

BREED ANCESTRY

 Australian Shepherd : 100.0%

GENETIC STATS

Predicted adult weight: **43 lbs**

Life stage: **Young adult**

Based on your dog's date of birth provided.

TEST DETAILS

Kit number: EM-47902098

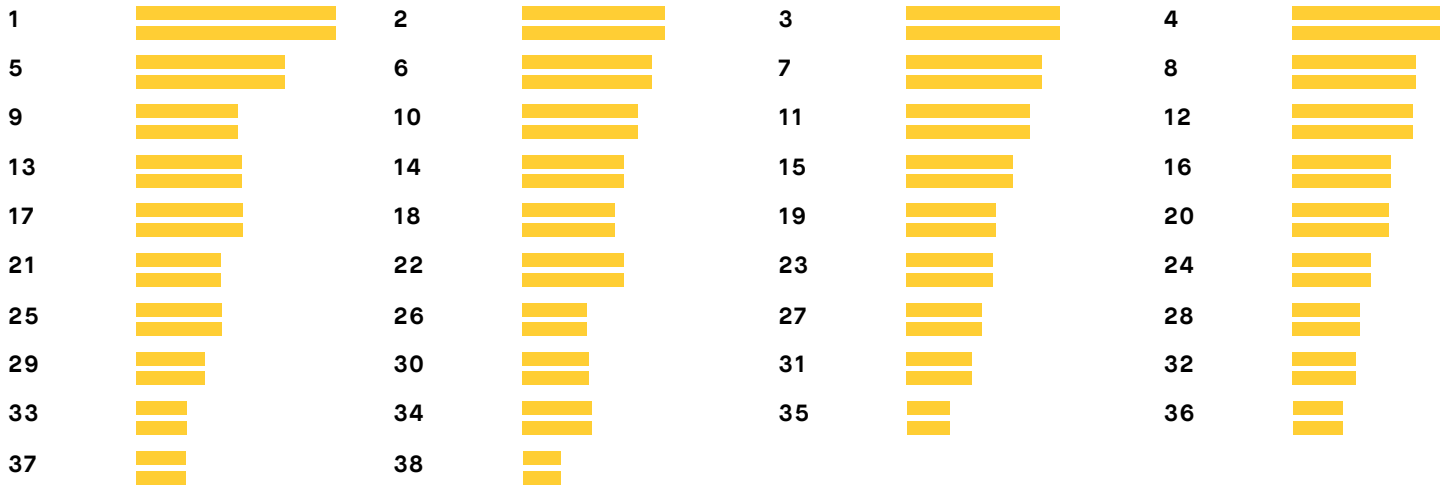
Swab number: 31220910602933

BREED ANCESTRY BY CHROMOSOME

Our advanced test identifies from where Smooch inherited every part of the chromosome pairs in her genome.

Breed colors:

Australian Shepherd



AUSTRALIAN SHEPHERD

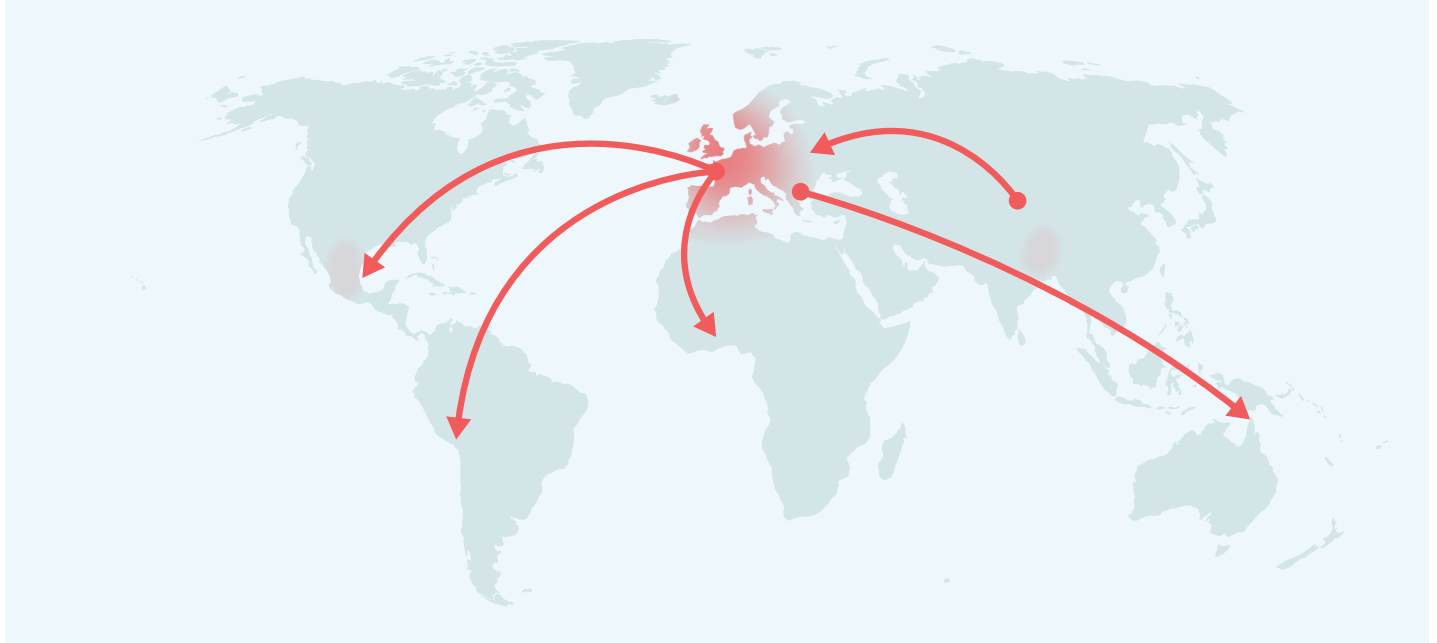


Fun Fact

Australian Shepherds rose to popularity and fame as rodeo stars. After the first World War, people flocked to the west and to watch exhibitions that showcased these very talented canines.

The Australian Shepherd, or Aussie, is the 17th most popular dog in the United States, and given their intelligence and temperament, it's no wonder they're so well-loved. Despite their name, the Australian Shepherd actually originated from the ranches of the United States around the 1800s. They are praised by stockmen and breeders for their trainability and intelligence. They have a medium build and a wide variation of different coat colors. Australian Shepherds have considerable energy and they usually need a job to do to keep themselves entertained, though they're also happy to spend time with the family and settle down at the end of the day. Australian Shepherds are often employed as guide dogs, rescue dogs, and therapy dogs. In addition to exercising an Aussie, it's equally important to keep their mind occupied, as if an Australian Shepherd gets bored they do have the tendency to invent their own games or activities, which sometimes involve destructive behaviors. This is a breed that thrives on close companionship. Aussies are at times called "Velcro Dogs" for their tendency to stay close to their owner.

