

**ORTHOPEDIC FOUNDATION FOR ANIMALS, INC.**

**OLAF**  
*registered name*

**POODLE**  
*breed*

*film/test/lab #*

**900020180725012**  
*tattoo/microchip/DNA profile*

**2616681**  
*application number*

**04/04/2025**  
*date of report*

**RESULTS:**

The elbows are normal. No radiographic evidence of elbow dysplasia is present.

**NOREG2616681**  
*registration no.*

**M**  
*sex*

**07/21/2019**  
*date of birth*

**67**  
*age at evaluation in months*



A Not-For-Profit Organization

**PO-EL9994M67-C-VPI**  
*O.F.A. NUMBER*

*This number issued with the right to correct or  
revoke by the Orthopedic Foundation for Animals.*

**NORMAL**

**owner**  
WAYNE MILLER  
NAOMI  
3750 TR 124  
MILLERSBURG OH 44654



G.G. KELLER, DVM, MS, DACVR  
CHIEF OF VETERINARY SERVICES

[www.ofa.org](http://www.ofa.org)

This electronic OFA certificate was generated on: 04/04/2025

This certification can be verified on the OFA website by entering the dog's registration number into the orange search box located at the top of the page or by scanning the QR code above.

If there are any errors on this certificate, please email [CORRECTIONS@OFA.ORG](mailto:CORRECTIONS@OFA.ORG) to request a correction.

Orthopedic Foundation for Animals, Inc.  
2300 E. Nifong Blvd.  
Columbia, MO 65201-3806

OFA website: [www.ofa.org](http://www.ofa.org)  
E-mail address: [ofa@ofa.org](mailto:ofa@ofa.org)  
Phone number: 573-442-0418  
Fax number: 573-875-5073

ORTHOPEDIC FOUNDATION FOR ANIMALS, INC.

OLAF  
*registered name*

POODLE  
*breed*

*film/test/lab #*

900020180725012  
*tattoo/microchip/DNA profile*

2616681  
*application number*

04/04/2025  
*date of report*

RESULTS:

No radiographic evidence of hip dysplasia is present. The consensus evaluation is: GOOD

NOREG2616681  
*registration no.*

M  
*sex*

07/21/2019  
*date of birth*

67  
*age at evaluation in months*



A Not-For-Profit Organization

PO-35629G67M-C-VPI  
O.F.A. NUMBER

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revoke by the Orthopedic Foundation for Animals.*

owner  
WAYNE MILLER  
NAOMI  
3750 TR 124  
MILLERSBURG OH 44654

OFA eCert



Verify QR scan

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ORTHOPEDIC FOUNDATION FOR ANIMALS, INC.

OLAF  
*registered name*

POODLE, MINIATURE  
*breed*

*film/test/lab #*

900020180725012  
*tattoo/microchip/DNA profile*

2616681  
*application number*

07/17/2025  
*date of report*

**RESULTS:**

Based upon the exam dated 05/09/2025, this dog has been found to be free of observable inherited eye disease and has been issued an Eye Certification Registry Number which is valid for one year from the time of the exam.

NOREG2616681  
*registration no.*

M  
*sex*

07/21/2019  
*date of birth*

69  
*age at evaluation in months*



A Not-For-Profit Organization



PO-EYE13610/69M-VPI  
*O.F.A. NUMBER*

*This number issued with the right to correct or revoke by the Orthopedic Foundation for Animals.*

NORMAL

**owner**  
WAYNE MILLER  
NAOMI MILLER  
3750 TR 124  
MILLERSBURG OH 44654

OFA eCert



Verify QR scan

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**OLAF**



DNA Test Report

Test Date: September 26th, 2020

[embk.me/olaf61](https://embk.me/olaf61)

