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BREED ANCESTRY

Poodle (Standard) : 50.0%
Golden Retriever : 50.0%

GENETIC STATS

Predicted adult weight: 48 lbs

TEST DETAILS

Kit number: EM-19755908 Swab number: 31220412302682

BREED ANCESTRY BY CHROMOSOME

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

			Breed	colors:			
		Pood	le (Standard)	Golden F	Retriever		
1		2		3		4	
5		6		7		8	
9	_	10		11		12	
13		14		15		16	
17		18		19		20	
21		22		23		24	
25	_	26		27		28	
29		30		31		32	
33	_	34		35	=	36	
37	_	38	_				





Fun Fact

From 1989 to 1991, John Suter raced a team of Poodles in the Iditarod. Although his teams placed in the back half of the pack, he managed to win \$2,000 in prize money before retiring his poodle team. The Iditarod has since changed its rules to specify that only northern dog breeds can compete. Test Date: May 16th, 2023



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POODLE (STANDARD)

The Standard Poodle is a popular, water-loving dog used for centuries as a bird dog and popular pet. Poodles were established in Germany by the 15th century. Oddly enough, they are the national dog breed of France, and they were the most popular breed of dog in the United States throughout the 1960s and 70s. They're still quite popular today, owing to their intelligence, trainability, and non-shedding coats. Although well-known for their fancy fur, they're one of the most intelligent breeds of dog and require a lot of exercise and stimulation.







Fun Fact

A Golden Retriever is also pictured in the Guinness Book of World's Records for "Most tennis balls held in mouth" (with 6). Test Date: May 16th, 2023



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GOLDEN RETRIEVER

The Golden Retriever was developed in the early 19th century as an ideal hunting companion, able to retrieve birds on both land and water in the marshy Scottish countryside. Their friendliness and intelligence makes the both a popular family pet and an excellent working dog, well suited for being a service dog, therapy dog or for search and rescue. The third most popular breed in the US, the American and Canadian Goldens are generally lankier and darker than their British counterparts. Their wavy, feathered topcoat is water resistant, their undercoat helps them with thermoregulation and both coats have a tendency for heavy seasonal shedding. Goldens need lots of exercise (especially when younger), and their love of play and water means their owners usually get a lot of exercise too! In 2013, the 100th anniversary of Britain's Golden Retriever Club, Goldens from around the world came made the pilgrimage to the breed's birthplace in Scotland, where 222 of them posed in a single record-breaking photo. At the same time, the Golden Retriever Lifetime Study was getting started in the United States, recruiting 3,000 Golden Retrievers for a lifetime study aimed at understanding how genetics, lifestyle and environment influences healthy aging and cancer risk in Goldens.







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MATERNAL LINE



Through Jingles'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: A1a

A1a is the most common maternal lineage among Western dogs. This lineage traveled from the site of dog domestication in Central Asia to Europe along with an early dog expansion perhaps 10,000 years ago. It hung around in European village dogs for many millennia. Then, about 300 years ago, some of the prized females in the line were chosen as the founding dogs for several dog breeds. That set in motion a huge expansion of this lineage. It's now the maternal lineage of the overwhelming majority of Mastiffs, Labrador Retrievers and Gordon Setters. About half of Boxers and less than half of Shar-Pei dogs descend from the A1a line. It is also common across the world among village dogs, a legacy of European colonialism.

HAPLOTYPE: A261

Part of the large A1a haplogroup, this haplotype occurs in village dogs in Peru. Among breeds, it is most common in Golden Retrievers, Gordon Setters, and Labrador Retrievers.





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RESULT

TRAITS: COAT COLOR

TRAIT

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.

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**^y**k**^y genotype will show a coat color pattern based on the genotype they have at the A Locus. Dogs who test as **K**^B**k**^y may be brindle rather than black or brown.

No dark hairs anywhere (ee)

Not expressed (K^BK^B)







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RESULT

TRAITS: COAT COLOR (CONTINUED)

TRAIT

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.

Any pigmented hair likely apricot or red (Intense 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**^y**k**^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.

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.

Not expressed (a^ta)

Not expressed (DD)







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RESULT

TRAITS: COAT COLOR (CONTINUED)

TRAIT

Cocoa (HPS3)

Dogs with the coco genotype will produce dark brown pigment instead of black in both their hair and skin.No coDogs with the Nco genotype will produce black pigment, but can pass the co allele on to their puppies.expreDogs that have the coco genotype as well as the bb genotype at the B locus are generally a lighter brownthan dogs that have the Bb or BB genotypes at the B locus.

