affect enzyme maximal speed (Grisart et al., 2004) and specificity (Schennink et al., 2007)





DGAT1 issue

What are the molecular mechanisms underlying variation of milk fat content and fatty acid composition with the DGAT1 K232a mutation?





Variation of milk production traits

With the aa, low fat content genotype:

significant reduction of fat content and milk fat globule size

tendency of increased milk production and lactose yield

no difference for protein content and yield and fat yield





Variation in fatty acid composition

With the aa, low fat content genotype:

more saturated middle chain and unsaturated long chain fatty acids

less unsaturated middle chain fatty acids

=> Comparable to schennink et al., 2007 and 2008 results





Gene expression profiling: microarray analysis

22k CRB* slide

~ 9 973 genes

4x44k Agilent slide

~ 13 746 genes

"Super slide"



25 % of overlapping between the 2 slides

= 14 567 genes with Ingenuity annotation

~55 % of the bovine genome







Strategy



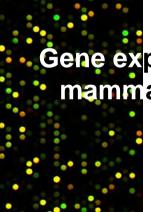
KK, high fat content



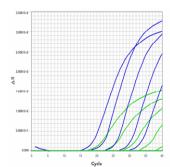
6 pairs (full or half sibs)
Pin-au-Haras
(Holstein x Normande)

aa, low fat content

UE Theix



Gene expression in mammary gland













Differential analysis, FDR<0,1, cutoff at 1,5

General microarray results

228 down-regulated genes with aa, low fat content

Cytoskeleton proteins

(actins, keratins, tropomyosin)

Post-translational modifications of the proteins

(LBP, B4GALT5)

Immune response

(complement, interleukins, interferon, chemokine, integrin)

Lipid metabolism

(LDL and chylomicron captation, catabolism)

197 up-regulated genes with aa, low fat content

Milk lipid biosynthesis and secretion

FA captation, FA, DG and PL synthesis

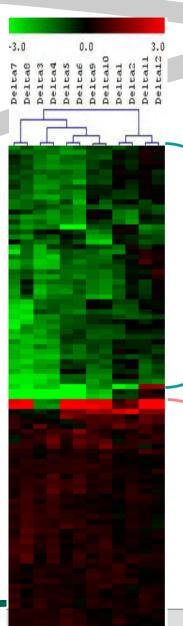
Lipids secretion

Regulation of lipid synthesis

Milk proteins (CSN1S1, CSN1S2, LALBA)







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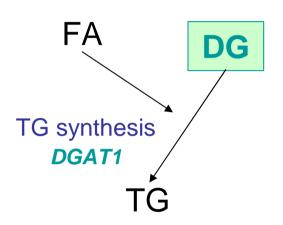
EAAP – Barcelona – 24th august 2009

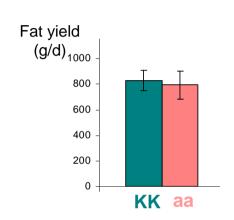




Decrease expression of DGAT1 gene but similar fat yield

aa, Low fat content KK, High fat content





DGAT1 catalyzes the production of TG (main lipid component in milk) from DG and FA.

With aa, Low fat content genotype: less efficient (Grisart et al., 2004) and DGAT1 is down regulated

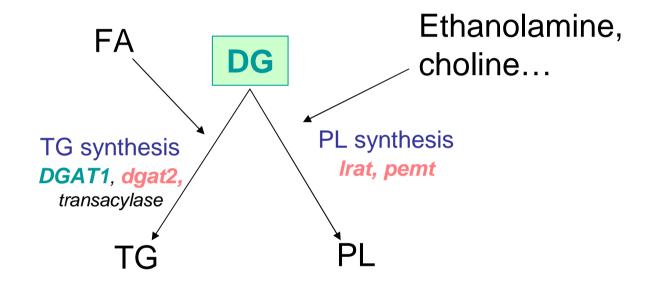
However, secreted fat yield is quite the same between genotypes

⇒ Alternative pathways to synthesize TG from DG ?





Differentially expressed genes



- + saturated middle chain FA
- + unsaturated long chain FA

Smaller milk fat globules

aa, Low fat content KK, High fat content





Conclusions

❖ Variation of milk FA content with K232a mutation can be explained by:

alteration of enzyme specificity of DGAT1(Schennink et al., 2007) alternative milk lipid biosynthesis pathways (TG, PL)

- ❖ Each milk component synthesis pathway seem affected by K232a mutation! Is this representative of an increased insulin sensitivity of the mammary tissue?
- Validation: genes redundancy between the 2 slides and qRT-PCR (in progress)
- functional annotation of some genes involved in the decrease of milk fat content.





Thanks for your attention!



