



Orivet

Genetic Summary Report

Animal Name: Lotto Played In Paris

Owner:

Sabrina Rogers

Membership Number : Lagniappe Frenchies

Member Body/Breed Club: Not assigned

Approved Collection Method: No



Accredited and Compliant with



Members of

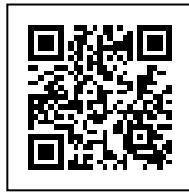


Harmonization of
Genetic Testing
for Dogs

orivet.com



Genetic Summary Report



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Owner's details

Name: Sabrina Rogers

Animal's Details

Registered Name : Lotto Played In Paris

Pet Name : Lotto Played In Paris

Registration Number :

Breed : French Bulldog

Microchip Number : 981020051726358

Sex : Female

Date of Birth : 17th Sep 2022

Colour : lilac and tan

Sample Collection Details

Case Number : 24A106599

Collected By :

Approved Collection : No

Sample Type : SWAB

Test Details

Test Requested : French Bulldog - Full Breed Profile

Pet Name : Lotto played in Paris

Date of Test : 18th Jun 2024

Authorisation

Sample with Lab ID Number 24A106599 was received at Orivet Genetics, DNA was extracted and analysed with the following result reported:

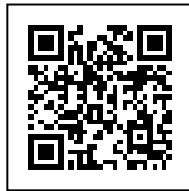
.....
George Sofronidis BSc (Hons)

.....
Dr Noam Pik BVSc, MAVS





Genetic Summary Report



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Animal's Details

Registered Name :	Lotto Played In Paris
Pet Name :	Lotto Played In Paris
Registration Number :	
Breed :	French Bulldog
Microchip Number :	981020051726358
Sex :	Female
Date of Birth :	17th Sep 2022
Colour :	lilac and tan

Tests Reported

Diseases	Result
Airway Distress Syndrome (ADAMTS3) - Risk Marker	NEGATIVE - NO COPIES OF THE ADAMTS3 VARIANT DETECTED
Cone-Rod Dystrophy I - PRA (crd -4/cord I)	NEGATIVE / CLEAR [NO VARIANT DETECTED]
Cystinuria Type 3 [Bulldog Risk Factor Variant 2&3]	NEGATIVE FOR HIGH RISK VARIANT 2 [SLC3A1] / NEGATIVE FOR LOW RISK VARIANT 3 [SLC7A9]
Degenerative Myelopathy	CARRIER [ONE COPY OF THE VARIANT DETECTED]
Hereditary Cataract	NEGATIVE / CLEAR [NO VARIANT DETECTED]
Hyperuricosuria	NEGATIVE / CLEAR [NO VARIANT DETECTED]

Owner's Name : Sabrina Rogers

Pet Name : Lotto played in Paris

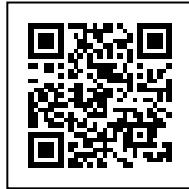
Microchip Number 981020051726358

Approved Collection Method : No





Genetic Summary Report



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Tests Reported

Diseases	Result
Multifocal Retinopathy CMR1 (Mastiff/Bull Breeds Type)	NEGATIVE / CLEAR [NO VARIANT DETECTED]

Traits	Result
E Locus - (Cream/Red/Yellow)	E/e - BLACK CARRIES EXTENSION [YELLOW/WHITE/APRICOT/RUBY/RED]
EM (MC1R) Locus - Melanistic Mask	E ⁿ /E ⁿ - NO MELANISTIC MASK (E ⁿ) EXTENSION ALLELE
I Pheomelanin Locus Colour Intensity	i/i - TWO COPIES OF THE MFSD12 INTENSITY ALLELE (LIKELY TO SHOW EXTREME DILUTION)
Brown Deletion = Bd	B ^d /B ^d - DOES NOT CARRY BROWN/RED/LIVER or CHOCOLATE [DELETION]
Brown Stop Codon = Bs	B ^s /B ^s - DOES NOT CARRY BROWN/RED/LIVER or CHOCOLATE [STOP CODON]
Brown Insertion = Bc	B ^c /B ^c - DOES NOT CARRY BROWN/RED/LIVER or CHOCOLATE [INSERTION]
Cocoa/Brown HPS3 [French Bulldog Type] = Co	b ^{co} /b ^{co} - TWO COPIES OF HPS3 COCOA VARIANT
D (Dilute) Locus	d/d - GREY, GREY BLUE, OR SILVER - COLOUR IS DILUTED
Dilute D2 Variant (Chow Chow Type)	D ² /D ² - NO COPY OF d2 ALLELE (DILUTE) - PIGMENT IS NORMAL
K Locus (Dominant Black)	k ^y /k ^y - RECESSIVE NON-BLACK [COLOUR PATTERN DETERMINED BY A LOCUS]

Owner's Name : Sabrina Rogers

Pet Name : Lotto played in Paris

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Approved Collection Method : No





Genetic Summary Report



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Tests Reported

Traits	Result
A Locus (Fawn/Sable;Tri/Tan Points)	a ^t /a ^t - TAN POINTS/BLACK & TAN or TRICOLOUR MAYBE BRINDLED [SEE K LOCUS]
Pied (BOTH SINE and REPEAT VARIANTS)	S/S - NO PIEBALD, WHITE SPOTTING, FLASH OR PARTI COAT COLOUR
M Locus (Merle/Dapple)	m [171bp] / m [171bp] - NON MERLE SOLID COAT (NO CHANGE TO COAT or EYE COLOUR)
Oculocutaneous Albinism (Small Breed Type)	NEGATIVE - NOT SHOWING THE PHENOTYPE
Long Hair Gene (Canine C95F)	NEGATIVE - NOT SHOWING THE PHENOTYPE
Long Hair Gene (Canine DUP GG)	NEGATIVE - NOT SHOWING THE PHENOTYPE
Body Size IGSF1 "Bulky Gene"	HOMOZYGOUS FOR INSULIN LIKE GROWTH FACTOR (IGF1R) - ASSOCIATED WITH AN INCREASE of BODY(BULKY) SIZE
Screw Tail (DVL2)	TWO COPIES OF THE DVL2 (SCREW TAIL PHENOTYPE) GENE VARIANT DETECTED

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Glossary of Genetic Terms (Results)



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NEGATIVE / CLEAR [NO VARIANT DETECTED]

No presence of the variant (mutation) has been detected. The animal is clear of the disease and will not pass on any disease-causing mutation.

CARRIER [ONE COPY OF THE VARIANT DETECTED]

This is also referred to as HETEROZYGOUS. One copy of the normal gene and copy of the affected (mutant) gene has been detected. The animal will not exhibit disease symptoms or develop the disease. Consideration needs to be taken if breeding this animal - if breeding with another carrier or affected or unknown then it may produce an affected offspring.

POSITIVE / AT RISK [TWO COPIES OF THE VARIANT DETECTED]

Two copies of the disease gene variant (mutation) have been detected also referred to as HOMOZYGOUS for the variant. The animal may show symptoms (affected) associated with the disease. Appropriate treatment should be pursued by consulting a Veterinarian.

POSITIVE HETEROZYGOUS [ONE COPY OF THE DOMINANT VARIANT DETECTED]

Also referred to as POSITIVE ONE COPY or POSITIVE HETEROZYGOUS. This result is associated with a disease that has a dominant mode of inheritance. One copy of the normal gene (wild type) and affected (mutant) gene is present. Appropriate treatment should be pursued by consulting a Veterinarian. This result can still be used to produce a clear offspring.

NORMAL BY PARENTAGE HISTORY

The sample submitted has had its parentage verified by DNA. By interrogating the DNA profiles of the Dam, Sire and Offspring this information together with the history submitted for the parents excludes this animal from having this disease. The controls run confirm that the dog is NORMAL for the disease requested.

NORMAL BY PEDIGREE

The sample submitted has had its parentage verified by Pedigree. The pedigree has been provided and details (genetic testing reports) of the parents have been included. Parentage could not be determined via DNA profile as no sample was submitted.

NO RESULTS AVAILABLE

Insufficient information has been provided to provide a result for this test. Sire and Dam information and/or sample may be required. This result is mostly associated with tests that have a patent/license and therefore certain restrictions apply. Please contact the laboratory to discuss.

INDETERMINABLE

The sample submitted has failed to give a conclusive result. This result is mainly due to the sample failing to "cluster" or result in the current grouping. A recollection is required at no charge.

DNA PROFILE

Also known as a DNA fingerprint. This is unique for the animal. No animal shares the same DNA profile. An individual's DNA profile is inherited from both parents and can be used for verifying parentage (pedigrees). This profile contains no disease or trait information and is simply a unique DNA signature for that animal.

Glossary of Genetic Terms (Results)



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PARENTAGE VERIFICATION/ QUALIFIES/CONFIRMED OR DOES NOT QUALIFY/EXCLUDED

Parentage is determined by examining the markers on the DNA profile. A result is generated and stated for all DNA parentage requests. Parentage confirmation reports can only be generated if a DNA profile has been carried out for Dam, Offspring and possible Sire/s.

PENDING

PENDING

TRAIT (PHENOTYPE)

A feature that an animal is born with (a genetically determined characteristic). Traits are a visual phenotype that range from colour to hair length, and also includes certain features such as tail length. If an individual is AFFECTED for a trait then it will show that characteristic eg. AFFECTED for the B (Brown) Locus or bb will be brown/chocolate.

POSITIVE – SHOWING THE PHENOTYPE

The animal is showing the trait or phenotype tested.

CLARIFICATION OF GENETIC TESTING

The goal of genetic testing is to provide breeders with relevant information to improve breeding practices in the interest of animal health. However, genetic inheritance is not a simple process, and may be complicated by several factors. Below is some information to help clarify these factors.

The goal of genetic testing is to provide breeders with relevant information to improve breeding practices in the interest of animal health. However, genetic inheritance is not a simple process, and may be complicated by several factors. Below is some information to help clarify these factors.

- 1) Some diseases may demonstrate signs of what Geneticists call "genetic heterogeneity". This is a term to describe an apparently single condition that may be caused by more than one mutation and/or gene
- 2) It is possible that there exists more than one disease that presents in a similar fashion and segregates in a single breed. These conditions –although phenotypically similar – may be caused by separate mutations and/or genes.
- 3) It is possible that the disease affecting your breed may be what Geneticists call an "oligogenic disease". This is a term to describe the existence of additional genes that may modify the action of a dominant gene associated with a disease. These modifier genes may for example give rise to a variable age of onset for a particular condition, or affect the penetrance of a particular mutation such that some animals may never develop the condition.

The range of hereditary diseases continues to increase and we see some that are relatively benign and others that can cause severe and/or fatal disease. Diagnosis of any disease should be based on pedigree history, clinical signs, history (incidence) of the disease and the specific genetic test for the disease. Penetrance of a disease will always vary not only from breed to breed but within a breed, and will vary with different diseases. Factors that influence penetrance are genetics, nutrition and environment. Although genetic testing should be a priority for breeders, we strongly recommend that temperament and phenotype also be considered when breeding.

Orivet Genetic Pet Care aims to frequently update breeders with the latest research from the scientific literature. If breeders have any questions regarding a particular condition, please contact us on (03) 9534 1544 or admin@orivet.com and we will be happy to work with you to answer any relevant questions.