No co alleles, not expressed (NN)

B Locus (TYRP1)

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".

Likely black colored nose/feet (BB)

Saddle Tan (RALY)

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.

Not expressed (NN)

S Locus (MITF)

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.

Likely solid colored, but may have small amounts of white (Ssp)







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RESULT

TRAITS: COAT COLOR (CONTINUED)

TRAIT

M Locus (PMEL)

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. Dogs with an **mm** 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.

R Locus (USH2A)

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.

Likely no impact on coat pattern (rr)

No merle alleles (mm)

H Locus (Harlequin)

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.

No harlequin alleles (hh)







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RESULT

TRAITS: OTHER COAT TRAITS

TRAIT

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 furnished (mustache, beard, and/or eyebrows) (FI)







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TRAITS: OTHER COAT TRAITS (CONTINUED)

TRAIT

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.

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.

RESULT

Likely long coat (LhLh)







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RESULT

TRAITS: OTHER COAT TRAITS (CONTINUED)

TRAIT

Shedding (MC5R)

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.Likely

Likely light shedding (CC)

Coat Texture (KRT71)

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.

Hairlessness (FOXI3)

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.

Hairlessness (SGK3)

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.

Very unlikely to be hairless (NN)







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TRAITS: OTHER COAT TRAITS (CONTINUED)

TRAIT

RESULT

Oculocutaneous Albinism Type 2 (SLC45A2)

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.

Likely not albino (NN)







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RESULT

TRAITS: OTHER BODY FEATURES

TRAIT

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.

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.

Likely normal-length tail (CC)

Unlikely to have hind dew claws (CC)







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RESULT

TRAITS: OTHER BODY FEATURES (CONTINUED)

TRAIT

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" largebreed 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)





DNA Test Report	Test Date: May 16th, 2023	embk.me/jingles29
TRAITS: BODY SIZE		
TRAIT		RESULT
Body Size (IGF1)		Smaller (II)
The I allele is associated with smaller body size.		Smaller (II)
Body Size (IGFR1)		Larger (GG)
The A allele is associated with smaller body size.		
Body Size (STC2)		Larger (TT)
The A allele is associated with smaller body size.		
Body Size (GHR - E191K)		Intermediate (GA)
The A allele is associated with smaller body size.		internediate (SA)
Body Size (GHR - P177L)		Larger (CC)
The ${\bf T}$ allele is associated with smaller body size.		





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RESULT

TRAITS: PERFORMANCE

TRAIT

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.

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 motivation (NN) 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.





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HEALTH REPORT

How to interpret Jingles's genetic health results:

If Jingles 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 Jingles 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

Jingles is not at increased risk for the genetic health conditions that Embark tests.

Clear results

Breed-relevant (16)

Other (239)







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BREED-RELEVANT RESULTS

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

Congenital Myasthenic Syndrome, CMS (COLQ, Golden Retriever Variant)	Clear
O Degenerative Myelopathy, DM (SOD1A)	Clear
Opstrophic Epidermolysis Bullosa (COL7A1, Golden Retriever 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
Ichthyosis, ICH1 (PNPLA1, Golden Retriever Variant)	Clear
Intervertebral Disc Disease (Type I) (FGF4 retrogene - CFA12)	Clear
Muscular Dystrophy (DMD, Golden Retriever Variant)	Clear
Neonatal Encephalopathy with Seizures, NEWS (ATF2)	Clear
Neuronal Ceroid Lipofuscinosis 5, NCL 5 (CLN5 Exon 4 Deletion, Golden Retriever Variant)	Clear
Osteochondrodysplasia (SLC13A1, Poodle Variant)	Clear
Osteogenesis Imperfecta (COL1A1, Golden Retriever Variant)	Clear
Progressive Retinal Atrophy, prcd (PRCD Exon 1)	Clear
Retina Dysplasia and/or Optic Nerve Hypoplasia (SIX6 Exon 1, Golden Retriever Variant)	Clear
⊘ Von Willebrand Disease Type I, Type I vWD (VWF)	Clear





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OTHER RESULTS

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

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
ALT Activity (GPT)	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, cmr1 (BEST1 Exon 2)	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



