# 

### BLUP Animal Model for prediction of breeding values for hip dysplasia in Norwegian dog breeds





#### 

# Canine Hip Dysplasia (HD)

- HD is a malformation of the coxofemoral joint a developmental orthopedic disease in many dog breeds
- >Many studies have shown that it is partly genetic controlled, with a polygenic background
- >Mild forms seldom affect the functional health of the dog
- >In more severe form HD might cause crippling lameness

<section-header><section-header><list-item><list-item><list-item><list-item>













# A Radiographic screening for HD has been performed in many breeds in Norway for more than 30 years > Official HD result is obligatory for dogs used for breeding in 70 breeds (for most of the breeds since 20 years) > For most of breeds 40-60% of the dogs has official HD result > Selection has until 2008 been based on mass selection

# Data HD-scores collected between 1987 and 2008 on 133.821 dogs from 32 breeds was available from the Norwegian Kennel Club (NKC) No. of records per breeds varied from: 739 for Hamiltonstöver and 786 for Chow Chow to 14.464 for Golden Retriever and 23.183 for German Shepherd

Frequencies of HD-score (extreme breeds)						
(0)		HD-score (%)				
Breed	# obs,	А	В	С	D	E
Vorster	2.738	91,2	5,0	2,7	0,9	0,2
Tervuren	2.648	91,1	3,3	3,3	1,6	0,7
Flat Coated Retriever	7.020	89,9	4,1	3,9	1,5	0,6
Chow Chow	822	74,9	5,0	9,6	5,1	5,4
German Shepherd	23.093	67,0	6,9	12,4	8,1	5,6
Newfoundland	2.584	59,9	3,5	13,3	14,1	9,2
						13



#### 

# Heritabilities (h<sup>2</sup>) and litter effect (c<sup>2</sup>)

For breeds with >1.000 records:

h<sup>2</sup> varied from: 0,08 for Newfoundland to 0,49 for Saint Bernhard with and average of ~0,30

c<sup>2</sup> varied from: 0,00 for Border Collie to 0,11 for Saint Bernhard with an average of ~0,05



# ACILITY OF AGRICULTURAL SCIENCES

# Model validation

Model validation has been conducted by prediction of EBV's based on all data ( $EBV_{All}$ ) and on a reduced data set, where HD records from dogs born in 2004 and later was deleted ( $EBV_{Red}$ )

The regression of EBV<sub>All</sub> on EBV<sub>Red</sub> has an expectation of 1,0

Dogs with deleted HD record in the reduced data will get  ${\sf EBV}_{\sf Red}$  based on pedigree information. The regression of  ${\sf EBV}_{\sf All}$  on  ${\sf EBV}_{\sf Red}$  for this group of dogs will therefore be at test of bias in pedigree indexes

For 3 breeds (Chow Chow, Irish Setter and Doberman) the regression was substantial below 1,0 (0,79, 0,70 and 0,71)



# 

# Summing up

- >The estimated heritability varies between breeds, but most estimates are between 0,2 and 0,3
- ▶BLUP AM was introduced by NKC in 2008
- ≻EBV's for 32 breeds are computed 4 times per year
- ≻Published on NKC's DogWeb
- >Breeding advice: The mean value of the combination should be above breed average (100)