The genetic background of clinical mastitis in Holstein-Friesian cattle

_J. Szyda^{1,2}, M. Mielczarek^{1,2}, M. Frąszczak¹, G. Minozzi^{3,4}, R. Giannico⁴, J L. Williams⁵, K. Wojdak-Maksymiec⁶

¹Biostatistics Group, Department of Genetics, Wroclaw University of Environmental and Life Sciences, Wroclaw, Poland; ²National Research Institute of Animal Production, Wroclaw, Poland; ³Department of Veterinary Medicine, University of Milan, Milan, Italy; ⁴Fondazione Parco Tecnologico Padano, Lodi, Italy; ⁵Davies Research Centre, School of Animal and Veterinary Sciences, University of Adelaide, Roseworthy, Australia; ⁶Department of Animal Genetics, West Pomeranian University of Technology, Szczecin, Poland.

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Objectives

Characterization of links between single nucleotide polymorphisms (SNPs), copy number variants (CNVs) and the incidence of clinical mastitis (CM)

Conclusions

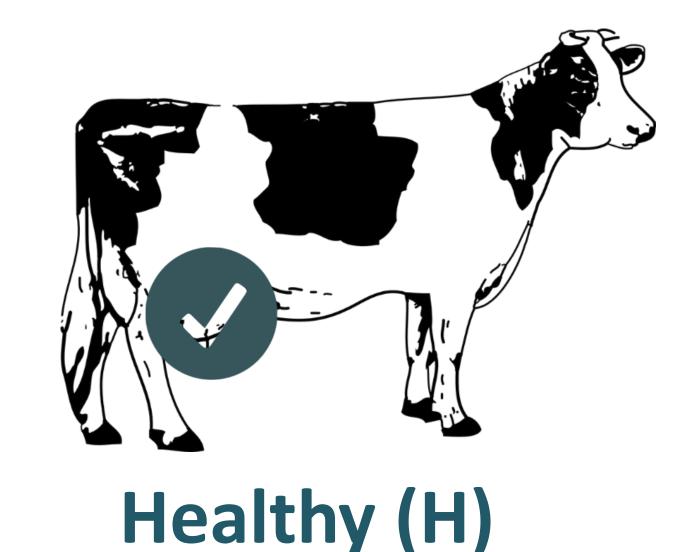
- CNVs play an important role in the susceptibility to CM
- Identified genes are involved in immune response

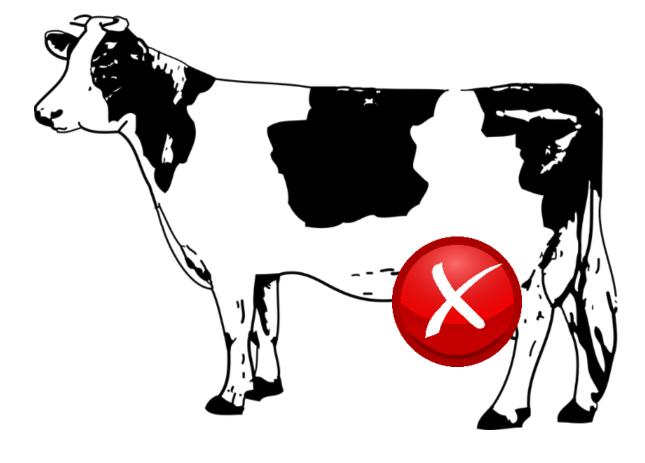
Presenting author: magda.mielczarek@upwr.edu.pl, magdam@illinois.edu

- Deletions more severe consequences on reducing resistance against CM, than...
- Duplications on increasing resistance to CM

Data set

- 32 Polish Holstein
 Friesian cows
- 16 paternal half-sibs
- Illumina HiSeq 2000
- Coverage 5x 17x
- NCBI BioProject
 PRJNA359667





Sick (S)

Results

SNPs

Introns of genes with differences in SNP genotypes between sick and healthy cows

RefSeq coding ring finger protein 122 (10/16 half-sibs)

cell viability and immune response

MET proto-oncogene receptor tyrosine kinase (9/16 half-sibs)

 processes related to inflammation, cancerogenesis

WRN gene (8/16 half-sibs)

 premature aging in humans, increased susceptibility to infection

CNV DELETIONS

Exons deleted in a sick cow, but present in its healthy half-sib exhibits potential causal influence CM (at least 7 half-sibs pairs)

FOXL2: forkhead box L2

role in inflammation

SSFA2: sperm specific antigen 2

- associated with SCS = indicator of CM
- overlapps with QTL for bovine immunoglobulin G

ADORA2A: adenosine A2a receptor

- modulating tissue response to inflammation
- in mice highly expressed in mammary gland

DGVa: Hou et al. 2011

NDUFS6: NADH: ubiquinone oxidoreductase subunit S6

• QTL for SCS = CM indicator DGVa: Boussaha *et al.* 2015

TXNRD2: thioredoxin reductase 2

 candidate for influencing susceptibility to S.aureus

DGVa: Hou et al. 2011

APP: amyloid beta precursor

- bactericidal and antifungal activities in human
 - molecular markers for SCM in ruminants

OTUD3: deubiquitinase 3

 associated with inflammatory bowel disease in humans

DGVa: Keel et al. 2017

Methods

Alignment to the reference genome

Post-alignment processes

SNP detection

CNV detection

Half-sibs SNP comparison

Half-sibs CNV comparison

Functional annotation

Enrichmentanalysis

Acknowledgements