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OTHER RESULTS

Canine Multiple System Degeneration (SERAC1 Exon 4, Chinese Crested Variant)	Clear
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
Collie Eye Anomaly (NHEJ1)	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 (TPO, Tenterfield Terrier Variant) Congenital Hypothyroidism with Goiter (TPO Intron 13, French Bulldog Variant) 	Clear Clear



DNA Test Report	Test Date: May 16th, 2023	embk.me/jingles29
OTHER RESULTS		
Congenital Myasthenic Syndrom	me, CMS (COLQ, Labrador Retriever Variant)	Clear
Ocongenital Myasthenic Syndrom	me, CMS (CHAT, Old Danish Pointing Dog Variant)	Clear
Congenital Myasthenic Syndrom	me, CMS (CHRNE, Jack Russell Terrier Variant)	Clear
Congenital Stationary Night Bli	ndness (LRIT3, Beagle Variant)	Clear
Congenital Stationary Night Bli	ndness (RPE65, Briard Variant)	Clear
🔗 Craniomandibular Osteopathy,	CMO (SLC37A2)	Clear
Craniomandibular Osteopathy,	CMO (SLC37A2 Intron 16, Basset Hound Variant)	Clear
🔗 Cystinuria Type I-A (SLC3A1, Ne	ewfoundland Variant)	Clear
🔗 Cystinuria Type II-A (SLC3A1, A	ustralian Cattle Dog Variant)	Clear
🔗 Cystinuria Type II-B (SLC7A9, N	Iiniature Pinscher Variant)	Clear
O Day Blindness (CNGB3 Deletion	n, Alaskan Malamute Variant)	Clear
Oay Blindness (CNGA3 Exon 7, 0	German Shepherd Variant)	Clear
Oay Blindness (CNGA3 Exon 7, I	Labrador Retriever Variant)	Clear
Oay Blindness (CNGB3 Exon 6,	German Shorthaired Pointer Variant)	Clear
O Deafness and Vestibular Syndro	ome of Dobermans, DVDob, DINGS (MYO7A)	Clear
Oemyelinating Polyneuropathy	(SBF2/MTRM13)	Clear
O Dental-Skeletal-Retinal Anoma	ly (MIA3, Cane Corso Variant)	Clear
Diffuse Cystic Renal Dysplasia	and Hepatic Fibrosis (INPP5E Intron 9, Norwich Terrier Variant)	Clear



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OTHER RESULTS		
Dilated Cardiomyopathy, DCM	I (RBM20, Schnauzer Variant)	Clear
Oilated Cardiomyopathy, DCM	11 (PDK4, Doberman Pinscher Variant 1)	Clear
Oilated Cardiomyopathy, DCM	12 (TTN, Doberman Pinscher Variant 2)	Clear
Disproportionate Dwarfism (P	PRKG2, Dogo Argentino Variant)	Clear
Ory Eye Curly Coat Syndrome	(FAM83H Exon 5)	Clear
Oystrophic Epidermolysis Bul	losa (COL7A1, Central Asian Shepherd Dog Variant)	Clear
Sarly Bilateral Deafness (LOX	HD1 Exon 38, Rottweiler Variant)	Clear
🔗 Early Onset Adult Deafness, E	OAD (EPS8L2 Deletion, Rhodesian Ridgeback Variant)	Clear
🔗 Early Onset Cerebellar Ataxia	(SEL1L, Finnish Hound Variant)	Clear
Ehlers Danlos (ADAMTS2, Dob	berman Pinscher Variant)	Clear
🔗 Enamel Hypoplasia (ENAM De	eletion, Italian Greyhound Variant)	Clear
🔗 Enamel Hypoplasia (ENAM SN	IP, Parson Russell Terrier Variant)	Clear
Episodic Falling Syndrome (B	CAN)	Clear
Exercise-Induced Collapse, E	IC (DNM1)	Clear
Sactor VII Deficiency (F7 Exor	n 5)	Clear
Sactor XI Deficiency (F11 Exor	n 7, Kerry Blue Terrier Variant)	Clear
Samilial Nephropathy (COL4A	4 Exon 3, Cocker Spaniel Variant)	Clear
Samilial Nephropathy (COL4A	4 Exon 30, English Springer Spaniel Variant)	Clear

