

Inqaba Biotechnical Industries (Pty) Ltd **Animal Genetics Division**

ZO2023/5142/20230116/#39006

Co. Reg. No: 2001/011245/07

VAT No: 4150197251

B Locus (TYRP1)

Report No:

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Profile: DG2023/42217 Species: Canis lupus familiaris / Canine / Dog

981 020 000 817 821 Name: Gilda Microchip #:

Registration #: ZA009944B21 Breed: Rhodesian Ridgeback

Test: [B Locus (Can)] B Locus (TYRP1)

c.121T>A **BCBC** Results:

c.991C>T CT **BSbS** c.1033delCCT CCT/CCT **BDBD**

Final Conclusion Carrying brown

Extraction Method: DNA Extraction: D4069 Sample Type: Whole Blood (EDTA) Test Type: Genetic Colour

[B Locus (Can)] B Locus (TYRP1)

The B locus or Tyrosinase Related Protein TYRP1 gene is associated with eumelanin production.

The mutation described in this report contains SNP mutation c.121 T>A in the TYRP1 gene, designated BC for T and bC for A. Similarly, SNP mutation c.991C>T in the TYRP1 gene is designated BS for C and bS for T. The INDEL mutation c.1033_1036delCCT in the TYRP1 gene is designated BD for CCT and bD for the deletion. The SNP mutation c.555T>G uniquely identified in Australian Shepherds, designates BA for the T and bA for the G.

*A compound heterozygote result occurs when there are two b-alleles at different mutation sites. The final colour of the individual is dependent on whether:

- a) the b-alleles come from the same parent (the individual does not show the brown colour but carries brown) OR
- b) the b-alleles come from both parents (the individual is brown in colour)

Both parents will need to be tested for the B-locus to conclude the phenotype of a compound heterozygote.

As the mutations present with autosomal recessive inheritance, two mutant alleles are required to confer the phenotype.

References: Schmutz et al 2002. TYRP1 and MC1R genotypes and their effects on coat color in dogs. Mammalian Genome 7, pp380-387. Schmutz and Berryere, 2007. Genes affecting coat colour and pattern in domestic dogs: a review. Animal Genetics 38, pp539-549. Jancuskova et al 2018. TYRP:c.555T>G is a recurrent mutation found in Australian Shepherd and Miniature American Shepherd dogs. Animal Genetics 49, 496-501.

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