

Boerboel Genetic Evaluation

SA Stud Book
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WELTEVREDEN PARK
ROODEPOORT
1709



Kennel Number: 639381



Estimated Breeding Values (EBVs)

Litter size and conformation traits are influenced by many genes, with every gene making a small contribution to the expression of the trait. These traits are, however, also influenced by physiological and physical environmental factors, for example the age and sex of the animal, the feeding and management the animal received and year x season effects. It is therefore not possible to know what the animal's genetic potential for the trait is by merely using the measurement.

With BLUP (*Best Linear Unbiased Prediction*) methodology an animal's genetic potential is predicted by making use of the genetic parameters (heritabilities and genetic correlations) of the breed, pedigree information, as well as the measurements of the animal as deviation from the average of the animals that were measured with the animal. Adjustments are made for differences in sex, age, classifier, management and year x season effects to ensure that fair comparisons are made amongst the animals. These predictions are called **EBVs (Estimated Breeding Values)** which indicates the animal's genetic potential for the specific trait and predicts the type of progeny that the animal will breed.

Pedigree information determines the genetic level at which the animal enters the genetic evaluation, it links the information of all relatives together and it is used to determine the genetic ranking of the breed. For young animals that were not yet measured, mid-parent EBVs are estimated, which are the average of the EBVs of the sire and dam. These breeding values are still inaccurate and can change a lot when more information (measurements) is included in subsequent genetic evaluations.

Breeding values therefore enables a breeder to compare his/her animals genetically to one another, as well as to all other animals of the breed. This Report gives a summary of the genetic levels of the active males and females of the relevant breeder in comparison to that of the breed for linearly classified traits and litter size. The genetic change that occurred over time can also easily be seen.

Breeding Value Indices (EBVi)

For easy interpretation, EBVs are standardised and expressed relative to the average (100) of the active animals of the breed, using 12 as standard deviation (SD). This means that an EBV of 100 is equal to the active average of the breed, and more or less 50% of the animals will have EBVi's of between 90 and 110, which indicates an average animal. Only 0.5% of the EBVi's will be lower than 64 and 0.5% will be higher than 136. Higher EBVi's indicate better genetic potential for the trait.

Accuracy

EBVs are published with an **accuracy value (%)**, which indicates how much info was available for the estimation of the EBV and with how much trust it can therefore be applied. Factors that influence the accuracy include heritability, completeness and correctness of pedigree information, number of own and measured relatives and the size of the group in which the animal was measured. When an accuracy of 85% and more is reached, the EBV should not change significantly with the inclusion of more information in subsequent genetic evaluations.

Kennel and Breed Averages: The different traits' averages of the breeder's active animals are compared to those of the breed in the Kennel and Breed Average Graphs and Tables. The breeder can therefore easily see for which traits the animals are above or below the average of the breed.

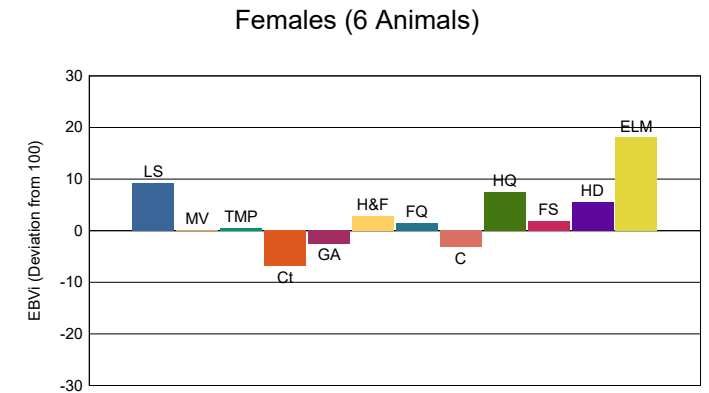
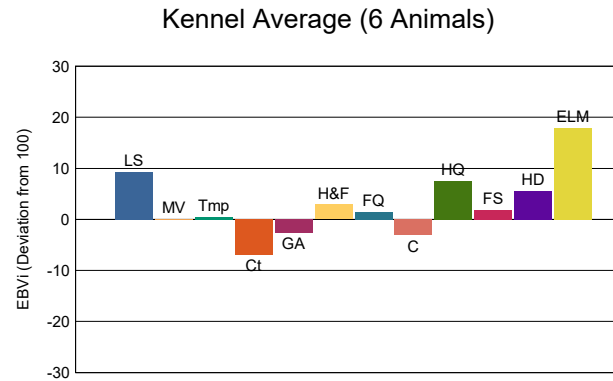
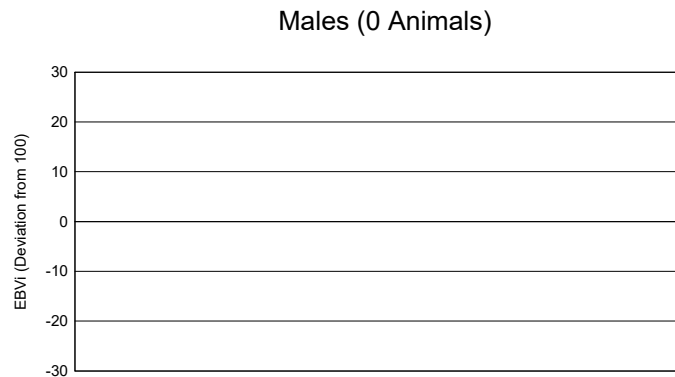
Genetic Trends: The Genetic Trends indicate the genetic change that occurred over time in the animals bred by the breeder compared to that of the breed. These are calculated as the average of the breeding values of all measured animals bred by the breeder, per year of birth.