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OTHER RESULTS

Sanconi Syndrome (FAN1, Basenji Variant)	Clear
Setal-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
Goniodysgenesis and Glaucoma, Pectinate Ligament Dysplasia, PLD (OLFM3)	Clear
Hemophilia A (F8 Exon 11, German Shepherd Variant 1)	Clear
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



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OTHER RESULTS

Hemophilia B (F9 Exon 7, Rhodesian Ridgeback Variant)	Clear
Hereditary Ataxia, Cerebellar Degeneration (RAB24, Old English Sheepdog and Gordon Setter Variant)	Clear
Hereditary Cataracts (HSF4 Exon 9, Australian Shepherd 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, Acatalasemia (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
Inflammatory Myopathy (SLC25A12)	Clear
Inherited Myopathy of Great Danes (BIN1)	Clear
Inherited Selected Cobalamin Malabsorption with Proteinuria (CUBN, Komondor Variant)	Clear



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OTHER RESULTS		
🔗 Intestinal Lipid Malabsorptic	on (ACSL5, Australian Kelpie)	Clear
🧭 Junctional Epidermolysis Bu	llosa (LAMA3 Exon 66, Australian Cattle Dog Variant)	Clear
🧭 Junctional Epidermolysis Bu	llosa (LAMB3 Exon 11, Australian Shepherd 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 (AT	G4D)	Clear
🔗 Laryngeal Paralysis (RAPGEF	F6, Miniature Bull Terrier Variant)	Clear
Zate Onset Spinocerebellar A	Ataxia (CAPN1)	Clear
O Late-Onset Neuronal Ceroid	Lipofuscinosis, NCL 12 (ATP13A2, Australian Cattle Dog Variant)	Clear
O Leonberger Polyneuropathy	1 (LPN1, ARHGEF10)	Clear
O Leonberger Polyneuropathy	2 (GJA9)	Clear
🔗 Lethal Acrodermatitis, LAD (I	MKLN1)	Clear
O Leukodystrophy (TSEN54 Ex	on 5, Standard Schnauzer Variant)	Clear
🧭 Ligneous Membranitis, LM (F	PLG)	Clear
S Limb Girdle Muscular Dystro	phy (SGCD, Boston Terrier Variant)	Clear
C Limb-Girdle Muscular Dystro	ophy 2D (SGCA Exon 3, Miniature Dachshund Variant)	Clear



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OTHER RESULTS

Comp QT Syndrome (KCNQ1)	Clear
Lundehund Syndrome (LEPREL1)	Clear
Macular Corneal Dystrophy, MCD (CHST6)	Clear
Malignant Hyperthermia (RYR1)	Clear
May-Hegglin Anomaly (MYH9)	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
Mucopolysaccharidosis Type VII, Sly Syndrome, MPS VII (GUSB Exon 5, Terrier Brasileiro Variant)	Clear
Multiple Drug Sensitivity (ABCB1)	Clear
Muscular Dystrophy (DMD, Cavalier King Charles Spaniel Variant 1)	Clear
Musladin-Lueke Syndrome, MLS (ADAMTSL2)	Clear
Myasthenia Gravis-Like Syndrome (CHRNE, Heideterrier Variant)	Clear



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OTHER RESULTS		
🧭 Myotonia Congenita (CLCN1 E	Exon 23, Australian Cattle Dog Variant)	Clear
🔗 Myotonia Congenita (CLCN1 E	Exon 7, Miniature Schnauzer Variant)	Clear
Narcolepsy (HCRTR2 Exon 1, I	Dachshund Variant)	Clear
Narcolepsy (HCRTR2 Intron 4,	, Doberman Pinscher Variant)	Clear
Narcolepsy (HCRTR2 Intron 6	, Labrador Retriever Variant)	Clear
Nemaline Myopathy (NEB, Am	ierican Bulldog Variant)	Clear
Neonatal Cerebellar Cortical	Degeneration (SPTBN2, Beagle Variant)	Clear
Neonatal Interstitial Lung Dise	ease (LAMP3)	Clear
Neuroaxonal Dystrophy, NAD ((VPS11, Rottweiler Variant)	Clear
Neuroaxonal Dystrophy, NAD ((TECPR2, Spanish Water Dog Variant)	Clear
Neuronal Ceroid Lipofuscinos	sis 1, NCL 1 (PPT1 Exon 8, Dachshund Variant 1)	Clear
Neuronal Ceroid Lipofuscinos	sis 10, NCL 10 (CTSD Exon 5, American Bulldog Variant)	Clear
Neuronal Ceroid Lipofuscinos	sis 2, NCL 2 (TPP1 Exon 4, Dachshund Variant 2)	Clear
Neuronal Ceroid Lipofuscinos	sis 5, NCL 5 (CLN5 Exon 4 SNP, Border Collie Variant)	Clear
Neuronal Ceroid Lipofuscinos	sis 6, NCL 6 (CLN6 Exon 7, Australian Shepherd Variant)	Clear
Neuronal Ceroid Lipofuscinos	sis 7, NCL 7 (MFSD8, Chihuahua and Chinese Crested Variant)	Clear
Neuronal Ceroid Lipofuscinos	sis 8, NCL 8 (CLN8, Australian Shepherd Variant)	Clear
Neuronal Ceroid Lipofuscinos	sis 8, NCL 8 (CLN8 Exon 2, English Setter Variant)	Clear



