



# Genetic Profile Test Results

HORSE ID: 112120\_014

**Horse:** Haap Hazard

PACK: Etalon DNA Minipanel

**Owner:** Steve Hugus

## Horse and Owner Information

<b>Horse</b>	Haap Hazard	<b>DOB</b>	2020-08-13
<b>Breed</b>	Arabian	<b>Age</b>	0 years, 3 months
<b>Color</b>	Black	<b>Sex</b>	Stallion
<b>Discipline</b>	.....	<b>Height</b>	.....
<b>Registry</b>	Arabian Horse Association	<b>Reg Number</b>	682337
<b>Sire</b>	Unknown	<b>Dam</b>	Unknown
<b>Sire Reg &amp; No.</b>	Arabian Horse Association 653110	<b>Dam Reg &amp; No.</b>	Arabian Horse Association 655188
<b>Comments</b>	Known Issues: Black W-19 1 suspect		

<b>Owner</b>	Steve Hugus	<b>Address</b>	323 Sheep Camp
<b>Phone</b>	307-856-0212 3078560212	<b>City, State</b>	Pavillion, WY
<b>Email</b>	shugus@wyoming.com	<b>Postal Code</b>	82523



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### Results Summary

**Coat Color:** Haap Hazard has one Red variant and one Black variant, indicating the base coat color appears Black. One Dominant White 19 variant was detected which may result in White markings. As a result of the variant count in each of the following, he has a minimum 50% chance of passing Red or Black, and 50% Dominant White 19 to any offspring.

**Variant Summary:** aa, Ee, nd1/nd2, W19/n  
**Myostatin: Endurance Type**

**3 panel negative: CA n/n, LFS n/n, SCID n/n**

**Traits:** Haap Hazard has not tested positive for any known disease variants on this panel. \*The DNA was also tested on our discovery/validation platform for non-Dun Primitive Markings. Preliminary results indicate this horse is heterozygous for non-Dun Primitive Markings (nd1/nd2) and may pass it to 50% of any offspring.

**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	-/-	<b>ASIP</b>	aa - No dominant Agouti variants detected; restricts any Black base to appear Bay.	<a href="#">More about A</a>
Black/Red	+/-	<b>MC1R</b>	Ee - One Black variant detected and one Red.	<a href="#">More about E</a>

### Modifier

Brindle/IP	-/-	<b>IKBK</b>	No Brindle/IP variants detected.	<a href="#">More about IP</a>
Grey	-/-	<b>STX17A</b>	No Grey variants detected.	<a href="#">More about G</a>

### Dilution

Champagne	-/-	<b>SLC36A1</b>	No Champagne variants detected.	<a href="#">More about CH</a>
Cream	-/-	<b>SLC45A2</b>	No Cream variants detected.	<a href="#">More about CR</a>
Dun	-/-,+/-	<b>TBX3</b>	nd1/nd2 (non-dun with possible primitive markings). One non-dun1 variant and one non-dun2 variant detected. No Dun variants detected.	<a href="#">More about Dun</a>
Pearl	-/-	<b>SLC45A2</b>	No Pearl variants detected.	<a href="#">More about prl</a>
Silver	-/-	<b>PMEL17</b>	No Silver variants detected.	<a href="#">More about Z</a>



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### Coat Color Results, continued

#### White Patterns Results

Dominant White	+/-	<i>KIT</i>	W19/n - One Dominant White 19 variant detected (DW1-21).	<a href="#">More about DW</a>
Frame Overo (LWO)	-/-	<i>EDNRB</i>	No Frame Overo (LWO) variants detected.	<a href="#">More about LWO</a>
Leopard Complex Spotting (LP)	-/-	<i>TRPM1</i>	No Leopard Complex Spotting (LP) variants detected.	<a href="#">More about LP</a>
Pattern 1 (LP modification)	-/-	<i>RFWD3</i>	No Pattern 1 (LP modification) variants detected.	<a href="#">More about PATN1</a>
Splashed White (MITF)	-/-,-/-	<i>MITF</i>	No Splashed White 1 or Splashed White 3 variants detected.	<a href="#">More about SW (MITF)</a>
Splashed White (PAX3)	-/-,-/-	<i>PAX3</i>	No Splashed White 2 or Splashed White 4 variants detected.	<a href="#">More about SW (PAX3)</a>
Sabino 1	-/-	<i>KIT</i>	No Sabino variants detected.	<a href="#">More about SB1</a>
Tobiano	-/-	<i>ECA3</i>	No Tobiano variants detected.	<a href="#">More about TO</a>



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## Health Genetics 1

### Immune System

<b>Foal Immunodeficiency Syndrome</b>	-/-	<b>SLC5A3</b>	No Foal Immunodeficiency Syndrome variants detected.	<a href="#">More about fis</a>
<b>Severe Combined Immunodeficiency</b>	-/-	<b>DNAPK</b>	No Severe Combined Immunodeficiency variants detected.	<a href="#">More about scid</a>
<b>West Nile Virus Susceptibility*</b>	+/-	<b>OAS1</b>	WNVR*/n - Increased susceptibility to West Nile Virus symptoms if contracted.	<a href="#">More about WNVR*</a>
<b>Immune-mediated Myositis*</b>	**	<b>MYH1</b>	**Upon request only, inquire about upgrade.	<a href="#">More about IMM*</a>

### Muscle Disorders

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



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## Health Genetics 2

### Neurologic Disorders

Cerebellar Abiotrophy	-/-	<b>MUTYH</b>	No Cerebellar Abiotrophy variants detected.	<a href="#">More about ca</a>
Lavender Foal Syndrome	-/-	<b>MYO5A</b>	No Lavender Foal Syndrome variants detected.	<a href="#">More about lfs</a>

### Reproductive Disorders

Androgen Insensitivity	-/-	<b>AR</b>	No Androgen Insensitivity variants detected.	<a href="#">More about as</a>
IAR - Subfertility*	-/-,+/-	<b>FKBP6</b>	One IAR Subfertility* variant detected; likely no effect.	<a href="#">More about iar*</a>

### Skin Disorders

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



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## Other Genetics

### Trait Genetics

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

### New Additions

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



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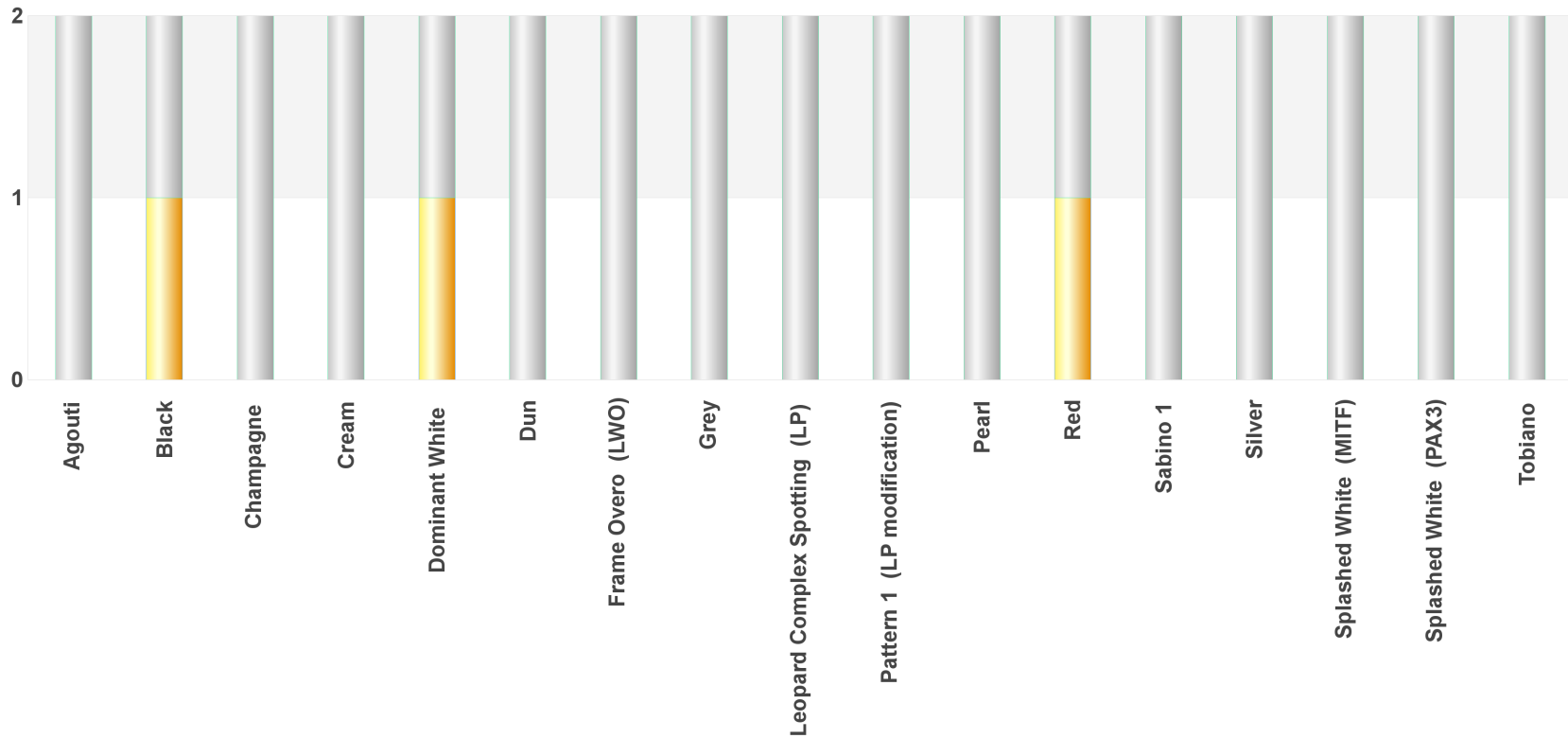
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## Inheritance Probabilities

### 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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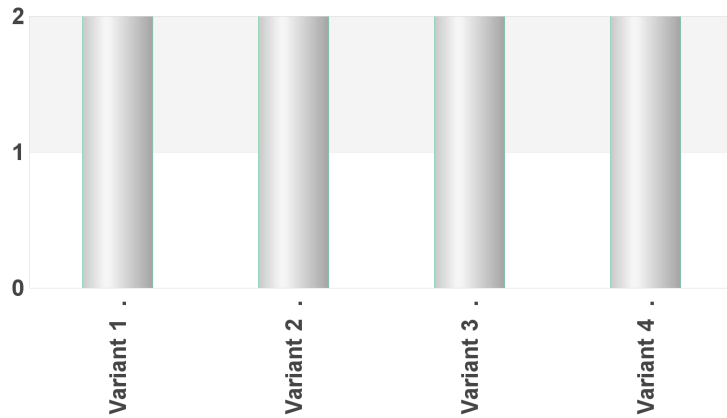
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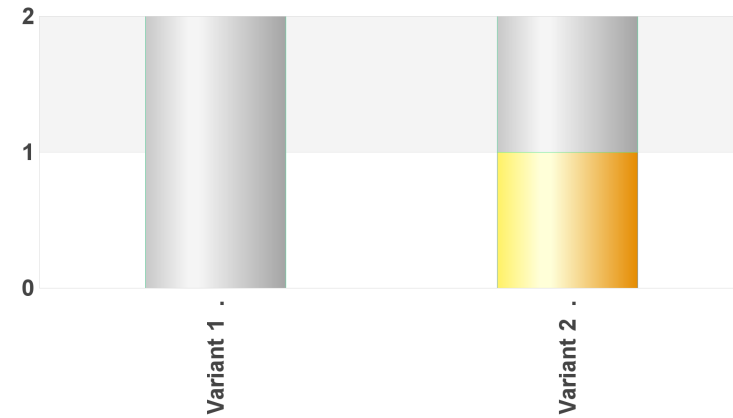
## Inheritance Probabilities

### 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

<b>Variant:</b>	One of two or more alternative forms of a gene that arise by mutation and are found at the same place on a chromosome.
<b>Variants: Heterozygous vs. Homozygous?</b>	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).
<b>Gene:</b>	A unit of heredity that is transferred from a parent to offspring and is thought to determine some characteristic of the offspring.
<b>Genotype:</b>	The genetic constitution or make up of an individual organism.
<b>Heterozygous:</b>	A pair of genes which are different (not the same). One is typically dominant and one recessive.
<b>Homozygous:</b>	A pair of genes that are identical (of one type).
<b>Phenotype:</b>	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!