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A genome-wide association scan for loci affecting osteochondrosis in German Warmblood horses

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Introduction

Osteochondrosis - Definition

- <u>Common</u> joint disorder in young domestic animals and humans
- Focal disturbance of enchondral ossification
- Osteochondrosis dissecans (OCD): Presence of an osteochondral fragment (chip)
- Predilection sites*
 - Caud. aspect of the prox. articular surface of the humerus
 - Medial aspect of the condyle of the humerus
 - Lateral trochlear ridge of the femur
 - Medial condyle of the femur
 - Intermediate ridge of the distal tibia
 - Lateral trochlear ridge of talus
 - Dorsal aspect of the distal metacarpus and metatarsus
- Multifactorial: <u>Hereditiy</u>, rapid growth, anatomic conformation, trauma, dietary imbalances...





Osteochondrosis – Prevalence and Heritability

Author	Breed	Joint	Prevalence	Heritability
Winter <i>et al.</i> (1996)	German Riding Horse		11 %	0.06-0.07
Willms et al. (1999)	German Riding Horse	Toe, hock	5 %	0.45-0.64
Pieramati <i>et al.</i> (2003)	Maremmanos	Stifle, hock, fetlock	16.6 %	0.19-0.58
Stock <i>et al</i> . (2005)	Hanoverian Warmblood	Fetlock Hock	20.8 % 9.6 %	0.08-0.14 0.19



Animals and phenotypes

- X-ray protocols of 1,162 stallions presented for licensing in the years 2005 to 2008 in four German breeding organizations: Hanoverian Society, Holsteiner Verband, Oldenburger Horse Breeders Society, Trakehner Verband
- Osteochondrotic findings (OC and OCD) were recorded for stifle, hock, fetlock and toe and coded as a binary trait (1 = at least one finding in the respective localization vs. 0 = no finding)

 \rightarrow Assignment to predilection sites?



 \rightarrow The localization ,,toe" includes the fetlock joint and is not a common definition!



Influencing factors

- Logistic regression (procedure glm in R), logit-link function
- Significant covariates and fixed effects were included in the respective GWAS

	Age	Height	Year	Principal consultant	Breeding Organisation
OCD _{TOE}					
OCD _{TOE-FRONT}					
OCD _{TOE-REAR}					
OCD _{FET}					
OCD _{FET-FRONT}					
OCD _{FET-REAR}					
OCD _{HOCK}					



GWAS

- Genotyping: 943 stallions typed with Illumina EquineSNP50 Bead Chip featuring 54,602 SNPs
- Preprocessing: Individual callrate ≥ 0.95 , SNP callrate ≥ 0.9 , MAF ≥ 0.05 , $p_{HWE} \leq 10^{-9}$, filtering for individuals with a pairwise IBS > 0.95

 \rightarrow 916 individuals

→ 44,410 SNPs

- Principal components to correct for population structure*
- Correction of p-values using genomic control







GWAs results for OCD_{TOE}

- Phenotype: at least one OCD-finding in one of the toes including fetlock
- Fixed effects: Year of licensing



Horse Chromosome



GWAs results for $\mathsf{OCD}_{\mathsf{FET}}$

- Phenotype: at least one OCD-finding in one of the fetlock joints
- Fixed effects: Year of licensing and breeding organization
- Estimated inflation factor λ : 1.05





GWAs results for OCD_{HOCK}

- Phenotype: at least one OCD-finding in one of the hock joints
- Covariate: Age
- Estimated inflation factor λ : 1.05





Results and Discussion

A major QTL on ECA20?





- The use of X-ray protocols as a source of phenotypic information has some disadvantages making it necessary to define more general phenotypic classes.
- One significant and several suggestive QTL affecting the liablity to OCD have been identified within the German Warmblood population.
- We hypothesize that a major QTL resides on ECA20 that generally affects the liability to OCD in the distal limb.



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