

Result certificate #012345

Detection of c.578C>T FGF5 gene variant influencing coat length in 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: N/M5

Explanation

Presence of c.578C>T (M5) of FGF5 gene variant influencing coat length in dogs of breeds Akita, Siberian Husky and Samoyed was examined.

Long coat phenotype is inherited in autosomal recessive trait. Dogs with long coat (result M5/M5) carry c.578T (M5) gene variant on both alleles; they inherited M5 from both parents. Individuals N/M5 are carriers of long coat. By mating two N/M5 short coated dogs, 25% M5/M5 long coated offspring can be expected. Dogs with N/N result have short coat and they do not carry any long coat variant.

Method: SOPAgriseq_canine, ngs

Date of issue: 06.02.2021

Date of testing: 01.02.2021 - 06.02.2021

Approved by: Mgr. Martina Šafrová, Laboratory Manager



Genomia s.r.o, Republikánská 6, 31200 Plzeň, Czech Republic www.genomia.cz, laborator@genomia.cz, tel: +420 373 749 999