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Polycystic Kidney Disease, PKD (PKD1)
 Pompe's Disease (GAA, Finnish and Swedish Lapphund, Lapponian Herder Variant)
 Prekallikrein Deficiency (KLKB1 Exon 8)
 Primary Ciliary Dyskinesia, PCD (NME5, Alaskan Malamute Variant)
 Primary Ciliary Dyskinesia, PCD (CCDC39 Exon 3, Old English Sheepdog Variant)
 Clear



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OTHER RESULTS

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
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



DNA Test Report Test Date: May 16th, 2023 embk.me/jingles29 **OTHER RESULTS** Pyruvate Dehydrogenase Deficiency (PDP1, Spaniel Variant) Clear \oslash Pyruvate Kinase Deficiency (PKLR Exon 5, Basenji Variant) Clear \bigcirc Pyruvate Kinase Deficiency (PKLR Exon 7, Beagle Variant) \oslash Clear \oslash Pyruvate Kinase Deficiency (PKLR Exon 10, Terrier Variant) Clear Pyruvate Kinase Deficiency (PKLR Exon 7, Labrador Retriever Variant) Clear (\checkmark) Pyruvate Kinase Deficiency (PKLR Exon 7, Pug Variant) Clear \bigcirc Raine Syndrome (FAM20C) Clear \oslash Recurrent Inflammatory Pulmonary Disease, RIPD (AKNA, Rough Collie Variant) Clear \oslash Renal Cystadenocarcinoma and Nodular Dermatofibrosis (FLCN Exon 7) Clear (~) $\langle \rangle$ Sensory Neuropathy (FAM134B, Border Collie Variant) Clear Severe Combined Immunodeficiency, SCID (PRKDC, Terrier Variant) Clear $\langle \rangle$ Severe Combined Immunodeficiency, SCID (RAG1, Wetterhoun Variant) Clear \oslash 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 (>) \oslash Skin Fragility Syndrome (PKP1, Chesapeake Bay Retriever Variant) Clear Spinocerebellar Ataxia (SCN8A, Alpine Dachsbracke Variant) Clear \oslash Spinocerebellar Ataxia with Myokymia and/or Seizures (KCNJ10) Clear \oslash



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OTHER RESULTS

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
O Thrombopathia (RASGRP1 Exon 5, American Eskimo Dog Variant)	Clear
O Thrombopathia (RASGRP1 Exon 5, Basset Hound Variant)	Clear
O Thrombopathia (RASGRP1 Exon 8, Landseer Variant)	Clear
Trapped Neutrophil Syndrome, TNS (VPS13B)	Clear
O Ullrich-like Congenital Muscular Dystrophy (COL6A3 Exon 10, Labrador Retriever Variant)	Clear
O Ullrich-like Congenital Muscular Dystrophy (COL6A1 Exon 3, Landseer Variant)	Clear
O Unilateral Deafness and Vestibular Syndrome (PTPRQ Exon 39, Doberman Pinscher)	Clear
Urate Kidney & Bladder Stones (SLC2A9)	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



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OTHER RESULTS

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





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RESULT

INBREEDING AND DIVERSITY

CATEGORY

Coefficient Of Inbreeding

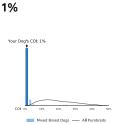
Our genetic COI measures the proportion of your dog's genome where the genes on the mother's side are identical by descent to those on the father's side.

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.

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 mixed breed dogs:



High Diversity

How common is this amount of diversity in mixed breed dogs:



