

Detection of g.15295122-15295126del
mutation in SACS gene causing neuronal
degeneration in Great Pyrenees dogs

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 g.15295122-15295126del mutation in SACS gene causing neuronal degeneration in Great Pyrenees dogs was tested. The disease is characterised by degeneration of the central nervous system that leads to progressive cerebellar ataxia (movement coordination disorder) and spasticity (condition with increased muscle tone). The clinical symptoms appear very soon at the age of 4 months and sometimes even earlier. The disease gradually progresses during next several years and the symptoms get worse. The affected dogs have abnormal gait - clumsy or uncoordinated movement with a wide-base stance, slipping and sliding manoeuvring.

Mutation that causes neuronal degeneration is inherited autosomally recessively which means that the disease develops only in those dogs who inherit mutated allele from both parents; 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: SOP188-MPS-canine, MPS, accredited method

Date of issue: 06.02.2021

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Approved by: Mgr. Martina Šafrová, Laboratory Manager



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