

Result certificate #012345

Detection of c.660C>T mutation in ADAMTSL2 gene causing Musladin-Lueke syndrom in Beagle breed

Customer: Jan Novák, Dlouhá 1, 30000 Plzeň, Czech Republic

Sample: Sample: 21-12345 Date received: 01.02.2021 Sample type: blood

Information provided by the customer Name: Lassie DEMO Breed: Plemeno

Tattoo number: 1392013 Microchip: 123 456 789 012 345 Reg. number: REGQ12345 Date of birth: 1.1.2020 Sex: female Date of sampling: 01.02.2021 The identity of the animal has been checked.

Result: Mutation was not detected (N/N)

Legend: N/N = wild-type genotype. N/P = carrier of the mutation. P/P = mutated genotype (individual will be most probably affected with the disease). (N = negative, P = positive)

Explanation

Presence or absence of c.660C>T mutation in exon 7 of ADAMTSL2 gene causing Musladin-Lueke syndrom (MLS) in Beagle breed was tested. ADAMTSL2 gene is coding the fibrillin-1 protein which forms the main part of the tissue microfibriles. MLS is an inherited disease that affects development and structure of connective tissues and manifests itself by extensive skin and joint fibrosis and affects even heart.

Mutation that causes MLS in Beagle breed is inherited as an autosomal recessive trait. That means the disease affects dogs with P/P genotype only. The dogs with N/P genotype are considered carriers of the disease (heterozygotes). In offspring of two heterozygous animals following genotype distribution can be expected: 25 % N/N, 25 % P/P and 50 % N/P.

Method: SOPAgriseq_canine, ngs, accredited method

Date of issue: 06.02.2021 Date of testing: 01.02.2021 - 06.02.2021 Approved by: Mgr. Martina Šafrová, Laboratory Manager



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