



Genetic Profile Test Results

HORSE ID: 032120_008

Horse: CPH Taiga Kimi Play

PACK: APHA

Owner: Cecile Penverne

Horse and Owner Information

Horse	CPH Taiga Kimi Play	DOB	2020-02-24
Breed	American Paint Horse	Age	0 years, 1 months
Color	Brown	Sex	mare
Discipline	Cutting, Ranch Horse, Reining	Height
Registry	American Paint Horse Association	Reg Number
Sire	Giant heart rof	Dam	Silbury kimimila
Sire Reg & No.	American Paint Horse Association 1060215	Dam Reg & No.	American Paint Horse Association 981133
Comments	Description: 4 white socks		

Owner	Cecile Penverne	Address	la villeneuve zinsec
Phone	662307475	City, State	Bern, EUROPE
Email	cymelapainthorse@free.fr	Postal Code	56240



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Results Summary

Coat Color: CPH Taiga Kimi Play has two Black variants and no Red variants, indicating the base coat color appears Black. One copy of the Dominant Agouti variant was detected; invisible on a Red base, it pushes/restricts Black out to points; legs, ear tips, etc. appearing Bay. One Dominant White 20 variant and one Tobiano variant was detected which may result in White markings. One Dun variant was detected which may dilute base coat color. One Cream variant was detected which may dilute base coat color, possibly appearing Buckskin. As a result of the variant count in each of the following, she has a minimum 100% chance of passing Black, and 50% Dominant Agouti and/or Cream and/or Dun and/or Dominant White 20 and/or Tobiano to any offspring.

Variant Summary: **Aa, EE, CR/n, D/nd2, W20/n, TO/n**

Myostatin: Sprint Type

6 panel negative: GBED n/n, HERDA n/n, HYPP n/n, MH n/n, PSSM1 n/n, LWO n/n

Traits: CPH Taiga Kimi Play has not tested positive for any known disease variants on this panel.

Please note: Your analysis is ongoing and may include some regions marked with an asterisk denoting the following.
* Discovery - This gene detection is in the early stages of discovery and will have varying reliability results.
** Inconclusive - Not a bad omen! Simply put, the gene of interest did not reveal itself (neither a positive nor a negative; no result, therefore unknown).



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Coat Color Results

Base

Agouti	+/-	ASIP	Aa - One dominant Agouti variant detected; restricts any Black base to appear Bay.	More about A
Black/Red	+/+	MC1R	EE - Two Black variants detected and no Red.	More about E

Modifier

Brindle/IP	-/-	IKBKG	No Brindle/IP variants detected.	More about IP
Grey	-/-	STX17A	No Grey variants detected.	More about G

Dilution

Champagne	-/-	SLC36A1	No Champagne variants detected.	More about CH
Cream	+/-	SLC45A2	CR/n - One Cream variant detected.	More about CR
Dun	+/-,-/-	TBX3	D/nd2 (Dun). One Dun variant and one non-dun2 variant detected. Can produce non-dun offspring without primitive markings.	More about Dun
Pearl	-/-	SLC45A2	No Pearl variants detected.	More about prl
Silver	-/-	PMEL17	No Silver variants detected.	More about Z



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Coat Color Results, continued

White Patterns Results

Dominant White	+/-	<i>KIT</i>	W20/n - One Dominant White 20 variant detected (DW1-21).	More about DW
Frame Overo (LWO)	-/-	<i>EDNRB</i>	No Frame Overo (LWO) variants detected.	More about LWO
Leopard Complex Spotting (LP)	-/-	<i>TRPM1</i>	No Leopard Complex Spotting (LP) variants detected.	More about LP
Pattern 1 (LP modification)	-/-	<i>RFWD3</i>	No Pattern 1 (LP modification) variants detected.	More about PATN1
Splashed White (MITF)	-/-,-/-	<i>MITF</i>	No Splashed White 1 or Splashed White 3 variants detected.	More about SW (MITF)
Splashed White (PAX3)	-/-,-/-	<i>PAX3</i>	No Splashed White 2 or Splashed White 4 variants detected.	More about SW (PAX3)
Sabino 1	-/-	<i>KIT</i>	No Sabino variants detected.	More about SB1
Tobiano	+/-	<i>ECA3</i>	TO/n - One Tobiano variant detected.	More about TO



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Health Genetics 1

Immune System

Foal Immunodeficiency Syndrome	-/-	SLC5A3	No Foal Immunodeficiency Syndrome variants detected.	More about fis
Severe Combined Immunodeficiency	-/-	DNAPK	No Severe Combined Immunodeficiency variants detected.	More about scid
West Nile Virus Susceptibility*	-/-	OAS1	Normal susceptibility to West Nile Virus symptoms.	More about WNVR*
Immune-mediated Myositis*	**	MYH1	**Upon request only, inquire about upgrade.	More about IMM*

Muscle Disorders

Glycogen Branching Enzyme Deficiency	-/-	GBE1	No Glycogen Branching Enzyme Deficiency variants detected.	More about gbed
Hyperkalemic Periodic Paralysis	-/-	SCN4A	No Hyperkalemic Periodic Paralysis variants detected.	More about HYPP
Malignant Hyperthermia	-/-	RYR1	No Malignant Hyperthermia variants detected.	More about MH
Myotonia	-/-	CLCN4	No Myotonia variants detected.	More about myt
Polysaccharide Storage Myopathy (type 1)	-/-	GYS1	No Polysaccharide Storage Myopathy type 1 variants detected.	More about PSSM1



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Health Genetics 2

Neurologic Disorders

Cerebellar Abiotrophy	-/-	<i>MUTYH</i>	No Cerebellar Abiotrophy variants detected.	More about ca
Lavender Foal Syndrome	-/-	<i>MYO5A</i>	No Lavender Foal Syndrome variants detected.	More about lfs

Reproductive Disorders

Androgen Insensitivity	-/-	<i>AR</i>	No Androgen Insensitivity variants detected.	More about as
IAR - Subfertility*	-/-,+/-	<i>FKBP6</i>	One IAR Subfertility* variant detected; likely no effect.	More about iar*

Skin Disorders

Hereditary Equine Regional Dermal Asthenia	-/-	<i>PPIB</i>	No Hereditary Equine Regional Dermal Asthenia variants detected.	More about herda
Junctional Epidermolysa Bullosis (type 1)	-/-	<i>LAMC2</i>	No Junctional Epidermolysa Bullosis (type 1) variants detected.	More about jeb1
Junctional Epidermolysa Bullosis (type 2*)	-/-	<i>LAMA3</i>	No Junctional Epidermolysa Bullosis (type 2*) variants detected.	More about jeb2*



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Other Genetics

Trait Genetics

Lordosis*	+/-,-/-,-/-,+/-	<i>ECA20</i>	No pattern of Lordosis* variants detected.	More about L*
Curiosity/Vigilance*	+/-	<i>DRD4</i>	One Curiosity and one Vigilance variant detected; likely both curious and vigilant.	More about Cur/Vig
Myostatin/Speed	+/+	<i>MSTN</i>	Two Sprint variants detected; likely Sprint ability over Endurance.	More about MSTN
DMRT3	-/-	<i>DMRT3</i>	No DMRT3 variants detected.	More about DMRT3

New Additions

Equine Recurrent Uveitis (Risk)*	***	<i>ECA18</i>	***DNA Minipanel PLUS only, inquire about upgrade.	More about ERU
Equine Recurrent Uveitis (Severity)*	***	<i>ECA20</i>	***DNA Minipanel PLUS only, inquire about upgrade.	More about ERU
Equine Metabolic Syndrome*	***	<i>FAM174A</i>	***DNA Minipanel PLUS only, inquire about upgrade.	More about EMS
Laminitis Risk*	***	<i>FAM174A</i>	***DNA Minipanel PLUS only, inquire about upgrade.	More about LAM
Squamous Cell Carcinoma*	***	<i>DDB2</i>	***DNA Minipanel PLUS only, inquire about upgrade.	More about SCC
Tiger Eye*	***	<i>SLC24A5</i>	***DNA Minipanel PLUS only, inquire about upgrade.	More about Tiger Eye
Dwarfism*	***	<i>ACAN</i>	***DNA Minipanel PLUS only, inquire about upgrade.	More about Dwarfism



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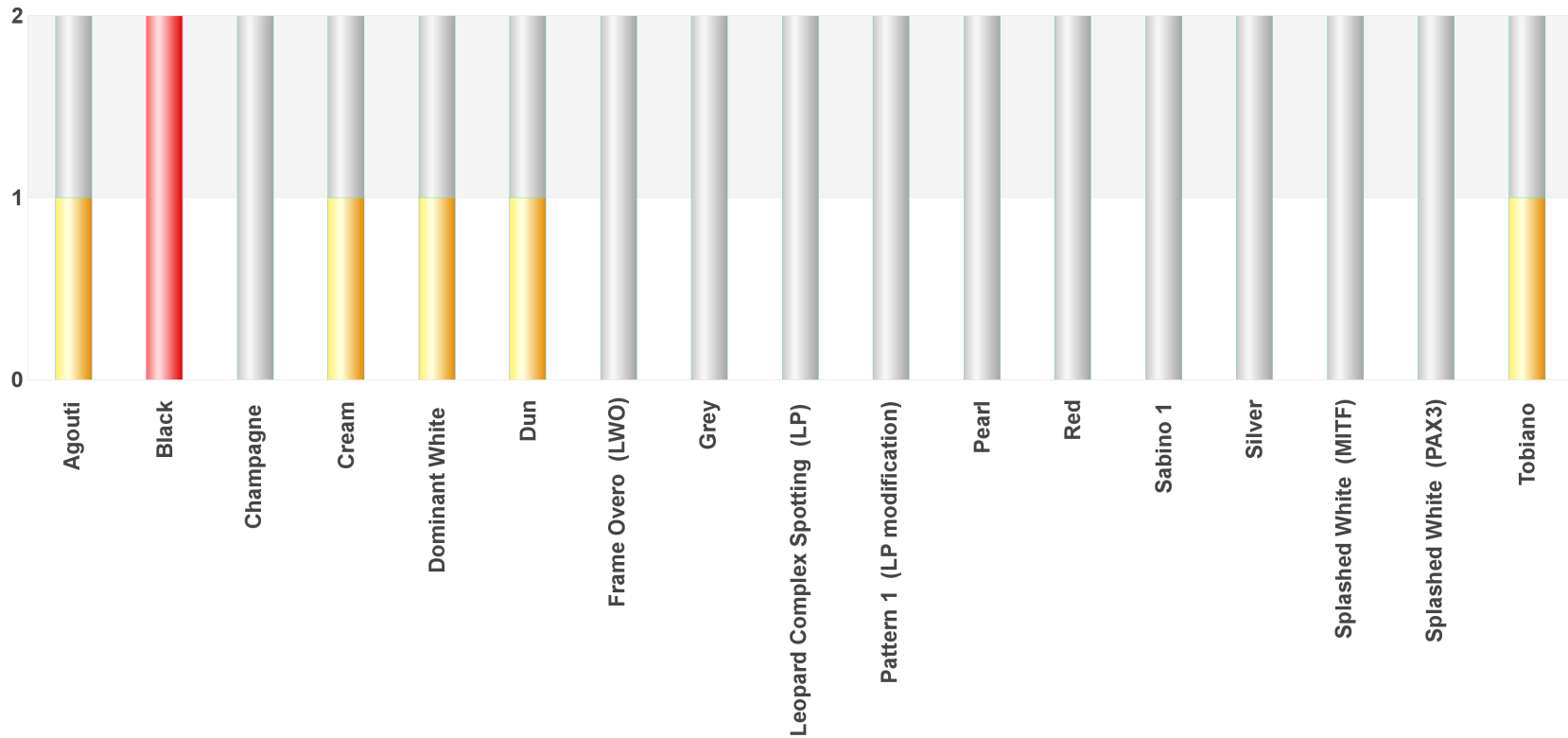
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Inheritance Probabilities

Coat Color



Coat Color Inheritance Probabilities: The bar graph above depicts the number of variants for specific coat color phenotypes based upon your horse's genetic testing results. Completely filled red bar represents two such variants (homozygous) and a half-filled yellow bar represents one such variant (heterozygous).



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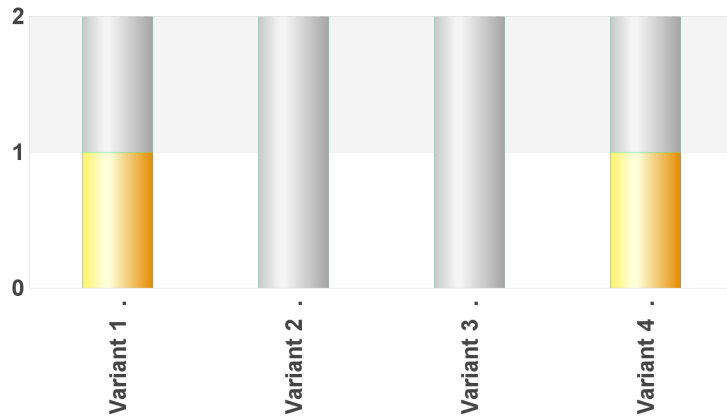
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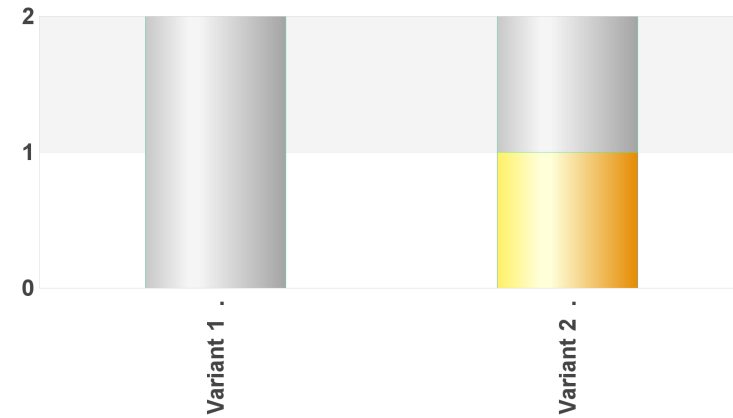
Inheritance Probabilities

Lordosis



Not affected

IAR Subfertility*



Not affected

Multi-variant Risk Charts: Each chart represents a trait, and each bar indicates a distinct risk or variant presence. These act in combination to produce the trait. A red bar indicates the horse carries 2 risk variants at the site; a partly-yellow bar indicates 1 risk variant; and a fully-grey bar indicates 0 risk variants. If all bars are red, then the horse carries two risk variants at each risk site and is likely affected. If all bars contain yellow or red, but are not all red, then the horse is likely a carrier. Otherwise, the horse is not a likely carrier of the tested trait.



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Defining Genetics & More Info

Variant:	One of two or more alternative forms of a gene that arise by mutation and are found at the same place on a chromosome.
Variants: Heterozygous vs. Homozygous?	Variant calls are written in a way that denotes their origin and whether they are DOMINANT (uppercase) or recessive (lowercase). For example, at MC1R (also known as extension), Black is dominant and thus written as "E" whereas Red is recessive and thus denoted as "e". Therefore, an EE horse is homozygous for Black (and thus appears black), an ee horse is homozygous for Red (appears Red), and an Ee horse is heterozygous (shows the dominant variant, thus is Black).
Gene:	A unit of heredity that is transferred from a parent to offspring and is thought to determine some characteristic of the offspring.
Genotype:	The genetic constitution or make up of an individual organism.
Heterozygous:	A pair of genes which are different (not the same). One is typically dominant and one recessive.
Homozygous:	A pair of genes that are identical (of one type).
Phenotype:	The observable or visible characteristics of an individual resulting from their genotype or the interaction of their various genes and environment.

The results depicted in this report do not constitute veterinary or medical advice. Any medical or veterinary advice should be sought from your veterinarian regarding these results or any health issues or questions you may have about your animal. Breed, sex, gene interaction, unknown genes and individual variances may impact the results, phenotypes, and behaviors in any animal in unknown and unpredictable ways. Please be advised that your animals' health is important to us and you should feel free to contact us should you have any further questions or feedback on our diagnostic platform, results reporting, or general questions. We value your input and thank you!