

Genetic profile test results

HORSE ID:
042021_019PACKAGES:
APHA

Horse and owner information

Horse

Echo Eternal

Date of birth

06-06-2014

Breed

Paint (Tovero)

Age

7 y.o.

Color

Black

Sex

Stallion

Discipline

Breeding

Height

16 Hands

Registry

American Paint Horse Association

Reg number

1050037

Sire

BBF DR DIXON

Dam

PRISSY PUDDIN

Sire Reg & No.

**American Paint Horse Association
386443**

Dam Reg & No.

**American Paint Horse Association
521784**

Owner

Amy Ashby

Address

2118 PINE ROAD

Phone

7135031180

City, State

CLEVELAND, TX

Email

ashbyakgm@gmail.com

Postal code

77328

 Results Summary

Variant summaries:

Color: **a/a, E/E, TO/n, LWO/n, nd1/nd2**

Speed: **Sprint Type**

Temperament: **Curious**

Gait: **Neg for DMRT3**

Performance and Abilities:

Curious

Two Curiosity variants; horse may be more curious than vigilant.

Non-"Gaited" DMRT3

No DMRT3 variants; likely non-gaited (*variants for novel "gait" abilities are currently in research).

Sprint

Two Sprint type variants; horse may accel at short distance, quick bursts of speed over endurance type activity.

Coat color:

Black (base, E) - E/E

Black (E) is the base coat color for this horse and is a relatively uncommon coat color on its own (usually it is found in combination with other colors or modifiers such as in Bay horses). A visible difference between a true black (Ee or EE), a dark chestnut (ee) or a bay (Aa or AA + E) can sometimes be seen in the fine hairs around the eyes and muzzle. On a true black these hairs typically remain black even if the horse is sun-bleached, while on other colors they will be lighter. For a horse to be "homozygous black", it must have TWO copies of the Black variant (EE) and horse has 100% chance of passing Black to any offspring.

Tobiano (TO) - TO/n

TO/n - One Tobiano (TO) variant detected; may result in large white markings on body, face and legs (occasionally produces 'minimal' or 'slipped' Tobiano horse with very little white only on face and lower legs (ongoing heritability study for minimal Tobiano). Horse has 50% chance of passing to any offspring.

Overo (LWO) - LWO/n

LWO/n - One Frame/Lethal White Overo (LWO) variant detected; may result in White markings and possibly blue eye(s). Horse has 50% chance of passing to any offspring.

non-Dun Primitive Markings (nd) - nd1/nd2

nd1/nd2 - Non-Dun (nd) Primitive Markings may result in dorsal striping, leg barring and "shadows" on the face and shoulder. Still in research, it is suspected in some cases that nd1 may also cause mild coat color dilution, though not as extensive as the Dun (D) variant. Horse with one copy of nd1 variant will pass nd1 to 50% of any offspring.

Blue Eyes Possible

Horses with the Tobiano variant (TO/n or TO/TO) in addition to the Lethal White Overo variant (LWO/n) have an increased likelihood of having one or two blues eyes.

Tovero (TO/TO, LWO/n or TO/n, LWO/n)

Horses with at least one Tobiano variant (TO/n or TO/TO) and one Lethal White Overo variant (LWO/n) are often referred to as "Tovero"

Health Variants:

Lethal White Overo (LWO) - LWO/n

LWO/n - One Frame/Lethal White Overo (LWO) variant detected, resulting in "Carrier" status and may exhibit White markings. Caution is recommended when breeding to avoid another carrier and thus a 25% chance of foal fatality. Read more about LWO by clicking the name of the variant above.

 Coat color BaseAgouti (A) - a/a

ASIP



Negative

Gene or region: **ASIP**

a/a - No Dominant Agouti variants detected. Agouti (A) restricts black pigment to the outer regions of the body, the legs, mane & tail, nose, ear tips causing the otherwise black horse to appear Bay. Agouti is invisible on the red based coat.

Black (E) - E/E

MC1R



Black Based

Gene or region: **MC1R**

E/E - Two Black variants detected and no Red.

Modifiers

Grey (G) - n/n

STX17A



Negative

Gene or region: **STX17A**

No Grey (G) variants detected.

Brindle (BR1) - n/n

MBTPS2BR1



Negative

Gene or region: **MBTPS2BR1**

n/n - No Brindle (BR1) variants detected. Horse with Brindle (BR1) may display overall haircoat showing streaks of darker and lighter hair, similar to the brindle coat color in other species.

non-Dun Primitive Markings (nd) - nd1/nd2

TBX3



Possibly
Affected

Gene or region: **TBX3**

nd1/nd2 - One non-Dun Primitive Markings variant detected. Non-Dun Primitive Markings can appear as a dorsal stripe, leg barring, shadows on the face and shoulders even in the absence of the Dun variant.

Dilutes

Champagne (CH) - n/n

SLC36A1



Negative

Gene or region: **SLC36A1**

No Champagne (CH) variants detected.

Silver (Z) - n/n

PMEL17



Negative

Gene or region: **PMEL17**

No Silver (Z) variants detected. Silver affects only black base (E) coat colors and is invisible on a red-based coat. On black or bay, it has a disproportionate diluting effect on the mane and tail, and is sometimes called, "Chocolate Flaxen".

Cream (CR) - n/n

SLC45A2



Negative

Gene or region: **SLC45A2**

No Cream (CR) variants detected.

Pearl (PRL) - n/n

SLC45A2



Negative

Gene or region: **SLC45A2**

No Pearl (prl) variants detected.

Dun (D) - n/n

TBX3



Negative

Gene or region: **TBX3**

No Dun (D) variants detected. Dun is a modifier that dilutes the base coat color often revealing Primitive Markings such as a dorsal stripe, leg barring, shadows on the face and shoulders.

Whites

Frame/Lethal White Overo (LWO) - LWO/n

EDNRB



Likely Affected

Gene or region: **EDNRB**

LWO/n - One Frame/Lethal White Overo variant detected. May result result in White markings and possibly blue eye(s).

Tobiano (TO) - TO/n

ECA3



Likely Affected

Gene or region: **ECA3**

TO/n - One Tobiano variant detected; likely white markings. This horse may pass Tobiano to 50% of any offspring.

Leopard Spotting Complex (LP) - n/n

TRPM1



Negative

Gene or region: **TRPM1**

No Leopard Complex Spotting (SP) variants detected.

Pattern 1 (PATN1) - n/n

RFWD3



Negative

Gene or region: **RFWD3**

No Pattern (PATN1/n) 1 variants detected.

Sabino1 (SB1) - n/n

KIT



Negative

Gene or region: KIT

No Sabino (SB1) variants detected.

Splashed White (SW1) - n/n

MITF



Negative

Gene or region: MITF

No Splashed White 1 (SW1) variants detected.

Splashed White (SW2) - n/n

PAX3



Negative

Gene or region: PAX3

No Splashed White 2 (SW2/n) variants detected.

Splashed White (SW3) - n/n

MITF



Negative

Gene or region: MITF

No Splashed White 3 (SW3) variants detected.

Splashed White (SW4) - n/n

PAX3



Negative

Gene or region: PAX3

No Splashed White 4 (SW4) variants detected.

Dominant White (W) - n/n

KIT



Negative

Gene or region: KIT

No Dominant White (1-21) variants detected.

 Health Variants Color Related RiskLethal White Overo (LWO) - LWO/n

EDNRB



Carrier

Gene or region: **EDNRB**

LWO/n - One Frame/Lethal White Overo (LWO) variant detected, resulting in "Carrier" status and may exhibit White markings. Caution is recommended when breeding to avoid another carrier and thus a 25% chance of foal fatality. Read more about LWO by clicking the name of the variant above.

Congenital Stationary Night Blindness (CSNB) -
n/n

TRPM1



Negative

Gene or region: **TRPM1**

n/n - No Leopard Complex Spotting (LP) variants detected, which is related to the presence of Congenital Stationary Night Blindness (CSNB) if horse is LP/LP. Horses with one copy of the Leopard Complex Spotting (LP) variant are not currently known to suffer any ill effects as a result. Horses with Congenital Stationary Night Blindness (CSNB) which may experience the inability to see in low to no-light conditions. Read more about CSNB by clicking the name of the variant above.

Multiple Congenital Ocular Anomalies (MCOA)
- n/n

PMEL17



Negative

Gene or region: **PMEL17**

n/n - No Silver variants detected which is related to the presence of Multiple Congenital Ocular Anomalies (MCOA). Read more about MCOA by clicking the name of the variant above.

 Immune System

Foal Immunodeficiency Syndrome (FIS) - n/n SLC5A3



Negative

Gene or region: SLC5A3

No Foal Immunodeficiency Syndrome (FIS) variants detected. Read more about FIS by clicking the name of the variant above.

Severe Combined Immunodeficiency (SCID) - n/n

DNAPK



Negative

Gene or region: DNAPK

No Severe Combined Immunodeficiency (SCID) variants detected. Read more about SCID by clicking the name of the variant above.

West Nile Virus Symptom Susceptibility Risk (WNVR) - n/n

OAS1



Negative

Gene or region: OAS1

No West Nile Virus Symptom Susceptibility Risk (WNVR) variants detected. Normal susceptibility to West Nile Virus symptoms. (*NOT a test for West Nile Virus). Read more about WNVR by clicking the name of the variant above.

Immune-mediated Myositis (IMM)

MYH1

Not ordered



Muscle Disorders

Glycogen Branching Enzyme Deficiency (GBED) - n/n

GBE1



Negative

Gene or region: GBE1

No Glycogen Branching Enzyme Deficiency (GBED) variants detected. Read more about GBED by clicking the name of the variant above.

[Hyperkalemic Partial Paralysis \(HYPP\) - n/n](#)

SCN4A



Negative

Gene or region: **SCN4A**

No Hyperkalemic Partial Paralysis (HYPP) variants detected. Read more about HYPP by clicking the name of the variant above.

[Malignant Hyperthermia \(MH\) - n/n](#)

RYR1



Negative

Gene or region: **RYR1**

No Malignant Hyperthermia (MH) variants detected. Read more about MH by clicking the name of the variant above.

[Myotonia \(MYT\) - n/n](#)

CLCN4



Negative

Gene or region: **CLCN4**

No Myotonia (MYT) variants detected. Read more about MYT by clicking the name of the variant above.

[Polysaccharide Storage Myopathy type 1 \(PSSM1\) - n/n](#)

GYS1



Negative

Gene or region: **GYS1**

No Polysaccharide Storage Myopathy type 1 (PSSM1) variants detected. Read more about PSSM1 by clicking the name of the variant above.

Neurologic Disorders

[Cerebellar Abiotrophy \(CA\) - n/n](#)

MUTYH



Negative

Gene or region: **MUTYH**

No Cerebellar Abiotrophy (CA) variants detected. Read more about CA by clicking the name of the variant above.

[Lavender Foal Syndrome \(LFS\) - n/n](#)

MYO5A



Negative

Gene or region: MYO5A

No Lavender Foal Syndrome (LFS) variants detected. Read more about LFS by clicking the name of the variant above.

[Hydrocephalus \(HDC\)](#)

B3GALNT2

Not ordered

[Recurrent Laryngeal Neuropathy \(RLN\)](#)

ECA3

Not ordered



Reproductive Disorders

[Androgen Insensitivity Syndrome \(AIS\) - n/n](#)

AR



Negative

Gene or region: AR

No pattern of Androgen Insensitivity Syndrome (AIS) variants detected. Read more about AIS by clicking the name of the variant above.

[Impaired Acrosomal Reaction - Subfertility Risk \(IAR\) - iar/iar, iar/n](#)

FKBP6IAR1...



Not Affected

Gene or region: FKBP6IAR1, FKBP6IAR2

No pattern for Impaired Acrosomal Reaction (IAR) - Subfertility Risk variants detected. Read more about IAR by clicking the name of the variant above.



Skin, Hoof and Connective Tissue Disorders

Hereditary Equine Regional Dermal Asthenia

(HERDA) - n/n

PPIB



Negative

Gene or region: PPIB

No Hereditary Equine Regional Dermal Asthenia (HERDA) variants detected. Read more about HERDA by clicking the name of the variant above.

Junctional Epidermolysa Bullosis type 1 (JEB1)

- n/n

LAMC2



Negative

Gene or region: LAMC2

No Junctional Epidermolysa Bullosis type 1 (JEB1) variants detected. Read more about JEB1 by clicking the name of the variant above.

Junctional Epidermolysa Bullosis type 2 (JEB2)

- n/n

LAMA3



Negative

Gene or region: LAMA3

No Junctional Epidermolysa Bullosis type 2 (JEB2) variants detected. Read more about JEB2 by clicking the name of the variant above.

"Warmblood" Fragile Foal Syndrome (FFS)

PLOD1

Not ordered

Hoof Wall Separation Disease (HWSD)

SERPINB11

Not ordered

Naked Foal Syndrome (NFS)

st14nfs

Not ordered



Ocular Disorders

Equine Recurrent Uveitis Susceptibility Risk

(ERUR)

BIEC2536712WB

Not ordered

<u>Equine Recurrent Uveitis Symptom Severity (ERUS)</u>	BIEC2421990WB	Not ordered
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<u>Squamous Cell Carcinoma Susceptibility Risk (SCC)</u>	DDB2	Not ordered
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Skeletal Disorders

<u>Dwarfism (D)</u>	ACAND1...	Not ordered
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<u>Friesian Dwarfism (FD)</u>	B4GALT7	Not ordered
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<u>Lordosis</u>	ECA20...	 Not detected
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Gene or region: ECA20, ECA20, ECA20, ECA20

Horses with one copy in each of the four Lordosis regions are not currently known to suffer any ill effects as a result. Horses with two copies in each of the four Lordosis regions exhibit signs of swayback. Currently studies are only proven in the N. Am. Saddlebred breed.

Endocrine Disorders

<u>Equine Metabolic Syndrome Susceptibility Risk (EMS)</u>	BIEC2263524EMS	Not ordered
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<u>Laminitis Susceptibility Risk - Equine Metabolic Syndrome related (LAM)</u>	BIEC2263524_LAM	Not ordered
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Additional Traits

Height (LCORL)

LCORL

Not ordered

 Performance and Abilities Gait TypeNon-"Gaited" DMRT3

DMRT3



Detected

Gene or region: **DMRT3**

No DMRT3 variants; likely non-gaited (*variants for novel "gait" abilities are currently in research).

 PerformanceSprint

MSTN



Detected

Gene or region: **MSTN**

Two Sprint type variants; horse may accel at short distance, quick bursts of speed over endurance type activity.

Temperament

Curious

DRD4



Detected

Gene or region: **DRD4**

Two Curiosity variants; horse may be more curious than vigilant.