MATERNAL LINE



Through Smooch's mitochondrial DNA we can trace her mother's ancestry back to where dogs and people first became friends. This map helps you visualize the routes that her ancestors took to your home. Their story is described below the map.

HAPLOGROUP: B1

B1 is the second most common maternal lineage in breeds of European or American origin. It is the female line of the majority of Golden Retrievers, Basset Hounds, and Shih Tzus, and about half of Beagles, Pekingese and Toy Poodles. This lineage is also somewhat common among village dogs that carry distinct ancestry from these breeds. We know this is a result of B1 dogs being common amongst the European dogs that their conquering owners brought around the world, because nowhere on earth is it a very common lineage in village dogs. It even enables us to trace the path of (human) colonization: Because most Bichons are B1 and Bichons are popular in Spanish culture, B1 is now fairly common among village dogs in Latin America.

HAPLOTYPE: B84

Part of the large B1 haplogroup, this haplotype occurs most frequently in Golden Retrievers, Beagles, and Staffordshire Terriers.

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** variant do not produce dark hairs and will express a red pigment called pheomelanin over their entire body. The shade of red, which can range from a deep copper to white, depends on other genetic factors, including the Intensity loci. In addition to determining if a dog can develop dark hairs, the E Locus can give a dog a black "mask" or "widow's peak" unless the dog has overriding coat color genetic factors.

No dark mask or grizzle (EE)

Dogs with one or two copies of the **E^m** variant may have a melanistic mask (dark facial hair as commonly seen in the German Shepherd Dog and Pug). In the absence of **E^m**, dogs with the **E^g** variant can have a "grizzle" phenotype (darker color on the head and top with a melanistic "widow's peak" and a lighter underside, commonly seen in the Afghan Hound and Borzoi and also referred to as "domino"). In the absence of both **E^m** and **E** variants, dogs with the **E^a** or **E^h** variants can express the grizzle phenotype. Additionally, a dog with any combination of two of the **E^g**, **E^a**, or **E^h** variants (example: **E^gE^a**) is also expected to express the grizzle phenotype.

K Locus (CBD103)

The K Locus **K^B** allele "overrides" the A Locus, meaning that it prevents the A Locus genotype from affecting coat color. For this reason, the **K^B** allele is referred to as the "dominant black" allele. As a result, dogs with at least one **K^B** 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^Yk^Y** genotype will show a coat color pattern based on the genotype they have at the A Locus. Dogs who test as **K^Bk^Y** may be brindle rather than black or brown.

More likely to have a mostly solid black or brown coat (K^Bk^Y)

TRAITS: COAT COLOR (CONTINUED)

| | |
|--------------|---------------|
| TRAIT | RESULT |
|--------------|---------------|

Intensity Loci

Areas of a dog's coat where dark (black or brown) pigment is not expressed either contain red/yellow pigment, or no pigment at all. Five locations across five chromosomes explain approximately 70% of red pigmentation "intensity" variation across all dogs. Dogs with a result of **Intense Red Pigmentation** will likely have deep red hair like an Irish Setter or "apricot" hair like some Poodles, dogs with a result of **Intermediate Red Pigmentation** will likely have tan or yellow hair like a Soft-Coated Wheaten Terrier, and dogs with **Dilute Red Pigmentation** will likely have cream or white hair like a Samoyed. Because the mutations we test may not directly cause differences in red pigmentation intensity, we consider this to be a linkage test.

No impact on coat pattern (Intermediate Red Pigmentation)

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^Yk^Y** 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^aa^a)

D Locus (MLPH)

The D locus result that we report is determined by three different genetic variants that can work together to cause diluted pigmentation. These are the common **d** allele, also known as "**d1**", and the less common alleles known as "**d2**" and "**d3**". Dogs with two **d** alleles, regardless of which variant, 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 in certain breeds, dilute dogs have a higher incidence of Color Dilution Alopecia. Dogs with one **d** allele will not be dilute, but can pass the **d** allele on to their puppies.

Dark areas of hair and skin are not lightened (Dd)

TRAITS: COAT COLOR (CONTINUED)

| TRAIT | RESULT |
|--|--|
| <p>Cocoa (HPS3)</p> <p>Dogs with the coco genotype will produce dark brown pigment instead of black in both their hair and skin. Dogs with the Nco genotype will produce black pigment, but can pass the co allele on to their puppies. Dogs that have the coco genotype as well as the bb genotype at the B locus are generally a lighter brown than dogs that have the Bb or BB genotypes at the B locus.</p> | <p>No co alleles, not expressed (NN)</p> |
| <p>B Locus (TYRP1)</p> <p>Dogs with two copies of the b allele produce brown pigment instead of black in both their hair and skin. Dogs with one copy of the b allele will produce black pigment, but can pass the b allele on to their puppies. E Locus ee dogs that carry two 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".</p> | <p>Black or gray hair and skin (Bb)</p> |
| <p>Saddle Tan (RALY)</p> <p>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 II 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 a^t allele, so dogs that do not express a^t are not influenced by this gene.</p> | <p>Not expressed (II)</p> |
| <p>S Locus (MITF)</p> <p>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.</p> | <p>Likely to have little to no white in coat (SS)</p> |

TRAITS: COAT COLOR (CONTINUED)

| TRAIT | RESULT |
|--|--|
| <p>M Locus (PMEL)</p> <p>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 M*m result are likely to be phenotypically merle or could be "non-expressing" merle, meaning that the merle pattern is very subtle or not at all evident in their coat. Dogs with an M*M* result are likely to be phenotypically merle or double merle. Dogs with an mm result have no merle alleles and are unlikely to have a merle coat pattern.</p> <p>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.</p> | <p>One merle allele; may express merle (M*m)</p> <p>Note: This locus includes several alleles. At the time this dog was genotyped Embark we could not distinguish all of the possible alleles.</p> |
| <p>R Locus (USH2A)</p> <p>The R Locus regulates the presence or absence of the roan coat color pattern. Partial duplication of the USH2A gene is strongly associated with this coat pattern. Dogs with at least one R allele will likely have roaning on otherwise uniformly unpigmented white areas. Roan appears in white areas controlled by the S Locus but not in other white or cream areas created by other loci, such as the E Locus with ee along with Dilute Red Pigmentation by I Locus (for example, in Samoyeds). Mechanisms for controlling the extent of roaning are currently unknown, and roaning can appear in a uniform or non-uniform pattern. Further, non-uniform roaning may appear as ticked, and not obviously roan. The roan pattern can appear with or without ticking.</p> | <p>Likely no impact on coat pattern (rr)</p> |
| <p>H Locus (Harlequin)</p> <p>This pattern is recognized in Great Danes and causes dogs to have a white coat with patches of darker pigment. A dog with an Hh result will be harlequin if they are also M*m or M*M* at the M Locus and are not ee at the E locus. Dogs with a result of hh will not be harlequin. This trait is thought to be homozygous lethal; a living dog with an HH genotype has never been found.</p> | <p>No harlequin alleles (hh)</p> |

TRAITS: COAT COLOR (CONTINUED)

| TRAIT | RESULT |
|-------|--------|
|-------|--------|

Panda White Spotting

Panda White Spotting originated in a line of German Shepherd Dogs and causes a mostly symmetrical white spotting of the head and/or body. This is a dominant variant of the KIT gene, which has a role in pigmentation.

Dogs with one copy of the I allele will exhibit this white spotting. Dogs with two copies of the I allele have never been observed, as two copies of the variant is suspected to be lethal to the developing embryo. Dogs with the **NN** result will not exhibit white spotting due to this variant.

Not expected to display Panda pattern (NN)

TRAITS: OTHER COAT TRAITS

| TRAIT | RESULT |
|-------|--------|
|-------|--------|

Furnishings (RSPO2)

Dogs with one or two copies of the **F** allele have “furnishings”: the mustache, beard, and eyebrows characteristic of breeds like the Schnauzer, Scottish Terrier, and Wire Haired Dachshund. A dog with two **I** 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.

Likely unfurnished (no mustache, beard, and/or eyebrows) (II)

TRAITS: OTHER COAT TRAITS (CONTINUED)

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|--------------|---------------|
| TRAIT | RESULT |
|--------------|---------------|

Coat Length (FGF5)

The FGF5 gene affects hair length in many species, including cats, dogs, mice, and humans. In dogs, an **Lh** allele confers a long, silky hair coat across many breeds, including Yorkshire Terriers, Cocker Spaniels, and Golden Retrievers, while the **Sh** allele causes a shorter coat, as seen in the Boxer or the American Staffordshire Terrier. In certain breeds, such as the Pembroke Welsh Corgi and French Bulldog, the long haircoat is described as "fluffy". The coat length determined by FGF5, as reported by us, is influenced by four genetic variants that work together to promote long hair.

The most common of these is the **Lh1** variant (G/T, CanFam3.1, chr32, g.4509367) and the less common ones are **Lh2** (C/T, CanFam3.1, chr32, g.4528639), **Lh3** (16bp deletion, CanFam3.1, chr32, g.4528616), and **Lh4** (GG insertion, CanFam3.1, chr32, g.4528621). The FGF5_Lh1 variant is found across many dog breeds. The less common alleles, FGF5_Lh2, have been found in the Akita, Samoyed, and Siberian Husky, FGF5_Lh3 have been found in the Eurasier, and FGF5_Lh4 have been found in the Afghan Hound, Eurasier, and French Bulldog.

Likely long coat (LhLh)

The **Lh** alleles have a recessive mode of inheritance, meaning that two copies of the **Lh** alleles are required to have long hair. The presence of two Lh alleles at any of these FGF5 loci is expected to result in long hair. One copy each of **Lh1** and **Lh2** have been found in Samoyeds, one copy each of **Lh1** and **Lh3** have been found in Eurasiers, and one copy each of **Lh1** and **Lh4** have been found in the Afghan Hounds and Eurasiers.

Interestingly, the Lh3 variant, a 16 base pair deletion, encompasses the Lh4 variant (GG insertion). The presence of one or two copies of Lh3 influences the outcome at the Lh4 locus. When two copies of Lh3 are present, there will be no reportable result for the FGF5_Lh4 locus. With one copy of Lh3, Lh4 can have either one copy of the variant allele or the normal allele. The overall FGF5 result remains unaffected by this.

TRAITS: OTHER COAT TRAITS (CONTINUED)

| TRAIT | RESULT |
|---|--|
| Shedding (MC5R) <p>Dogs with at least one copy of the ancestral C allele, like many Labradors and German Shepherd Dogs, are heavy or seasonal shedders, while those with two copies of the T allele, including many Boxers, Shih Tzus and Chihuahuas, tend to be lighter shedders. Dogs with furnished/wire-haired coats caused by RSPO2 (the furnishings gene) tend to be low shedders regardless of their genotype at this gene.</p> | Likely heavy/seasonal shedding (CC) |
| Coat Texture (KRT71) <p>Dogs with a long coat and at least one copy of the T allele have a wavy or curly coat characteristic of Poodles and Bichon Frises. Dogs with two copies of the ancestral C 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 F allele for the Furnishings (RSPO2) gene then they are likely to have a curly coat. Dogs with short coats may carry one or two copies of the T allele but still have straight coats.</p> | Likely straight coat (CC) |
| Hairlessness (FOXI3) <p>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 NDup genotype are likely to be hairless while dogs with the NN genotype are likely to have a normal coat. The DupDup 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.</p> | Very unlikely to be hairless (NN) |
| Hairlessness (SGK3) <p>Hairlessness in the American Hairless Terrier arises from a mutation in the SGK3 gene. Dogs with the DD result are likely to be hairless. Dogs with the ND genotype will have a normal coat, but can pass the D variant on to their offspring.</p> | Very unlikely to be hairless (NN) |

TRAITS: OTHER COAT TRAITS (CONTINUED)

| TRAIT | RESULT |
|---|-------------------------------|
| Oculocutaneous Albinism Type 2 (SLC45A2) | |
| <p>Dogs with two copies DD of this deletion in the SLC45A2 gene have oculocutaneous albinism (OCA), 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 ND 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.</p> | Likely not albino (NN) |

TRAITS: OTHER BODY FEATURES

| TRAIT | RESULT |
|-------|--------|
|-------|--------|

Muzzle Length (BMP3)

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 **C** allele. Dogs in many short-length muzzle (brachycephalic) breeds such as the English Bulldog, Pug, and Pekingese have two copies of the derived **A** 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.

Likely medium or long muzzle (CC)

Tail Length (T)

Whereas most dogs have two **C** alleles and a long tail, dogs with one **G** 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 **GG** genotypes have not been observed, suggesting that dogs with the **GG** 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.

Likely normal-length tail (CC)

Hind Dewclaws (LMBR1)

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 **T** 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 **CC** or **TC** dogs will have hind dewclaws.

Unlikely to have hind dew claws (CC)

TRAITS: OTHER BODY FEATURES (CONTINUED)

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|--------------|---------------|
| TRAIT | RESULT |
|--------------|---------------|

Chondrodysplasia (Chr. 18 FGF4 Retrogene)

Dogs with one or two copies of the **I** allele will exhibit a short-legged trait known as chondrodysplasia (CDPA). CDPA is a breed-defining characteristic of many breeds exhibiting the "short-legged, long-bodied" appearance known as disproportionate dwarfism, including the corgi, dachshund and basset hound. The impact of the **I** allele on leg length is additive. Therefore, dogs with the **II** result display the largest reduction in leg length. Dogs with the **NI** genotype will have an intermediate leg length, while dogs with the **NN** result will not exhibit leg shortening due to this variant. Breeds that display disproportionate dwarfism also frequently inherit a genetic variant known as the chondrodystrophy (CDDY) variant. The CDDY variant also shortens legs (in a less significant amount than CDPA) but, secondarily, increases the risk of Type I Intervertebral Disc Disease (IVDD). Test results for CDDY are listed in this dog's health testing results under "Intervertebral Disc Disease (Type I)". In contrast, the CDPA variant has NOT been shown to increase the risk of IVDD.