Inbreeding: Inbreeding is the mating of related animals and is widely viewed as being harmful. The danger of inbreeding lies there in that it can gradually decrease fertility and survivability – a phenomenon known as inbreeding depression. It can cause lower conception rates, more abortions, more stillborn and weak offspring and a higher susceptibility to diseases. Indications in dogs include a low litter size as well as problems relative to fertility, such as invisible heat, behavioural problems when mating and low sperm quality. Overall, reduced litter size most likely reflects inbreeding depression in several traits, both physiological and behavioural, as well as a high frequency of non-viable embryos due to lethal alleles. Generally, the effects of inbreeding can become noticeable at a value of 6.25%, therefore it is generally recommended that individuals should have inbreeding coefficients of less than 6.25%. Breeders should therefore avoid matings causing inbreeding of more than 6.25% in the progeny. The theoretical maximum inbreeding coefficient is 50%. However, it should be noted that not all inbred animals show signs of inbreeding depression.

The data used in this evaluation was obtained from LOGIX from linear classifications that were done by the Breeder Society. All possible attempts have been made to present the information correctly. SA Stud Book does not accept responsibility for the use and interpretation of the information presented on animals in this publication.

Genetic Levels

Kennel Average relative to Breed Average, where Breed Average is indicated as 0



Average Breeding Value Indices (EBVi) where Breed Average = 100

Trait	All Animals	Males	Females
Litter size (LS)	109		109
Movement (Mv)	100		100
Temperament (Tmp)	101		101
Coat (Ct)	93		93
General Appearance	97		97
Head & Face (H&F)	103		103
Front Quarter (FQ)	101		101
Centre (C)	97		97
Hind Quarter (HQ)	107		107
Final Score (FS)	102		102
Hip Health (HH)	106		106
Elbow Health (ELM)	118		118

Inbreeding

Notes

Inbreeding primarily leads to decreased fertility, low survivability of pups and low disease resistance.

Inbreeding coefficients are calculated for all animals of the breed and indicates the probability that two genes of a pair are the same due to a common ancestor in the pedigree of the animal.

The adverse effects of inbreeding can be observed from 6.25% onwards.

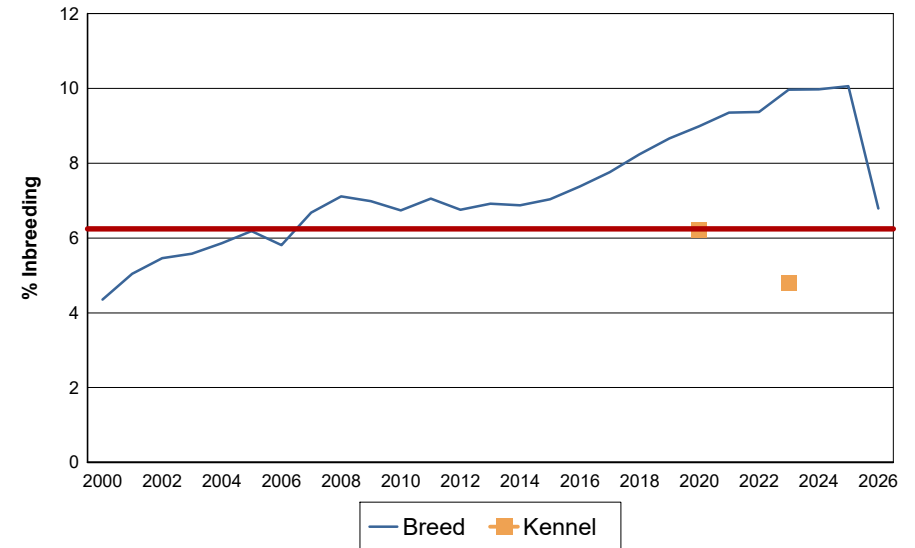
It is therefore highly recommended to record pedigree information as accurately. Inbreeding is dependent on pedigree information. Conventional methods for the calculation of inbreeding coefficients usually assume that unknown parents are unrelated, which is usually far from the truth. Inbreeding coefficients are therefore conservative indicators of the level of inbreeding.

Incomplete or incorrect pedigree information can be reported to SA Stud Book via the Breed Society.

Inbreeding Levels for Specific Matings

Progeny with Parents :	25%
Full Brother x Full Sister :	25%
Half Brother x Half Sister :	12.5%
One Common Grandparent :	6.25%
One Common Great-Grandparent :	3.125%

Rate of Inbreeding



Inbreeding Levels

	All Animals		Males		Females		Sires		Dams	
	Kennel	Breed	Kennel	Breed	Kennel	Breed	Kennel	Breed	Kennel	Breed
Number	6	49007	0	25141	6	23857	2	3371	2	6287
Average Inbreeding	5.42	8.49		8.46	5.42	8.53	4.31	8.01	8.57	8.00
Number inbred	6	48727		24990	6	23728	2	24990	2	23728
Number >6.25% Inbr.	1	30600		15537	1	15058		15537	2	15058

Estimated Breeding Value Indices (EBVi's)

Estimated Breeding Value Indices (EBVi's)

Females

Inbreeding >6.25% is high

< 80

>= 100

>= 120

Animal Information

Inbreed

Health

Fertility

General

Appearance

Progeny :

Appraisal

Health

Progeny with litters

Name	BYr	Sire Dam	%	Hips Acc.	Elbows Acc.	Litter Size Acc.	Move-ment	Tempe-rament	Coat	General Appear.	Head & Face	Front Quarter	Centre	Hind Quarter	Final Score	Progeny Progeny	Litters Litters	Progeny Kennels	Hips	Elbows	Progeny with litters
AFRIKA RAINSTORM	2021	MES RUBENSTEIN AFRIKA LAGITTA	8.52	116 33	126 50	117 40	110 54	111 34	99 66	102 53	102 66	105 55	102 56	115 58	107 62	10	1				
RAMOS STORM	2023	KADIMA MAVERICK JUNKER AFRIKA RAINSTORM	4.80	103 5	117 24	108 14	98 29	99 5	92 41	97 29	103 40	101 30	96 32	106 34	101 37						
RAMOS ARIEL	2023	KADIMA MAVERICK JUNKER AFRIKA RAINSTORM	4.80	103 5	117 24	108 14	98 29	99 5	92 41	97 29	103 40	101 30	96 32	106 34	101 37						
RAMOS BELA	2023	KADIMA MAVERICK JUNKER AFRIKA RAINSTORM	4.80	103 5	117 24	108 14	98 29	99 5	92 41	97 29	103 40	101 30	96 32	106 34	101 37						
RAMOS ANNABELLE	2023	KADIMA MAVERICK JUNKER AFRIKA RAINSTORM	4.80	103 5	117 24	108 14	98 29	99 5	92 41	97 29	103 40	101 30	96 32	106 34	101 37						
RAMOS SUNSHINE	2023	KADIMA MAVERICK JUNKER AFRIKA RAINSTORM	4.80	103 5	117 24	108 14	98 29	99 5	92 41	97 29	103 40	101 30	96 32	106 34	101 37						

Estimated Breeding Value Indices (EBVi's)

Sires

Inbreeding >6.25% is high

< 80

>= 100

>= 120

Sire Information			Inbreed	Health		Fertility	General			Appearance						Progeny :		Appraisal	Health		Progeny with litters
Name	BYr	Sire Dam	%	Hips Acc.	Elbows Acc.	Litter Size Acc.	Move-ment	Tempe-rament	Coat	General Appear.	Head & Face	Front Quarter	Centre	Hind Quarter	Final Score	Progeny Litters	Progeny Kennels	Hips	Elbows		
MES RUBENSTEIN		GROENBERG BBD RAMBO	4.08	111	59	133	122	134	109	99	101	114	104	128	116	314	52	101	21	20	
	2016	MES LIZE		58	67	73	86	71	92	86	92	87	88	89	90		83				
KADIMA MAVERICK JUNKER		MAXIPAL TROMPIE	4.53	91	108	99	87	86	85	92	105	96	89	97	94	11	2				
	2018	BREKER KALASHNIKOV		14	35	22	49	23	64	49	63	50	52	55	59						

Estimated Breeding Value Indices (EBVi's)

Dams

Inbreeding >6.25% is high

< 80

>= 100

>= 120

Animal Information			Inbreed	Health		Fertility	General			Appearance						Progeny :		Appraisal	Health		Progeny with litters
Name	BYr	Sire Dam	%	Hips Acc.	Elbows Acc.	Litter Size Acc.	Move- ment	Tempe- r ament	Coat	General Appear.	Head & Face	Front Quarter	Centre	Hind Quarter	Final Score	Progeny Litters	Progeny Kennels		Hips	Elbows	
AFRIKA LAGITTA		CABARET BALBOA	8.63	108	138	83	94	92	85	106	99	100	99	102	97	9	2	2	1	1	3
	2019	AFRIKA SAKABEKKIE		5	23	37	56	28	72	55	71	56	59	61	66						
AFRIKA RAINSTORM		MES RUBENSTEIN	8.52	116	126	117	110	111	99	102	102	105	102	115	107	10	1				
	2021	AFRIKA LAGITTA		33	50	40	54	34	66	53	66	55	56	58	62						

