STAMBOOM
PEDIGREE

RAAD VAN BEHEER
DUTCH KENNEL CLUB

Naam hond / Name dog	INSABA ECLIPSE
Ras – Variëteit / Breed – Type	Rhodesian Ridgeback
FCI nr	146
Geslacht / Sex	Teef

Stamboeknummer / Pedigree nr	NHSB 3226261
Chipnummer / Microchip	528140000820630
Geboortedatum / Date of birth	16-2-2021
Kleur / Colour	TARWEKLEURIG

Opmerkingen / Remarks
•

Fokker / Breeder	E. Havinga
Kennelnaam / Kennel name	INSABA
Kenmerk / Litter nr	8290-2020-ne

Vader 1
ARESVUMA CHAM'MBALI-MUSANGA
NHSB 3096208 IMP (RU)
981020009174717

- 3 HAWKINSARA BOO FOR ARESVUMA
  RKF 3859025
  - 7 ZURITAMI YADI
    VDH 05/171977
  - 8 CLACHAN SHASHI
    VDH 13/17U3098
- 4 ASADI FUADI AZALI UCHANGA
  RKF 3411223
  - 9 LOBENGULA SAFARI GAHIJI
    RKF 2610374
  - 10 OGBONNA CAPTAIN
    RKF 2756420

Moeder 2
INSABA DARK STORM
NHSB 3114318
528140000724027

- 5 KISANGANI HIGH FIVE SYDNEY
  VDH 12/1098829
  - 11 ELANGENI BRAVEHEARTRULES
    ANKC 2100264292
  - 12 KISANGANI ENJENJE N'TOMBI
    VDH 06/1094936
- 6 INSABA BUNIWA
  NHSB 2959306
  - 13 CHIWAMBO BAHA DE KIUNGWANA
    LOF6 2341
  - 14 FAIRRAY INSABA
    NHSB 2804518 IMP (IT)

DNA CERTIFIED PARENTAGE

Directeur, Rony Doedijns

Afgiftedatum / Issue date

16-4-2021

Bij deze stamboom hoort een document met resultaten van mogelijke onderzoeken, titels en wedstrijden. This pedigree is accompanied by a document with results of possible health screenings, titles and competitions

RAAD VAN BEHEER
HOUDEN VAN HONDEN

CERTIFICAAT

Naam	Insaba Eclipse
Ras	Rhodesian Ridgeback
Stamboeknr	3226261
Identificatie nr	528140000820630

Het onderstaande onderzoek is uitgevoerd conform het Raad van Beheer onderzoeksreglement.

Onderzoek	22-08-2022
	Heupdysplasie
	2470-2022-hd

Het (voor)onderzoek is uitgevoerd door:

Dierenarts	Dierenkliniek Hellendoorn Nijverdal
	H. ten Kate
	Ommerweg 54
	7447 RG Hellendoorn

Beoordeeld	22-08-2022	
Onderzoekresultaat	Beoordeling op onderdelen:	
	Botafwijking:	0
	Norbergwaarde:	32.5
	Aansluiting:	
	Vorm:	
	Eindbeoordeling:	HD A
		Volgens Internationale FCI norm

RAAD VAN BEHEER
HOUDEN VAN HONDEN

CERTIFICAAT

Naam	Insaba Eclipse
Ras	Rhodesian Ridgeback
Stamboeknr.	3226261
Identificatie nr.	528140000820630

Onderzoek

Het onderstaande onderzoek is uitgevoerd conform het Raad van Beheer onderzoeksreglement.
22-08-2022

Elleboogdysplasie

1454-2022-ed

Dierenarts

Het (voor)onderzoek is uitgevoerd door:
Dierenkliniek Hellendoorn Nijverdal

H. ten Kate

Ommerweg 54

7447 RG Hellendoorn

Beoordeeld 22-08-2022

Onderzoekresultaat

Beoordeling op onderdelen:

	Links	Rechts
Artrose:	Vrij	Vrij
Diagnose:		
OCD	Vrij	Vrij
LPC	Vrij	Vrij
LPA	Vrij	Vrij
Inc	Vrij	Vrij
OV.	Vrij	Vrij
Eindbeoordeling:	VRIJ	Nederlandse norm
	FREE	International Elbow group

RAAD VAN BEHEER
HOUDEN VAN HONDEN

CERTIFICAAT

LABOKLIN
LABORATORIUM VOOR KLINISCHE DIAGNOSTIEK N.V.