Not indicative of chondrodysplasia (normal leg length) (NN)

Blue Eye Color (ALX4)

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 (**Dup**) 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. **NN** 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.

Less likely to have blue eyes (NN)

Back Muscling & Bulk, Large Breed (ACSL4)

The **T** 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" **T** allele is absent from leaner shaped large breed dogs like the Great Dane, Irish Wolfhound, and Scottish Deerhound, which are fixed for the ancestral **C** 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.

Likely normal muscling (CC)

TRAITS: BODY SIZE

| TRAIT | RESULT |
|--|--------------------------|
| Body Size (IGF1) The I allele is associated with smaller body size. | Intermediate (NI) |
| Body Size (IGFR1) The A allele is associated with smaller body size. | Larger (GG) |
| Body Size (STC2) The A allele is associated with smaller body size. | Intermediate (TA) |
| Body Size (GHR - E191K) The A allele is associated with smaller body size. | Larger (GG) |
| Body Size (GHR - P177L) The T allele is associated with smaller body size. | Larger (CC) |

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)

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)

HEALTH REPORT

How to interpret Smooch's genetic health results:

If Smooch inherited any of the variants that we tested, they will be listed at the top of the Health Report section, along with a description of how to interpret this result. We also include all of the variants that we tested Smooch for that we did not detect the risk variant for.

A genetic test is not a diagnosis

This genetic test does not diagnose a disease. Please talk to your vet about your dog's genetic results, or if you think that your pet may have a health condition or disease.

Summary

Of the 262 genetic health risks we analyzed, we found 3 results that you should learn about.

Increased risk results (1)

Multiple Drug Sensitivity

Notable results (2)

ALT Activity

Degenerative Myelopathy, DM














Clear results

Breed-relevant (11)

Other (247)



















BREED-RELEVANT RESULTS

Research studies indicate that these results are more relevant to dogs like Smooch, and may influence her chances of developing certain health conditions.



















| | |
|--|----------------|
|  Multiple Drug Sensitivity (ABCB1) | Increased risk |
|  Degenerative Myelopathy, DM (SOD1A) | Notable |
|  Canine Multifocal Retinopathy, cmr1 (BEST1 Exon 2) | Clear |
|  Collie Eye Anomaly (NHEJ1) | Clear |
|  Craniomandibular Osteopathy, CMO (SLC37A2) | Clear |
|  Day Blindness (CNGB3 Deletion, Alaskan Malamute Variant) | Clear |
|  Hereditary Cataracts (HSF4 Exon 9, Australian Shepherd Variant) | Clear |
|  Junctional Epidermolysis Bullosa (LAMB3 Exon 11, Australian Shepherd Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 6, NCL 6 (CLN6 Exon 7, Australian Shepherd Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 8, NCL 8 (CLN8, Australian Shepherd Variant) | Clear |
|  Primary Ciliary Dyskinesia, PCD (STK36, Australian Shepherd Variant) | Clear |
|  Progressive Retinal Atrophy, prcd (PRCD Exon 1) | Clear |
|  Urate Kidney & Bladder Stones (SLC2A9) | Clear |

OTHER RESULTS



















Research has not yet linked these conditions to dogs with similar breeds to Smooch. Review any increased risk or notable results to understand her potential risk and recommendations.

| | |
|--|---------|
|  ALT Activity (GPT) | Notable |
|  2-DHA Kidney & Bladder Stones (APRT) | Clear |
|  Acral Mutilation Syndrome (GDNF-AS, Spaniel and Pointer Variant) | Clear |
|  Alaskan Husky Encephalopathy (SLC19A3) | Clear |
|  Alaskan Malamute Polyneuropathy, AMPN (NDRG1 SNP) | Clear |
|  Alexander Disease (GFAP) | Clear |
|  Anhidrotic Ectodermal Dysplasia (EDA Intron 8) | Clear |
|  Autosomal Dominant Progressive Retinal Atrophy (RHO) | Clear |
|  Bald Thigh Syndrome (IGFBP5) | Clear |
|  Bernard-Soulier Syndrome, BSS (GP9, Cocker Spaniel Variant) | Clear |
|  Bully Whippet Syndrome (MSTN) | Clear |
|  Canine Elliptocytosis (SPTB Exon 30) | Clear |
|  Canine Fucosidosis (FUCA1) | Clear |
|  Canine Leukocyte Adhesion Deficiency Type I, CLAD I (ITGB2, Setter Variant) | Clear |
|  Canine Leukocyte Adhesion Deficiency Type III, CLAD III (FERMT3, German Shepherd Variant) | Clear |
|  Canine Multifocal Retinopathy, cmr2 (BEST1 Exon 5, Coton de Tulear Variant) | Clear |
|  Canine Multifocal Retinopathy, cmr3 (BEST1 Exon 10 Deletion, Finnish and Swedish Lapphund, Lapponian Herder Variant) | Clear |
|  Canine Multiple System Degeneration (SERAC1 Exon 4, Chinese Crested Variant) | Clear |



















OTHER RESULTS

| | |
|--|-------|
|  Canine Multiple System Degeneration (SERAC1 Exon 15, Kerry Blue Terrier Variant) | Clear |
|  Cardiomyopathy and Juvenile Mortality (YARS2) | Clear |
|  Centronuclear Myopathy, CNM (PTPLA) | Clear |
|  Cerebellar Hypoplasia (VLDLR, Eurasier Variant) | Clear |
|  Chondrodystrophy (ITGA10, Norwegian Elkhound and Karelian Bear Dog Variant) | Clear |
|  Cleft Lip and/or Cleft Palate (ADAMTS20, Nova Scotia Duck Tolling Retriever Variant) | Clear |
|  Cleft Palate, CP1 (DLX6 intron 2, Nova Scotia Duck Tolling Retriever Variant) | Clear |
|  Cobalamin Malabsorption (CUBN Exon 8, Beagle Variant) | Clear |
|  Cobalamin Malabsorption (CUBN Exon 53, Border Collie Variant) | Clear |
|  Complement 3 Deficiency, C3 Deficiency (C3) | Clear |
|  Congenital Cornification Disorder (NSDHL, Chihuahua Variant) | Clear |
|  Congenital Hypothyroidism (TPO, Rat, Toy, Hairless Terrier Variant) | Clear |
|  Congenital Hypothyroidism (TPO, Tenterfield Terrier Variant) | Clear |
|  Congenital Hypothyroidism with Goiter (TPO Intron 13, French Bulldog Variant) | Clear |
|  Congenital Hypothyroidism with Goiter (SLC5A5, Shih Tzu Variant) | Clear |
|  Congenital Macrothrombocytopenia (TUBB1 Exon 1, Cairn and Norfolk Terrier Variant) | Clear |
|  Congenital Myasthenic Syndrome, CMS (COLQ, Labrador Retriever Variant) | Clear |
|  Congenital Myasthenic Syndrome, CMS (COLQ, Golden Retriever Variant) | Clear |



















