

Result certificate #084537

Detection of gene variants influencing coat length in dogs

Sample

Sample: 16-27701 Name: Nikki Valatheca Breed: Great Swiss Mountain Dog Microchip: 953 010 000 206 316 Reg. number: CMKU/VSP/2310/14/16 Date of birth: 10.12.2014 Sex: female Date received: 05.10.2016 Sample type: blood The identity of the animal has been checked by MVDr. Aleš Jurovic Customer Lucie Macková Valšov 135 79201 Bruntál Czech Republic

Result: N/N

Explanation

Presence of FGF5 gene variants influencing coat length in dogs was examined.

If the result is N/N – the dog does not carry any variant specific for long hair – the dog has short hair
If the result is N/FGF5 – the dog carries one copy of the variant FGF5 gene – the dog is short-haired, but it can give birth to long-haired offsprings, if suitably crossed.

• If the result is FGF5/FGF5 – the dog carries two variant alleles in the FGF5 gene – the dog is long-haired

Long coat phenotype is inherited in autosomal recessive trait. Long coated dogs have two variant alleles in the FGF5 gene (each from different parent). In case of mating two FGF5 carriers, theoretically, 25% long coated offspring will be born.

In some breeds, variant for long coat phenotype was not elucidated.

Method: SOP172-FGF5-dog, direct DNA sequencing

Report date: 10.10.2016 Responsible person: Mgr. Markéta Dajbychová, Deputy Laboratory Manager



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