

Detection of c.1080\_1089delTCGCCTGGAC  
mutation in NDR1 gene causing  
polyneuropathy in Greyhound 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.1080\_1089delTCGCCTGGAC mutation in NDR1 gene causing polyneuropathy in Greyhound breed was tested. The polyneuropathy belongs to hereditary neurological diseases and the symptoms include decreased or absent reflexes, muscle weakness, muscle pain and cramps and an odd "bunny-hopping" gait, etc. First clinical symptoms appear in young dogs at three to nine months of age and progress with age.

Mutation that causes polyneuropathy in Greyhound 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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