OTHER RESULTS

| | |
|--|-------|
|  Congenital Myasthenic Syndrome, CMS (CHAT, Old Danish Pointing Dog Variant) | Clear |
|  Congenital Myasthenic Syndrome, CMS (CHRNE, Jack Russell Terrier Variant) | Clear |
|  Congenital Stationary Night Blindness (LRIT3, Beagle Variant) | Clear |
|  Congenital Stationary Night Blindness (RPE65, Briard Variant) | Clear |
|  Craniomandibular Osteopathy, CMO (SLC37A2 Intron 16, Basset Hound Variant) | Clear |
|  Cystinuria Type I-A (SLC3A1, Newfoundland Variant) | Clear |
|  Cystinuria Type II-A (SLC3A1, Australian Cattle Dog Variant) | Clear |
|  Cystinuria Type II-B (SLC7A9, Miniature Pinscher Variant) | Clear |
|  Darier Disease (ATP2A2, Irish Terrier Variant) | Clear |
|  Day Blindness (CNGA3 Exon 7, German Shepherd Variant) | Clear |
|  Day Blindness (CNGA3 Exon 7, Labrador Retriever Variant) | Clear |
|  Day Blindness (CNGB3 Exon 6, German Shorthaired Pointer Variant) | Clear |
|  Deafness and Vestibular Syndrome of Dobermans, DVDob, DINGS (MYO7A) | Clear |
|  Demyelinating Polyneuropathy (SBF2/MTRM13) | Clear |
|  Dental-Skeletal-Retinal Anomaly (MIA3, Cane Corso Variant) | Clear |
|  Diffuse Cystic Renal Dysplasia and Hepatic Fibrosis (INPP5E Intron 9, Norwich Terrier Variant) | Clear |
|  Dilated Cardiomyopathy, DCM (RBM20, Schnauzer Variant) | Clear |
|  Dilated Cardiomyopathy, DCM1 (PDK4, Doberman Pinscher Variant 1) | Clear |



















OTHER RESULTS

| | |
|---|-------|
|  Dilated Cardiomyopathy, DCM2 (TTN, Doberman Pinscher Variant 2) | Clear |
|  Disproportionate Dwarfism (PRKG2, Dogo Argentino Variant) | Clear |
|  Dry Eye Curly Coat Syndrome (FAM83H Exon 5) | Clear |
|  Dystrophic Epidermolysis Bullosa (COL7A1, Central Asian Shepherd Dog Variant) | Clear |
|  Dystrophic Epidermolysis Bullosa (COL7A1, Golden Retriever Variant) | Clear |
|  Early Bilateral Deafness (LOXHD1 Exon 38, Rottweiler Variant) | Clear |
|  Early Onset Adult Deafness, EOAD (EPS8L2 Deletion, Rhodesian Ridgeback Variant) | Clear |
|  Early Onset Cerebellar Ataxia (SEL1L, Finnish Hound Variant) | Clear |
|  Ehlers Danlos (ADAMTS2, Doberman Pinscher Variant) | Clear |
|  Ehlers-Danlos Syndrome (EDS) (COL5A1, Labrador Retriever Variant) | Clear |
|  Enamel Hypoplasia (ENAM Deletion, Italian Greyhound Variant) | Clear |
|  Enamel Hypoplasia (ENAM SNP, Parson Russell Terrier Variant) | Clear |
|  Episodic Falling Syndrome (BCAN) | Clear |
|  Exercise-Induced Collapse, EIC (DNM1) | Clear |
|  Factor VII Deficiency (F7 Exon 5) | Clear |
|  Factor XI Deficiency (F11 Exon 7, Kerry Blue Terrier Variant) | Clear |
|  Familial Nephropathy (COL4A4 Exon 3, Cocker Spaniel Variant) | Clear |
|  Familial Nephropathy (COL4A4 Exon 30, English Springer Spaniel Variant) | Clear |



















OTHER RESULTS

| | |
|--|-------|
|  Fanconi Syndrome (FAN1, Basenji Variant) | Clear |
|  Fetal-Onset Neonatal Neuroaxonal Dystrophy (MFN2, Giant Schnauzer Variant) | Clear |
|  Glanzmann's Thrombasthenia Type I (ITGA2B Exon 13, Great Pyrenees Variant) | Clear |
|  Glanzmann's Thrombasthenia Type I (ITGA2B Exon 12, Otterhound Variant) | Clear |
|  Globoid Cell Leukodystrophy, Krabbe disease (GALC Exon 5, Terrier Variant) | Clear |
|  Glycogen Storage Disease Type IA, Von Gierke Disease, GSD IA (G6PC, Maltese Variant) | Clear |
|  Glycogen Storage Disease Type IIIA, GSD IIIA (AGL, Curly Coated Retriever Variant) | Clear |
|  Glycogen storage disease Type VII, Phosphofructokinase Deficiency, PFK Deficiency (PFKM, Whippet and English Springer Spaniel Variant) | Clear |
|  Glycogen storage disease Type VII, Phosphofructokinase Deficiency, PFK Deficiency (PFKM, Wachtelhund Variant) | Clear |
|  GM1 Gangliosidosis (GLB1 Exon 2, Portuguese Water Dog Variant) | Clear |
|  GM1 Gangliosidosis (GLB1 Exon 15, Shiba Inu Variant) | Clear |
|  GM1 Gangliosidosis (GLB1 Exon 15, Alaskan Husky Variant) | Clear |
|  GM2 Gangliosidosis (HEXA, Japanese Chin Variant) | Clear |
|  GM2 Gangliosidosis (HEXB, Poodle Variant) | Clear |
|  Golden Retriever Progressive Retinal Atrophy 1, GR-PRA1 (SLC4A3) | Clear |
|  Golden Retriever Progressive Retinal Atrophy 2, GR-PRA2 (TTC8) | Clear |
|  Goniodysgenesis and Glaucoma, Pectinate Ligament Dysplasia, PLD (OLFM3) | Clear |
|  Hemophilia A (F8 Exon 11, German Shepherd Variant 1) | Clear |



