Industriestraat 29 . 6433JW Hoensbroek

Dierenkliniek
Hellendoorn-Nijverdal
Ommerweg 54
7447 RG Hellendoorn
Nederland

Report
No.: 2208-N-10000
Date of arrival: 23-08-2022
Date of report: 26-08-2022

Patient identification:	Dog female	* 16.02.21
	Rhodesian Ridgeback	
Owner / Animal-ID:	Havinga, Elisa	
Type of sample:	EDTA	
Date sample was taken:	22-08-2022	

Sampling:

The following impartial person (veterinarian, breed warden, or similar) signed the form for the sampling and identity check of the animal:

H. ten Kate

Name:	**Insaba Eclipse**
Stud book no.:	**3226261**
Chip no.:	**528140000820630**
Tattoo no.:	**---**

Degenerative Myelopathy - PCR

Result: Genotype N/N (exon 2)

Interpretation: The examined animal is homozygous for the wildtype-allele. It does not carry the high-risk factor for DM in exon 2 of the SOD1-gene.

Trait of inheritance: autosomal-recessive

Please note: In the Bernese Mountain Dog breed the mutation in exon 1 of the SOD1-gene also occurs in correlation with DM.

sample ID: 2208-N-10000

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Hemophilia B (Factor IX) - PCR

Result: Genotype female X(N)/X(N), male X(N)/Y

Interpretation: The examined animal is homozygous for the wildtype-allele. It does not carry the causative mutation for Hemophilia B in the FIX-gene.

Trait of inheritance: X chromosomal-recessive

Scientific studies found correlation between the mutation and symptoms of the disease in the following breeds: Rhodesian Ridgeback

Juvenile Myoclonic Epilepsy (JME)

Result: Genotype N/N

Interpretation: The examined animal is homozygous for the wildtype-allele. It does not carry the causative mutation for JME in the DIRAS1-gene.

Trait of inheritance: autosomal-recessive

Scientific studies found correlation between the mutation and symptoms of the disease in the following breeds: Rhodesian Ridgeback

D-locus D1 (dilution)

Result for d1: Genotype N/N (before D/D)

Interpretation: No d1-allele was found for this sample.

The overall genotype for the D-locus-complex can only be deduced if all known variants on the D-locus (d1, d2 and d3) are analysed. Some of these alleles only exist in specific breeds.

Please note: The nomenclature of the results has been changed due to harmonizing efforts for genetic tests.

B-locus (brown, chocolate, liver(nose))

This genetic analysis of the B-locus includes the three variants bd, bc and bs described for all breeds so far, as well as the corresponding wildtypes as allele N.

sample ID: 2208-N-10000

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Variant bd

Result for bd: Genotype N/N (before B/B)

Interpretation: No bd-allele was found for this sample.

Variant bc

Result for bc: Genotype N/N (before B/B)

Interpretation: No bc-allele was found for this sample.

Variant bs

Result for bs: Genotype N/N (before B/B)

Interpretation: No bs-allele was found for this sample.

When one of the variants is found homozygous, dark pigment (eumelanin) changes in colour accordingly. When several variants of the B-locus are found in heterozygous state, it is not possible to directly determine the influence on the eumelanin.

The overall genotype for the B-locus-complex can only be deduced if all known variants on the B-locus (bd, bc, bs, b4 and be) are analysed. Some of these alleles only exist in specific breeds.

Please note: The nomenclature of the results has been changed due to harmonizing efforts for genetic tests.

Haemophilia A (factor VIII deficiency) - PCR

pending

The current result is only valid for the sample submitted to our laboratory. The sender is responsible for the correct information regarding the sample material.The laboratory can not be made liable. Furthermore, any obligation for compensation is limited to the value of the tests performed.

There is a possibility that other mutations may have caused the disease/phenotype. The analysis was performed according to the latest knowledge and technology.

The laboratory is accredited for the performed tests according to DIN EN ISO/IEC 17025:2018. (except partner lab tests).

sample ID: 2208-N-10000

LABOKLIN
LABORATORIUM VOOR KLINISCHE DIAGNOSTIEK N.V.

*** END of report ***

Drs. N. Van Zon

**** LET OP! ***

Per 28-02-'22 is ons nieuwe adres:
Industriestraat 29
6433 JW Hoensbroek