

**The genomic region responsible for
E. coli F4ab/ac susceptibility in pigs:
 Characterization and Application**

SABRE CUTTING EDGE GENOMICS FOR SUSTAINABLE ANIMAL BREEDING

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Acknowledgements

Denmark
 Mette Juul Jacobsen
 Steffen Kracht
 Bjarne Nielsen
 Susanna Cirera
 Peter Karlskov Mortensen
 Merete Fredholm

Sweden
 Inger Edfors-Lilja
 Leif Andersson

Switzerland
 Peter Vogeli
 David Joller

Spain (SABRE travelgrant)
 Gloria Esteso

UK
 Alan Archibald
 Pig team at Sanger Institute

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***E. coli* F4ac in the pig**

- 10 % of all piglets suffer from *E. coli* diarrhoea.
- 20% of infected piglets die
- 37% caused by *E. coli* O149,F4ac
- 170.000 piglets/year in Denmark
- Resistance is autosomal recessive (SSC13)

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Adhesion of *E. coli* F4ac

Adhesion

- Fimbriae type
- Susceptible cell

Colonisation

- Toxin production

Python *et al.* 2002

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Characterization

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Mapping of the F4ac-receptor

- F4ac receptor locus has firmly been linkage mapped between the SW207 and SW1876 (7 cM). (Python *et al.*, 2002; Jørgensen *et al.*, 2003).
- This region contains approx. 150 genes (comparative mapping).
- SNP in *MUC4* shows complete LD with the phenotype (*Xba*I polymorphism) (PCT application, Jørgensen *et al.*, 2004)
- Fine mapping of the *MUC4* flanking region
- Detect ancestral haplotype

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Fine mapping of the *MUC4* region

The 3 generation material consists of:

2 Boars 10 founders 8 sows

X

26 F1 and 200 F2

18 informative founder chromosomes (12 susceptible and 6 resistant)

SNP discovery in EST-data and by genomic re-sequencing in founders

Fine mapping of the *MUC4* region

Identified 177 polymorphic sites, mostly SNPs and a few small deletions/insertions

Mette Juul Jacobsen

SNP-typing in founders

- Large unique haplotype block on susceptible chromosomes.
- *MUC4* in the center
- Resistant haplotype pattern is complex
- Resistant is wildtype
- SNP-typing in additional Swiss founders
- Crucial to determine minimal shared haplotype in susceptible

Sequencing of *MUC4*-BAC Sanger Institute

Application

Dansk Svineproduktion

MUC4 marker in use by Danish Pig Production

Initially selection only on Yorkshire and Landrace boars

Selection of Yorkshire boars

- RR ~ 20%

Selection of Landrace boars

- SR ~ 10%
- RR ~ 2%

Resistens mod smågrieseidarré inddrages i avlsmålet