OTHER RESULTS

| | |
|--|-------|
|  Hemophilia A (F8 Exon 1, German Shepherd Variant 2) | Clear |
|  Hemophilia A (F8 Exon 10, Boxer Variant) | Clear |
|  Hemophilia B (F9 Exon 7, Terrier Variant) | Clear |
|  Hemophilia B (F9 Exon 7, Rhodesian Ridgeback Variant) | Clear |
|  Hereditary Ataxia, Cerebellar Degeneration (RAB24, Old English Sheepdog and Gordon Setter Variant) | Clear |
|  Hereditary Footpad Hyperkeratosis (FAM83G, Terrier and Kromfohrlander Variant) | Clear |
|  Hereditary Footpad Hyperkeratosis (DSG1, Rottweiler Variant) | Clear |
|  Hereditary Nasal Parakeratosis (SUV39H2 Intron 4, Greyhound Variant) | Clear |
|  Hereditary Nasal Parakeratosis, HNPk (SUV39H2) | Clear |
|  Hereditary Vitamin D-Resistant Rickets (VDR) | Clear |
|  Hypocatalasia, Acatlasemia (CAT) | Clear |
|  Hypomyelination and Tremors (FNIP2, Weimaraner Variant) | Clear |
|  Hypophosphatasia (ALPL Exon 9, Karelian Bear Dog Variant) | Clear |
|  Ichthyosis (NIPAL4, American Bulldog Variant) | Clear |
|  Ichthyosis (ASPRV1 Exon 2, German Shepherd Variant) | Clear |
|  Ichthyosis (SLC27A4, Great Dane Variant) | Clear |
|  Ichthyosis, Epidermolytic Hyperkeratosis (KRT10, Terrier Variant) | Clear |
|  Ichthyosis, ICH1 (PNPLA1, Golden Retriever Variant) | Clear |



















OTHER RESULTS

| | |
|--|-------|
|  Ichthyosis, ICH2 (ABHD5, Golden Retriever Variant) | Clear |
|  Inflammatory Myopathy (SLC25A12) | Clear |
|  Inherited Myopathy of Great Danes (BIN1) | Clear |
|  Inherited Selected Cobalamin Malabsorption with Proteinuria (CUBN, Komondor Variant) | Clear |
|  Intervertebral Disc Disease (Type I) (FGF4 retrogene - CFA12) | Clear |
|  Intestinal Lipid Malabsorption (ACSL5, Australian Kelpie) | Clear |
|  Junctional Epidermolysis Bullosa (LAMA3 Exon 66, Australian Cattle Dog Variant) | Clear |
|  Juvenile Epilepsy (LGI2) | Clear |
|  Juvenile Laryngeal Paralysis and Polyneuropathy (RAB3GAP1, Rottweiler Variant) | Clear |
|  Juvenile Myoclonic Epilepsy (DIRAS1) | Clear |
|  L-2-Hydroxyglutaricaciduria, L2HGA (L2HGDH, Staffordshire Bull Terrier Variant) | Clear |
|  Lagotto Storage Disease (ATG4D) | Clear |
|  Laryngeal Paralysis (RAPGEF6, Miniature Bull Terrier Variant) | Clear |
|  Late Onset Spinocerebellar Ataxia (CAPN1) | Clear |
|  Late-Onset Neuronal Ceroid Lipofuscinosis, NCL 12 (ATP13A2, Australian Cattle Dog Variant) | Clear |
|  Leonberger Polyneuropathy 1 (LPN1, ARHGEF10) | Clear |
|  Leonberger Polyneuropathy 2 (GJA9) | Clear |
|  Lethal Acrodermatitis, LAD (MKLN1) | Clear |



















OTHER RESULTS

| | |
|---|-------|
|  Leukodystrophy (TSEN54 Exon 5, Standard Schnauzer Variant) | Clear |
|  Ligneous Membranitis, LM (PLG) | Clear |
|  Limb Girdle Muscular Dystrophy (SGCD, Boston Terrier Variant) | Clear |
|  Limb-Girdle Muscular Dystrophy 2D (SGCA Exon 3, Miniature Dachshund Variant) | Clear |
|  Long QT Syndrome (KCNQ1) | Clear |
|  Lundehund Syndrome (LEPREL1) | Clear |
|  Macular Corneal Dystrophy, MCD (CHST6) | Clear |
|  Malignant Hyperthermia (RYR1) | Clear |
|  May-Hegglin Anomaly (MYH9) | Clear |
|  Medium-Chain Acyl-CoA Dehydrogenase Deficiency, MCADD (ACADM, Cavalier King Charles Spaniel Variant) | Clear |
|  Methemoglobinemia (CYB5R3, Pit Bull Terrier Variant) | Clear |
|  Methemoglobinemia (CYB5R3) | Clear |
|  Microphthalmia (RBP4 Exon 2, Soft Coated Wheaten Terrier Variant) | Clear |
|  Mucopolysaccharidosis IIIB, Sanfilippo Syndrome Type B, MPS IIIB (NAGLU, Schipperke Variant) | Clear |
|  Mucopolysaccharidosis Type IIIA, Sanfilippo Syndrome Type A, MPS IIIA (SGSH Exon 6, Dachshund Variant) | Clear |
|  Mucopolysaccharidosis Type IIIA, Sanfilippo Syndrome Type A, MPS IIIA (SGSH Exon 6, New Zealand Huntaway Variant) | Clear |
|  Mucopolysaccharidosis Type VI, Maroteaux-Lamy Syndrome, MPS VI (ARSB Exon 5, Miniature Pinscher Variant) | Clear |
|  Mucopolysaccharidosis Type VII, Sly Syndrome, MPS VII (GUSB Exon 3, German Shepherd Variant) | Clear |



















OTHER RESULTS

| | |
|---|-------|
|  Mucopolysaccharidosis Type VII, Sly Syndrome, MPS VII (GUSB Exon 5, Terrier Brasileiro Variant) | Clear |
|  Muscular Dystrophy (DMD, Cavalier King Charles Spaniel Variant 1) | Clear |
|  Muscular Dystrophy (DMD, Golden Retriever Variant) | Clear |
|  Musladin-Lueke Syndrome, MLS (ADAMTSL2) | Clear |
|  Myasthenia Gravis-Like Syndrome (CHRNE, Heideterrier Variant) | Clear |
|  Myotonia Congenita (CLCN1 Exon 23, Australian Cattle Dog Variant) | Clear |
|  Myotonia Congenita (CLCN1 Exon 19, Labrador Retriever Variant) | Clear |
|  Myotonia Congenita (CLCN1 Exon 7, Miniature Schnauzer Variant) | Clear |
|  Narcolepsy (HCRTR2 Exon 1, Dachshund Variant) | Clear |
|  Narcolepsy (HCRTR2 Intron 4, Doberman Pinscher Variant) | Clear |
|  Narcolepsy (HCRTR2 Intron 6, Labrador Retriever Variant) | Clear |
|  Nemaline Myopathy (NEB, American Bulldog Variant) | Clear |
|  Neonatal Cerebellar Cortical Degeneration (SPTBN2, Beagle Variant) | Clear |
|  Neonatal Encephalopathy with Seizures, NEWS (ATF2) | Clear |
|  Neonatal Interstitial Lung Disease (LAMP3) | Clear |
|  Neuroaxonal Dystrophy, NAD (VPS11, Rottweiler Variant) | Clear |
|  Neuroaxonal Dystrophy, NAD (TECPR2, Spanish Water Dog Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 1, NCL 1 (PPT1 Exon 8, Dachshund Variant 1) | Clear |



