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## GENETIC STATS

Predicted adult weight: **23 lbs**

Genetic age: **20 human years**

Based on the date of birth you provided

## TEST DETAILS

Kit number: EM-29149056

Swab number: 31200760602361





**OLAF**

## TRAITS: COAT COLOR

**TRAIT**

**RESULT**

**E Locus (MC1R)**

The E Locus determines if and where a dog can produce dark (black or brown) hair. Dogs with two copies of the recessive **e** allele do not produce dark hairs at all, and will be "red" over their entire body. The shade of red, which can range from a deep copper to yellow/gold to cream, is dependent on other genetic factors including the Intensity (I) Locus, which has yet to be genetically mapped. In addition to determining if a dog can develop dark hairs at all, the E Locus can give a dog a black "mask" or "widow's peak," unless the dog has overriding coat color genetic factors. Dogs with one or two copies of the **Em** allele usually have a melanistic mask (dark facial hair as commonly seen in the German Shepherd and Pug). Dogs with no copies of **Em** but one or two copies of the **Eg** allele usually have a melanistic "widow's peak" (dark forehead hair as commonly seen in the Afghan Hound and Borzoi, where it is called either "grizzle" or "domino").

**No dark hairs anywhere (ee)**

**K Locus (CBD103)**

The K Locus **K<sup>B</sup>** allele "overrides" the A Locus, meaning that it prevents the A Locus genotype from affecting coat color. For this reason, the **K<sup>B</sup>** allele is referred to as the "dominant black" allele. As a result, dogs with at least one **K<sup>B</sup>** allele will usually have solid black or brown coats (or red/cream coats if they are **ee** at the E Locus) regardless of their genotype at the A Locus, although several other genes could impact the dog's coat and cause other patterns, such as white spotting. Dogs with the **k<sup>Y</sup>k<sup>Y</sup>** genotype will show a coat color pattern based on the genotype they have at the A Locus. Dogs who test as **K<sup>B</sup>k<sup>Y</sup>** may be brindle rather than black or brown.

**Not expressed (k<sup>Y</sup>k<sup>Y</sup>)**

**A Locus (ASIP)**

The A Locus controls switching between black and red pigment in hair cells, but it will only be expressed in dogs that are not **ee** at the E Locus and are **k<sup>Y</sup>k<sup>Y</sup>** at the K Locus. Sable (also called "Fawn") dogs have a mostly or entirely red coat with some interspersed black hairs. Agouti (also called "Wolf Sable") dogs have red hairs with black tips, mostly on their head and back. Black and tan dogs are mostly black or brown with lighter patches on their cheeks, eyebrows, chest, and legs. Recessive black dogs have solid-colored black or brown coats.

**Not expressed (a<sup>t</sup>a<sup>t</sup>)**



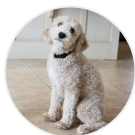
## TRAITS: COAT COLOR (CONTINUED)

TRAIT	RESULT
<b>D Locus (MLPH)</b>  Dogs with two copies of the <b>d</b> allele will have all black pigment lightened ("diluted") to gray, or brown pigment lightened to lighter brown in their hair, skin, and sometimes eyes. There are many breed-specific names for these dilute colors, such as "blue", "charcoal", "fawn", "silver", and "Isabella". Note that dilute dogs have a higher incidence of Color Dilution Alopecia, especially in certain breeds. Dogs with one copy of the <b>d</b> allele will not be dilute, but can pass the <b>d</b> allele on to their puppies.	<b>Not expressed (DD)</b>
<b>B Locus (TYRP1)</b>  Dogs with two copies of the <b>b</b> allele produce brown pigment instead of black in both their hair and skin. Dogs with one copy of the <b>b</b> allele will produce black pigment, but can pass the <b>b</b> allele on to their puppies. E Locus <b>ee</b> dogs that carry two <b>b</b> alleles will have red or cream coats, but have brown noses, eye rims, and footpads (sometimes referred to as "Dudley Nose" in Labrador Retrievers). "Liver" or "chocolate" is the preferred color term for brown in most breeds; in the Doberman Pinscher it is referred to as "red".	<b>Likely black colored nose/feet (Bb)</b>
<b>Saddle Tan (RALY)</b>  The "Saddle Tan" pattern causes the black hairs to recede into a "saddle" shape on the back, leaving a tan face, legs, and belly, as a dog ages. The Saddle Tan pattern is characteristic of breeds like the Corgi, Beagle, and German Shepherd. Dogs that have the <b>II</b> genotype at this locus are more likely to be mostly black with tan points on the eyebrows, muzzle, and legs as commonly seen in the Doberman Pinscher and the Rottweiler. This gene modifies the A Locus <b>a<sup>t</sup></b> allele, so dogs that do not express <b>a<sup>t</sup></b> are not influenced by this gene.	<b>Not expressed (II)</b>



## TRAITS: COAT COLOR (CONTINUED)

TRAIT	RESULT
<b>S Locus (MITF)</b>  The S Locus determines white spotting and pigment distribution. MITF controls where pigment is produced, and an insertion in the MITF gene causes a loss of pigment in the coat and skin, resulting in white hair and/or pink skin. Dogs with two copies of this variant will likely have breed-dependent white patterning, with a nearly white, parti, or piebald coat. Dogs with one copy of this variant will have more limited white spotting and may be considered flash, parti or piebald. This MITF variant does not explain all white spotting patterns in dogs and other variants are currently being researched. Some dogs may have small amounts of white on the paws, chest, face, or tail regardless of their S Locus genotype.	<b>Likely flash, parti, or piebald (Ssp)</b>
<b>M Locus (PMEL)</b>  Merle coat patterning is common to several dog breeds including the Australian Shepherd, Catahoula Leopard Dog, and Shetland Sheepdog, among many others. Merle arises from an unstable SINE insertion (which we term the "M*" allele) that disrupts activity of the pigmentary gene PMEL, leading to mottled or patchy coat color. Dogs with an <b>M*m</b> result are likely to be phenotypically merle or could be "phantom" merle, that is, they have a merle allele that does not affect coat color. Dogs with an <b>M*M*</b> result are likely to be phenotypically merle or double merle. Dogs with an <b>mm</b> result have no merle alleles and are unlikely to have a merle coat pattern.  Note that Embark does not currently distinguish between the recently described cryptic, atypical, atypical+, classic, and harlequin merle alleles. Our merle test only detects the presence, but not the length of the SINE insertion. We do not recommend making breeding decisions on this result alone. Please pursue further testing for allelic distinction prior to breeding decisions.	<b>No merle alleles (mm)</b>
<b>H Locus (Harlequin)</b>  This pattern is recognized in Great Danes and causes dogs to have a white coat with patches of darker pigment. A dog with an <b>Hh</b> result will be harlequin if they are also <b>M*m</b> or <b>M*M*</b> at the M Locus and are not <b>ee</b> at the E locus. Dogs with a result of <b>hh</b> will not be harlequin. This trait is thought to be homozygous lethal; a living dog with an <b>HH</b> genotype has never been found.	<b>No harlequin alleles (hh)</b>



# OLAF



DNA Test Report

Test Date: September 26th, 2020

embk.me/olaf61

## TRAITS: OTHER COAT TRAITS

TRAIT	RESULT
<b>Furnishings (RSP02) LINKAGE</b>	
Dogs with one or two copies of the <b>F</b> allele have "furnishings": the mustache, beard, and eyebrows characteristic of breeds like the Schnauzer, Scottish Terrier, and Wire Haired Dachshund. A dog with two <b>I</b> alleles will not have furnishings, which is sometimes called an "improper coat" in breeds where furnishings are part of the breed standard. The mutation is a genetic insertion which we measure indirectly using a linkage test highly correlated with the insertion.	<b>Likely furnished (mustache, beard, and/or eyebrows) (FF)</b>
<b>Coat Length (FGF5)</b>	
The FGF5 gene is known to affect hair length in many different species, including cats, dogs, mice, and humans. In dogs, the <b>T</b> allele confers a long, silky haircoat as observed in the Yorkshire Terrier and the Long Haired Whippet. The ancestral <b>G</b> allele causes a shorter coat as seen in the Boxer or the American Staffordshire Terrier. In certain breeds (such as Corgi), the long haircoat is described as "fluff."	<b>Likely long coat (TT)</b>
<b>Shedding (MC5R)</b>	
Dogs with at least one copy of the ancestral <b>C</b> allele, like many Labradors and German Shepherd Dogs, are heavy or seasonal shedders, while those with two copies of the <b>T</b> allele, including many Boxers, Shih Tzus and Chihuahuas, tend to be lighter shedders. Dogs with furnished/wire-haired coats caused by RSP02 (the furnishings gene) tend to be low shedders regardless of their genotype at this gene.	<b>Likely light shedding (TT)</b>
<b>Coat Texture (KRT71)</b>	
Dogs with a long coat and at least one copy of the <b>T</b> allele have a wavy or curly coat characteristic of Poodles and Bichon Frises. Dogs with two copies of the ancestral <b>C</b> allele are likely to have a straight coat, but there are other factors that can cause a curly coat, for example if they at least one <b>F</b> allele for the Furnishings (RSP02) gene then they are likely to have a curly coat. Dogs with short coats may carry one or two copies of the <b>T</b> allele but still have straight coats.	<b>Likely curly coat (TT)</b>



**OLAF**



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## TRAITS: OTHER COAT TRAITS (CONTINUED)

TRAIT	RESULT
<b>Hairlessness (FOXI3) LINKAGE</b>	
A duplication in the FOXI3 gene causes hairlessness over most of the body as well as changes in tooth shape and number. This mutation occurs in Peruvian Inca Orchid, Xoloitzcuintli (Mexican Hairless), and Chinese Crested (other hairless breeds have different mutations). Dogs with the <b>NDup</b> genotype are likely to be hairless while dogs with the <b>NN</b> genotype are likely to have a normal coat. The <b>DupDup</b> genotype has never been observed, suggesting that dogs with that genotype cannot survive to birth. Please note that this is a linkage test, so it may not be as predictive as direct tests of the mutation in some lines.	<b>Very unlikely to be hairless (NN)</b>
<b>Hairlessness (SGK3)</b>	
Hairlessness in the American Hairless Terrier arises from a mutation in the SGK3 gene. Dogs with the <b>ND</b> genotype are likely to be hairless while dogs with the <b>NN</b> genotype are likely to have a normal coat.	<b>Very unlikely to be hairless (NN)</b>
<b>Oculocutaneous Albinism Type 2 (SLC45A2) LINKAGE</b>	
Dogs with two copies <b>DD</b> of this deletion in the SLC45A2 gene have oculocutaneous albinism type 2 (OCA2), also known as Doberman Z Factor Albinism, a recessive condition characterized by severely reduced or absent pigment in the eyes, skin, and hair. Affected dogs sometimes suffer from vision problems due to lack of eye pigment (which helps direct and absorb ambient light) and are prone to sunburn. Dogs with a single copy of the deletion <b>ND</b> will not be affected but can pass the mutation on to their offspring. This particular mutation can be traced back to a single white Doberman Pinscher born in 1976, and it has only been observed in dogs descended from this individual. Please note that this is a linkage test, so it may not be as predictive as direct tests of the mutation in some lines.	<b>Likely not albino (NN)</b>



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## TRAITS: OTHER BODY FEATURES

TRAIT	RESULT
<b>Muzzle Length (BMP3)</b>  Dogs in medium-length muzzle (mesocephalic) breeds like Staffordshire Terriers and Labradors, and long muzzle (dolichocephalic) breeds like Whippet and Collie have one, or more commonly two, copies of the ancestral <b>C</b> allele. Dogs in many short-length muzzle (brachycephalic) breeds such as the English Bulldog, Pug, and Pekingese have two copies of the derived <b>A</b> allele. At least five different genes affect muzzle length in dogs, with BMP3 being the only one with a known causal mutation. For example, the skull shape of some breeds, including the dolichocephalic Scottish Terrier or the brachycephalic Japanese Chin, appear to be caused by other genes. Thus, dogs may have short or long muzzles due to other genetic factors that are not yet known to science.	<b>Likely medium or long muzzle (CC)</b>
<b>Tail Length (T)</b>  Whereas most dogs have two <b>C</b> alleles and a long tail, dogs with one <b>G</b> allele are likely to have a bobtail, which is an unusually short or absent tail. This mutation causes natural bobtail in many breeds including the Pembroke Welsh Corgi, the Australian Shepherd, and the Brittany Spaniel. Dogs with <b>GG</b> genotypes have not been observed, suggesting that dogs with the <b>GG</b> genotype do not survive to birth. Please note that this mutation does not explain every natural bobtail! While certain lineages of Boston Terrier, English Bulldog, Rottweiler, Miniature Schnauzer, Cavalier King Charles Spaniel, and Parson Russell Terrier, and Dobermans are born with a natural bobtail, these breeds do not have this mutation. This suggests that other unknown genetic mutations can also lead to a natural bobtail.	<b>Likely normal-length tail (CC)</b>
<b>Hind Dewclaws (LMBR1)</b>  Common in certain breeds such as the Saint Bernard, hind dewclaws are extra, nonfunctional digits located midway between a dog's paw and hock. Dogs with at least one copy of the <b>T</b> allele have about a 50% chance of having hind dewclaws. Note that other (currently unknown to science) mutations can also cause hind dewclaws, so some <b>TT</b> or <b>TC</b> dogs will have hind dewclaws.	<b>Likely to have hind dew claws (CT)</b>



**OLAF**



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## TRAITS: OTHER BODY FEATURES (CONTINUED)

TRAIT	RESULT
<p><b>Blue Eye Color (ALX4) LINKAGE</b></p> <p>Embark researchers discovered this large duplication associated with blue eyes in Arctic breeds like Siberian Husky as well as tri-colored (non-merle) Australian Shepherds. Dogs with at least one copy of the duplication (<b>Dup</b>) are more likely to have at least one blue eye. Some dogs with the duplication may have only one blue eye (complete heterochromia) or may not have blue eyes at all; nevertheless, they can still pass the duplication and the trait to their offspring. <b>NN</b> dogs do not carry this duplication, but may have blue eyes due to other factors, such as merle. Please note that this is a linkage test, so it may not be as predictive as direct tests of the mutation in some lines.</p>	<p><b>Less likely to have blue eyes (NN)</b></p>
<p><b>Back Muscling &amp; Bulk, Large Breed (ACSL4)</b></p> <p>The <b>T</b> allele is associated with heavy muscling along the back and trunk in characteristically "bulky" large-breed dogs including the Saint Bernard, Bernese Mountain Dog, Greater Swiss Mountain Dog, and Rottweiler. The "bulky" <b>T</b> allele is absent from leaner shaped large breed dogs like the Great Dane, Irish Wolfhound, and Scottish Deerhound, which are fixed for the ancestral <b>C</b> allele. Note that this mutation does not seem to affect muscling in small or even mid-sized dog breeds with notable back muscling, including the American Staffordshire Terrier, Boston Terrier, and the English Bulldog.</p>	<p><b>Likely normal muscling (CC)</b></p>



# OLAF



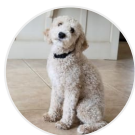
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## TRAITS: BODY SIZE

TRAIT	RESULT
<b>Body Size (IGF1)</b> The I allele is associated with smaller body size.	<b>Smaller (II)</b>
<b>Body Size (IGFR1)</b> The A allele is associated with smaller body size.	<b>Larger (GG)</b>
<b>Body Size (STC2)</b> The A allele is associated with smaller body size.	<b>Intermediate (TA)</b>
<b>Body Size (GHR - E191K)</b> The A allele is associated with smaller body size.	<b>Smaller (AA)</b>
<b>Body Size (GHR - P177L)</b> The T allele is associated with smaller body size.	<b>Larger (CC)</b>



# OLAF



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## TRAITS: PERFORMANCE

### TRAIT

### RESULT

#### Altitude Adaptation (EPAS1)

This mutation causes dogs to be especially tolerant of low oxygen environments (hypoxia), such as those found at high elevations. Dogs with at least one **A** allele are less susceptible to "altitude sickness." This mutation was originally identified in breeds from high altitude areas such as the Tibetan Mastiff.

**Normal altitude tolerance (GG)**

#### Appetite (POMC) LINKAGE

This mutation in the POMC gene is found primarily in Labrador and Flat Coated Retrievers. Compared to dogs with no copies of the mutation (**NN**), dogs with one (**ND**) or two (**DD**) copies of the mutation are more likely to have high food motivation, which can cause them to eat excessively, have higher body fat percentage, and be more prone to obesity. Read more about the genetics of POMC, and learn how you can contribute to research, in our blog post (<https://embarkvet.com/resources/blog/pomc-dogs/>). We measure this result using a linkage test.

**Normal food motivation (NN)**



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## CLINICAL TOOLS

These clinical genetic tools can inform clinical decisions and diagnoses. These tools do not predict increased risk for disease.

### Alanine Aminotransferase Activity (GPT)

Olaf's baseline ALT level is Low Normal

#### Why is this important to your vet?

Olaf has one copy of a variant associated with reduced ALT activity as measured on veterinary blood chemistry panels. Please inform your veterinarian that Olaf has this genotype, as ALT is often used as an indicator of liver health and Olaf is likely to have a lower than average resting ALT activity. As such, an increase in Olaf's ALT activity could be evidence of liver damage, even if it is within normal limits by standard ALT reference ranges.

#### What is Alanine Aminotransferase Activity?

Alanine aminotransferase (ALT) is a clinical tool that can be used by veterinarians to better monitor liver health. This result is not associated with liver disease. ALT is one of several values veterinarians measure on routine blood work to evaluate the liver. It is a naturally occurring enzyme located in liver cells that helps break down protein. When the liver is damaged or inflamed, ALT is released into the bloodstream.

#### How vets diagnose this condition

Genetic testing is the only way to provide your veterinarian with this clinical tool.

#### How this condition is treated

Veterinarians may recommend blood work to establish a baseline ALT value for healthy dogs with one or two copies of this variant.



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## INBREEDING AND DIVERSITY

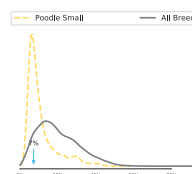
### CATEGORY

#### Coefficient Of Inbreeding

Our genetic COI measures the proportion of your dog's genome where the genes on the mother's side are identical by descent to those on the father's side.

### RESULT

7%

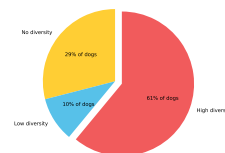


#### MHC Class II - DLA DRB1

A Dog Leukocyte Antigen (DLA) gene, DRB1 encodes a major histocompatibility complex (MHC) protein involved in the immune response. Some studies have shown associations between certain DRB1 haplotypes and autoimmune diseases such as Addison's disease (hypoadrenocorticism) in certain dog breeds, but these findings have yet to be scientifically validated.

### High Diversity

How common is this amount of diversity in purebreds:



#### MHC Class II - DLA DQA1 and DQB1

DQA1 and DQB1 are two tightly linked DLA genes that code for MHC proteins involved in the immune response. A number of studies have shown correlations of DQA-DQB1 haplotypes and certain autoimmune diseases; however, these have not yet been scientifically validated.

### High Diversity

How common is this amount of diversity in purebreds:

