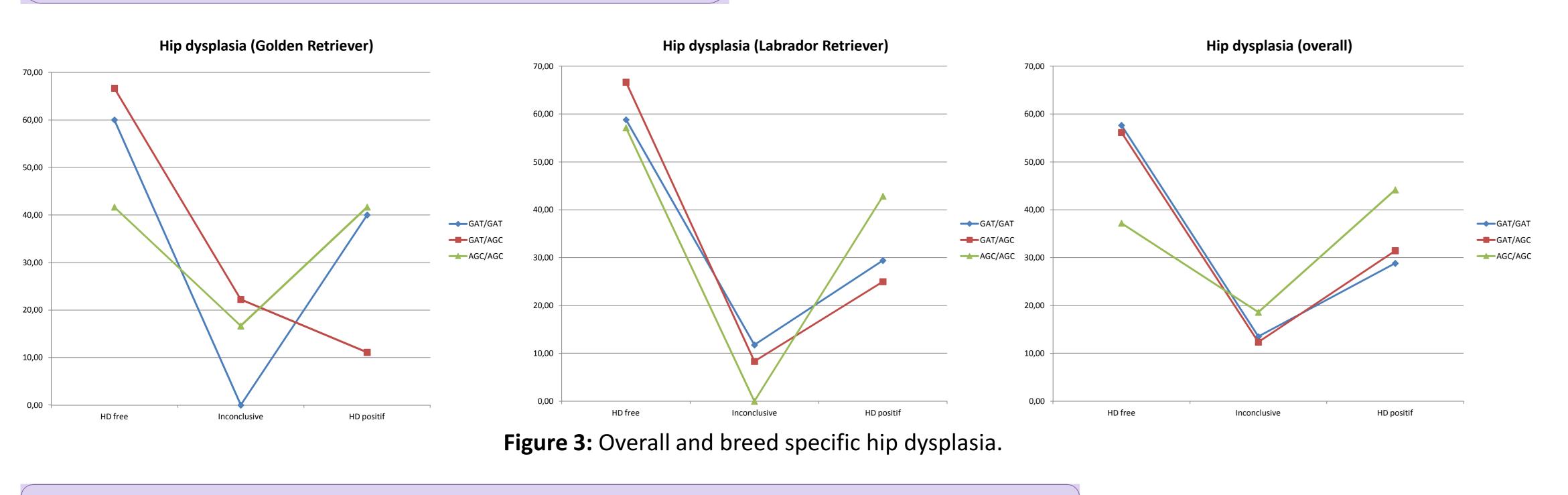


Introduction

Hip dysplasia is a highly prevalent complex disorder. Despite decades of selection against this disease based on radiographic methods, prevalence has hardly been Recently, focus reduced. has been redirected at the underlying genetics. This has led to the discovery of an association between the AGC haplotype in the FBN2 gene and hip dysplasia, based on an American population of dogs (Friedenberg et al., 2011). Aim of this study was to investigate the prevalence of this mutation in a Dutch and Belgian dog population and to confirm the association of the mutations with phenotypical hip dysplasia.



Results

271 dogs of 50 different breeds were genotyped (Table 1). X-rays of 243 dogs were available. There were no significant age differences between groups (FBN2 and HD, Figure 2).

All the radiographic assessments were highly correlated with each other (p < 0,001). The AGC haplotype was only significantly correlated with the general assessment (p< 0,05). Our model contained two significant coefficients (namely age and FBN2, p< 0,05) that explained 14 % of the variation.

Reference: Friendenberg S.G., et al. (2011). Evaluation of a fibrillin 2 gene haplotype associated with hip dysplasia and incipient osteoarthritis in dogs. AJVR 72 (4), p. 530 – 540.





FACULTEIT FARMACEUTISCHE WETENSCHAPPEN

Relation between the FBN2 haplotype and phenotypical hip dysplasia Bart J.G. Broeckx¹, Frank Coopman², Geert Verhoeven³, Tim Bosmans⁴, Ann C.C. Criel⁵, Ingrid Gielen³, Jimmy H. Saunders³, Sandra Soetaert¹, Henri Van Bree³, Wim Van Haeringen⁶,

Filip Van Nieuwerburgh¹, Bernadette Van Ryssen³, Dieter Deforce¹

¹ Laboratory for Pharmaceutical Biotechnology, Faculty of Pharmaceutical Sciences, Ghent University, Ghent, Belgium ² Faculty of Applied Bio-engineering, University College Ghent, Ghent, Belgium ³ Department of Medical Imaging, Faculty of Veterinary Medicine, Ghent University, Merelbeke, Belgium ⁴ Department of Medicine and Clinical Biology of Small Animals, Faculty of Veterinary Medicine, Ghent University, Merelbeke, Belgium

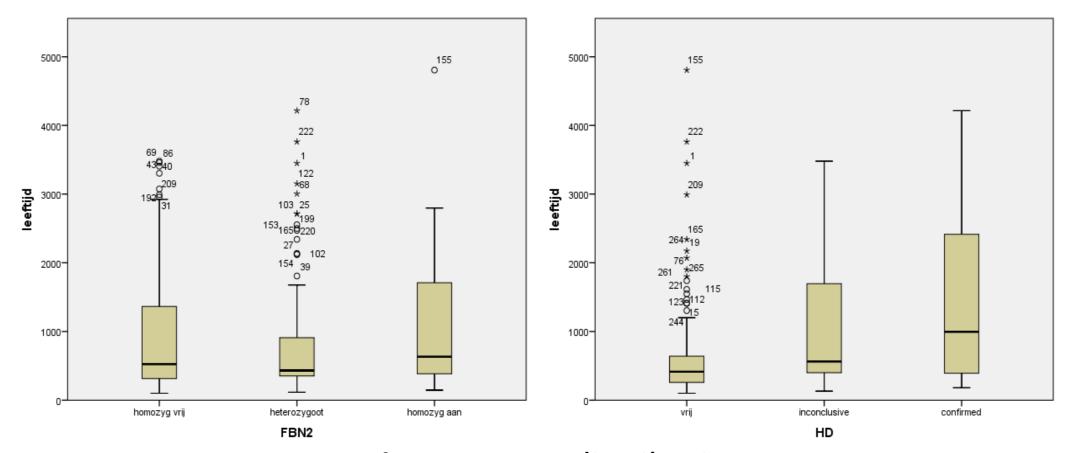
⁵ NCSA, Brussels, Belgium

⁶ Dr. Van Haeringen Laboratorium b.v., Wageningen, The Netherlands



Table 1: General and breed specific prevalence and allele frequencies.

					Allele frequency	
	N	GAT/GAT (%)	GAT/AGC (%)	AGC/AGC (%)	GAT	AGC
All dogs	271	46,5	36,2	17,3	64,6	35,4
German Shepherd	54	50	44,4	5,6	72,2	27,8
Golden Retriever	38	18,4	50	31,6	43,4	56,6
Labrador Retriever	39	48,7	33,3	17,9	65,4	34,6



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Figure 1:

Phenotypical categories (general assessment), from left to right: healthy, inconclusive, dysplastic.

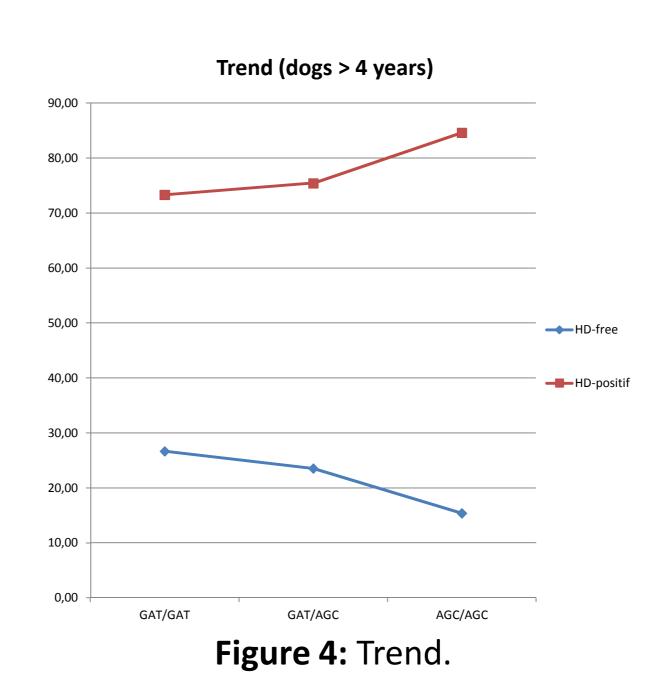


Figure 2: Age distribution.

Materials and Methods Blood samples and X-rays (standard hip extended) were collected from dogs of different breeds. Birth date and date of x-ray were noted. Based on the standard hip x-ray, hip status was assessed by consensus (BB, FC) four different radiographic based on categories:

- (Yes/No/Inconclusive).

The dogs were genotyped with KASPar. The prevalence of the haplotype was calculated in the general population and in 3 breeds. The an association between presence of phenotype and genotype was investigated. Statistical analysis was performed with SPSS Statistics 20.

Conclusions

Although significantly correlated, variation is only partly explained by FBN2 and age. Breed specific differences seem to be present (Figure 3). There is a trend towards worse hip conformation for the homozygous AGC dogs (Figure 4). A more refined phenotyping, a larger amount of animals tested and the inclusion of more environmental factors will be necessary to refine the model. These improvements are being tested at the moment. In general, we reach the same conclusions as presented by Friedenberg et al. (2011) but on a larger database and over more breeds.





- Hip dysplasia (yes/no/inconclusive) (Figure 1) - Deg. Joint Dis. (No/Mild/Moderate/Severe), - Morgan line (Yes/No/Inconclusive) - Circumferential femoral head osteophyte