OTHER RESULTS

| | |
|--|-------|
|  Neuronal Ceroid Lipofuscinosis 10, NCL 10 (CTSD Exon 5, American Bulldog Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 2, NCL 2 (TPP1 Exon 4, Dachshund Variant 2) | Clear |
|  Neuronal Ceroid Lipofuscinosis 5, NCL 5 (CLN5 Exon 4 SNP, Border Collie Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 5, NCL 5 (CLN5 Exon 4 Deletion, Golden Retriever Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 7, NCL 7 (MFSD8, Chihuahua and Chinese Crested Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 8, NCL 8 (CLN8 Exon 2, English Setter Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis 8, NCL 8 (CLN8 Insertion, Saluki Variant) | Clear |
|  Neuronal Ceroid Lipofuscinosis, Cerebellar Ataxia, NCL4A (ARSG Exon 2, American Staffordshire Terrier Variant) | Clear |
|  Oculocutaneous Albinism, OCA (SLC45A2 Exon 6, Bullmastiff Variant) | Clear |
|  Oculocutaneous Albinism, OCA (SLC45A2, Small Breed Variant) | Clear |
|  Oculoskeletal Dysplasia 2 (COL9A2, Samoyed Variant) | Clear |
|  Osteochondrodysplasia (SLC13A1, Poodle Variant) | Clear |
|  Osteogenesis Imperfecta (COL1A2, Beagle Variant) | Clear |
|  Osteogenesis Imperfecta (SERPINH1, Dachshund Variant) | Clear |
|  Osteogenesis Imperfecta (COL1A1, Golden Retriever Variant) | Clear |
|  P2Y12 Receptor Platelet Disorder (P2Y12) | Clear |
|  Pachyonychia Congenita (KRT16, Dogue de Bordeaux Variant) | Clear |
|  Paroxysmal Dyskinesia, PxD (PIGN) | Clear |



















OTHER RESULTS

| | |
|---|-------|
|  Persistent Mullerian Duct Syndrome, PMDS (AMHR2) | Clear |
|  Pituitary Dwarfism (POU1F1 Intron 4, Karelian Bear Dog Variant) | Clear |
|  Platelet Factor X Receptor Deficiency, Scott Syndrome (TMEM16F) | Clear |
|  Polycystic Kidney Disease, PKD (PKD1) | Clear |
|  Pompe's Disease (GAA, Finnish and Swedish Lapphund, Lapponian Herder Variant) | Clear |
|  Prekallikrein Deficiency (KLKB1 Exon 8) | Clear |
|  Primary Ciliary Dyskinesia, PCD (NME5, Alaskan Malamute Variant) | Clear |
|  Primary Ciliary Dyskinesia, PCD (CCDC39 Exon 3, Old English Sheepdog Variant) | Clear |
|  Primary Hyperoxaluria (AGXT) | Clear |
|  Primary Lens Luxation (ADAMTS17) | Clear |
|  Primary Open Angle Glaucoma (ADAMTS17 Exon 11, Basset Fauve de Bretagne Variant) | Clear |
|  Primary Open Angle Glaucoma (ADAMTS10 Exon 17, Beagle Variant) | Clear |
|  Primary Open Angle Glaucoma (ADAMTS10 Exon 9, Norwegian Elkhound Variant) | Clear |
|  Primary Open Angle Glaucoma and Primary Lens Luxation (ADAMTS17 Exon 2, Chinese Shar-Pei Variant) | Clear |
|  Progressive Retinal Atrophy (SAG) | Clear |
|  Progressive Retinal Atrophy (IFT122 Exon 26, Lapponian Herder Variant) | Clear |
|  Progressive Retinal Atrophy, Bardet-Biedl Syndrome (BBS2 Exon 11, Shetland Sheepdog Variant) | Clear |
|  Progressive Retinal Atrophy, CNGA (CNGA1 Exon 9) | Clear |















OTHER RESULTS

| | |
|--|-------|
|  Progressive Retinal Atrophy, crd1 (PDE6B, American Staffordshire Terrier Variant) | Clear |
|  Progressive Retinal Atrophy, crd4/cord1 (RPGRIP1) | Clear |
|  Progressive Retinal Atrophy, PRA1 (CNGB1) | Clear |
|  Progressive Retinal Atrophy, PRA3 (FAM161A) | Clear |
|  Progressive Retinal Atrophy, rcd1 (PDE6B Exon 21, Irish Setter Variant) | Clear |
|  Progressive Retinal Atrophy, rcd3 (PDE6A) | Clear |
|  Proportionate Dwarfism (GH1 Exon 5, Chihuahua Variant) | Clear |
|  Protein Losing Nephropathy, PLN (NPHS1) | Clear |
|  Pyruvate Dehydrogenase Deficiency (PDP1, Spaniel Variant) | Clear |
|  Pyruvate Kinase Deficiency (PKLR Exon 5, Basenji Variant) | Clear |
|  Pyruvate Kinase Deficiency (PKLR Exon 7, Beagle Variant) | Clear |
|  Pyruvate Kinase Deficiency (PKLR Exon 10, Terrier Variant) | Clear |
|  Pyruvate Kinase Deficiency (PKLR Exon 7, Labrador Retriever Variant) | Clear |
|  Pyruvate Kinase Deficiency (PKLR Exon 7, Pug Variant) | Clear |
|  Raine Syndrome (FAM20C) | Clear |
|  Recurrent Inflammatory Pulmonary Disease, RIPD (AKNA, Rough Collie Variant) | Clear |
|  Renal Cystadenocarcinoma and Nodular Dermatofibrosis (FLCN Exon 7) | Clear |
|  Retina Dysplasia and/or Optic Nerve Hypoplasia (SIX6 Exon 1, Golden Retriever Variant) | Clear |

OTHER RESULTS

| | |
|---|-------|
|  Sensory Neuropathy (FAM134B, Border Collie Variant) | Clear |
|  Severe Combined Immunodeficiency, SCID (PRKDC, Terrier Variant) | Clear |
|  Severe Combined Immunodeficiency, SCID (RAG1, Wetterhoun Variant) | Clear |
|  Shaking Puppy Syndrome (PLP1, English Springer Spaniel Variant) | Clear |
|  Shar-Pei Autoinflammatory Disease, SPAID, Shar-Pei Fever (MTBP) | Clear |
|  Skeletal Dysplasia 2, SD2 (COL11A2, Labrador Retriever Variant) | Clear |
|  Skin Fragility Syndrome (PKP1, Chesapeake Bay Retriever Variant) | Clear |
|  Spinocerebellar Ataxia (SCN8A, Alpine Dachsbracke Variant) | Clear |
|  Spinocerebellar Ataxia with Myokymia and/or Seizures (KCNJ10) | Clear |
|  Spongy Degeneration with Cerebellar Ataxia 1 (KCNJ10) | Clear |
|  Spongy Degeneration with Cerebellar Ataxia 2 (ATP1B2) | Clear |
|  Stargardt Disease (ABCA4 Exon 28, Labrador Retriever Variant) | Clear |
|  Succinic Semialdehyde Dehydrogenase Deficiency (ALDH5A1 Exon 7, Saluki Variant) | Clear |
|  Thrombopathia (RASGRP1 Exon 5, American Eskimo Dog Variant) | Clear |
|  Thrombopathia (RASGRP1 Exon 5, Basset Hound Variant) | Clear |
|  Thrombopathia (RASGRP1 Exon 8, Landseer Variant) | Clear |
|  Trapped Neutrophil Syndrome, TNS (VPS13B) | Clear |
|  Ullrich-like Congenital Muscular Dystrophy (COL6A3 Exon 10, Labrador Retriever Variant) | Clear |

OTHER RESULTS

| | |
|--|-----------|
|  Ullrich-like Congenital Muscular Dystrophy (COL6A1 Exon 3, Landseer Variant) | Clear |
|  Unilateral Deafness and Vestibular Syndrome (PTPRQ Exon 39, Doberman Pinscher) | Clear |
|  Von Willebrand Disease Type I, Type I vWD (VWF) | Clear |
|  Von Willebrand Disease Type II, Type II vWD (VWF, Pointer Variant) | Clear |
|  Von Willebrand Disease Type III, Type III vWD (VWF Exon 4, Terrier Variant) | Clear |
|  Von Willebrand Disease Type III, Type III vWD (VWF Intron 16, Nederlandse Kooikerhondje Variant) | Clear |
|  Von Willebrand Disease Type III, Type III vWD (VWF Exon 7, Shetland Sheepdog Variant) | Clear |
|  X-Linked Hereditary Nephropathy, XLHN (COL4A5 Exon 35, Samoyed Variant 2) | Clear |
|  X-Linked Myotubular Myopathy (MTM1, Labrador Retriever Variant) | Clear |
|  X-Linked Progressive Retinal Atrophy 1, XL-PRA1 (RPGR) | Clear |
|  X-linked Severe Combined Immunodeficiency, X-SCID (IL2RG Exon 1, Basset Hound Variant) | Clear |
|  X-linked Severe Combined Immunodeficiency, X-SCID (IL2RG, Corgi Variant) | Clear |
|  Xanthine Urolithiasis (XDH, Mixed Breed Variant) | Clear |
|  β -Mannosidosis (MANBA Exon 16, Mixed-Breed Variant) | Clear |
| Mast Cell Tumor | No result |

HEALTH REPORT

Increased risk result

Multiple Drug Sensitivity

Smooch inherited one copy of the variant we tested for MDR1 Drug Sensitivity
Smooch is at increased risk for MDR1

How to interpret this result

Smooch has one copy of a variant at the ABCB1 gene and is at risk for displaying adverse drug reactions. While she may not be as severely affected as a dog with two copies of the ABCB1 drug sensitivity allele, normal dosages of drugs could still have potentially severe effects on Smooch. Please inform your veterinarian that Smooch carries this variant; it is essential that they know this information before prescribing drugs.

What is MDR1 Drug Sensitivity?

Sensitivity to certain classes of drugs, notably the parasiticide ivermectin, as well as certain gastroprotectant and anti-cancer medications, occurs in dogs with a mutation in the ABCB1 gene.

When signs & symptoms develop in affected dogs

Symptoms arise after a dog has received an MDR1 problem drug or dosage, and can range from vomiting and diarrhea to lethargy, seizures, or coma.

Signs & symptoms

MDR1 often presents in young adulthood, only because this is most commonly when a dog is first exposed to a problem drug like high dose ivermectin or acepromazine.

How vets diagnose this condition

This is usually a retroactive diagnosis after a dog has an adverse reaction to a problem drug--however, genetic testing could help you avoid a first reaction altogether.

How this condition is treated

MDR1 is perfectly avoidable simply by avoiding the problem drugs, or problem dosages.

Actions to take if your dog is affected

- Review the MDR1 Problem Drug List as described by Washington State University and notify your veterinarian to flag this in your dog's file!
- Farm dogs with MDR1 may also benefit if they are either kept away from herds where ivermectin is used as a routine antiparasitic, or if another form of antiparasitic is used in areas that they are working.

HEALTH REPORT

Notable result

ALT Activity

Smooch inherited both copies of the variant we tested for Alanine Aminotransferase Activity

Why is this important to your vet?

Smooch has two copies of a variant in the GPT gene and is likely to have a lower than average baseline ALT activity. ALT is a commonly used measure of liver health on routine veterinary blood chemistry panels. As such, your veterinarian may want to watch for changes in Smooch's ALT activity above their current, healthy, ALT activity. As an increase above Smooch's baseline 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.

HEALTH REPORT

Notable result

Degenerative Myelopathy, DM

Smooch inherited one copy of the variant we tested for Degenerative Myelopathy, DM

What does this result mean?

This variant should not impact Smooch's health. This variant is inherited in an autosomal recessive manner, meaning that a dog needs two copies of the variant to show signs of this condition. Smooch is unlikely to develop this condition due to this variant because she only has one copy of the variant.

Impact on Breeding

Your dog carries this variant and will pass it on to ~50% of her offspring. You can email breeders@embarkvet.com to discuss with a genetic counselor how the genotype results should be applied to a breeding program.

What is Degenerative Myelopathy, DM?

The dog equivalent of Amyotrophic Lateral Sclerosis, or Lou Gehrig's disease, DM is a progressive degenerative disorder of the spinal cord. Because the nerves that control the hind limbs are the first to degenerate, the most common clinical signs are back muscle wasting and gait abnormalities.

When signs & symptoms develop in affected dogs

Affected dogs do not usually show signs of DM until they are at least 8 years old.

How vets diagnose this condition

Definitive diagnosis requires microscopic analysis of the spinal cord after death. However, veterinarians use clues such as genetic testing, breed, age, and other diagnostics to determine if DM is the most likely cause of your dog's clinical signs.

How this condition is treated

As dogs are seniors at the time of onset, the treatment for DM is aimed towards increasing their comfort through a combination of lifestyle changes, medication, and physical therapy.

Actions to take if your dog is affected

- Giving your dog the best quality of life for as long as possible is all you can do after receiving this diagnosis.

INBREEDING AND DIVERSITY

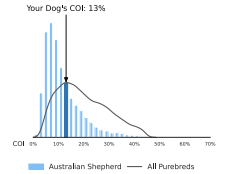
CATEGORY

RESULT

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.

13%

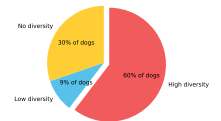


MHC Class II - DLA DRB1

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:

