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SANTA BARBARA • SANTA CRUZ

VETERINARY GENETICS LABORATORY SCHOOL OF VETERINARY MEDICINE ONE SHIELDS AVENUE DAVIS, CALIFORNIA 95616-8744 TELEPHONE: (530) 752-2211 FAX: (530) 752-3556

HORSE COAT COLOR TEST RESULTS

TONI PERDEW 3005 LEXINGTON CT BEDFORD, IA 50833 *Case:* DT37346

Date Received: 13-Jun-2014

Print Date: 17-Jun-2014

Report ID: 4910-2100-1551-3088

Verify report at www.vgl.ucdavis.edu/myvgl/verify.html

Horse: 2014 HI JACD SILVER'S FILLY Reg:

DOB: **05/12/2014** *Breed:* **QH** *Sex:* **M** *Alt. ID:*

Sire: SPIN WHIZ ME Reg:
Dam: HI JACD SILVER Reg:

Dam: HI JACD SILVER			Reg:			
	RED FACTOR	E/e	Both black and red factors detected. Either E or e transmitted to offspring. Basic color is black, bay or brown in the absence of other modifying genes.	W10 DOMINANT WHITE		Not requested.
	AGOUTI		Not requested.	SPLASHED WHITE		Not requested.
	CREAM	N/Cr	Heterozygous, dilute, one copy of Cream gene. Typical colors are palomino, buckskin and smoky black in the absence of other modifying genes.	TOBIANO		Not requested.
	PEARL		Not requested.	LEOPARD		Not requested.
	SILVER		Not requested.	GRAY		Not requested.
	DUN		Not requested.	ROAN		Not requested.
	CHAMPAGNE	MPAGNE Not requested.				
	LETHAL WHITE OVERO		Not requested.			
	SABINO 1		Not requested.			

For more detailed information on Horse Coat Color results, please visit: www.vgl.ucdavis.edu/services/coatcolorhorse.php

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HORSE COAT COLOR TEST RESULTS

TONI PERDEW 3005 LEXINGTON CT BEDFORD, IA 50833 *Case:* DT37347

Date Received: 13-Jun-2014

Print Date: 17-Jun-2014

Report ID: 5051-3363-6615-2017

Verify report at www.vgl.ucdavis.edu/myvgl/verify.html

Horse: KALAMITY DUNIT'S 2014 COLT Reg:

DOB: **04/12/2014** *Breed:* **QH** *Sex:* **S** *Alt. ID:*

Sire: SPIN WHIZ ME Reg:
Dam: KALAMITY DUNIT Reg:

Dam: KALAMITY DUNIT		LAWITY DUNIT	Reg:			
	RED FACTOR	E/E	No red factor detected. Horse cannot have red foals regardless of the color of mate. Basic color is black, bay or brown in the absence of other modifying genes.	W10 DOMINANT WHITE		Not requested.
	AGOUTI		Not requested.	SPLASHED WHITE		Not requested.
	CREAM	N/N	No evidence for the Cream dilution altered sequence detected. Basic color is sorrel or chestnut, bay or black in the absence of other modifying genes.	TOBIANO		Not requested.
	PEARL		Not requested.	LEOPARD		Not requested.
	SILVER		Not requested.	GRAY		Not requested.
	DUN	D/D	Horse is homozygous for the Dun gene. All offspring should be dun dilute.	ROAN		Not requested.
	CHAMPAGNE		Not requested.			
	LETHAL WHITE OVERO		Not requested.			
	SABINO 1		Not requested.			

For more detailed information on Horse Coat Color results, please visit: www.vgl.ucdavis.edu/services/coatcolorhorse.php

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HORSE COAT COLOR TEST RESULTS

TONI PERDEW 3005 LEXINGTON CT BEDFORD, IA 50833 *Case:* DT37348

Date Received: 13-Jun-2014

Print Date: 17-Jun-2014

Report ID: 7021-8306-8791-8131

Verify report at www.vgl.ucdavis.edu/myvgl/verify.html

Horse: 2014 HOLINDOC'S COLT Reg:

DOB: **04/01/2014** Breed: **QH** Sex: **S** Alt. ID:

Sire: HESA CHAMPAGNE WHIZ Reg:
Dam: HOLINDOC Reg:

Dam: HC	DLINDOC	Keg:			
RED FACTOR	E/E	No red factor detected. Horse cannot have red foals regardless of the color of mate. Basic color is black, bay or brown in the absence of other modifying genes.	W10 DOMINANT WHITE		Not requested.
AGOUTI	A/a	Black pigment distributed in points pattern. Basic color is bay or brown in the absence of other modifying genes.	SPLASHED WHITE		Not requested.
CREAM		Not requested.	TOBIANO		Not requested.
PEARL		Not requested.	LEOPARD		Not requested.
SILVER		Not requested.	GRAY		Not requested.
DUN		Not requested.	ROAN		Not requested.
CHAMPAGNE		Not requested.			
LETHAL WHITE OVERO		Not requested.			
SABINO 1		Not requested.			

For more detailed information on Horse Coat Color results, please visit: www.vgl.ucdavis.edu/services/coatcolorhorse.php

Horse Coat Color Results with Explanations

Red Factor

e/e - Only the red factor detected. Basic color is sorrel or chestnut in the absence of other modifying genes.

E/e - Both black and red factors detected. Either E or e transmitted to offspring. Basic color is black, bay or brown in the absence of other modifying genes.

E/E - No red factor detected. Horse cannot have red foals regardless of the color of mate. Basic color is black, bay or brown in the absence of other modifying genes.

Agouti

A/A - Black pigment distributed in points pattern. Basic color is bay or brown in the absence of other modifying genes.

A/a - Black pigment distributed in points pattern. Basic color is bay or brown in the absence of other modifying genes.

a/a - Only recessive allele detected. Black pigment distributed uniformly. Basic color is black in the absence of other modifying genes.

Cream

N/N - No evidence for the Cream dilution altered sequence detected. Basic color is sorrel or chestnut, bay or black in the absence of other modifying genes.

N/Cr - Heterozygous, dilute, one copy of Cream gene. Typical colors are palomino, buckskin and smoky black in the absence of other modifying genes.

Cr/Cr - Double dilute (two copies of Cream gene). Typical colors are cremello, perlino and smoky cream in the absence of other modifying genes.

Pearl Pearl

N/N - No evidence of the altered sequence detected.

N/Prl - One copy of the altered sequence detected. If Cream dilution is also present, a pseudo-double Cream dilute phenotype will result.

Prl/Prl - Two copies of the altered sequence detected. On a chestnut base color, a uniform apricot color of body hair, mane and tail will result.

<u>Tobiano</u>

N/N - No evidence of altered sequence detected. Horse is not Tobiano.

 $\ensuremath{N/TO}$ - One copy of altered sequence. Approximately 50% of the offspring will inherit Tobiano.

TO/TO - Two copies of altered sequence. Horse is homozygous for Tobiano. All offspring will inherit Tobiano.

W10 Dominant White

 $N\!/N$ - No evidence of altered sequence detected.

N/W10 - One copy of the W10 mutation detected. Horse will display some degree of white spotting but the specific pattern cannot be predicted.

W10/W10 - Two copies of the W10 gene detected.*

 \ast Homozygous W10/W10 horses may not be viable. This result may only be found in aborted fetuses produced in matings between two W10 carriers.

Leopard / Appaloosa

N/N - No copies of Leopard Complex (Appaloosa) spotting.

LP/N - 1 copy of Leopard Complex mutation, 50% of offspring will inherit the Leopard gene. $\!\!^*$

LP/LP - 2 copies of Leopard Complex mutation, all offspring will inherit the Leopard gene.* Horse has congenital stationary night blindness (CSNB).

* Expression of Leopard Complex is variable and white patterning may not be present in all horses that inherit the gene.

<u>Dun</u>

 $\ensuremath{D/D}\xspace$ - Horse is homozygous for the Dun gene. All offspring should be dun dilute.

D/N - Horse has one copy of the Dun gene. Horse will transmit Dun gene to 50% of the offspring.

N/N - Markers inconsistent with horse being Dun.

Silver

 $N\!/N$ - No evidence of the altered sequence detected.

N/Z - One copy of the altered sequence detected. Black-based horses will be chocolate with flaxen or lightened mane and tail. Bay-based horses will have lightened black pigment on lower legs, mane and tail. No effect on chestnut color.

Z/Z - Two copies of altered sequence detected. Black-based horses will be chocolate with flaxen or lightened mane and tail. Bay-based horses will have lightened black pigment on lower legs, mane and tail. No effect on chestnut color.

Lethal White Overo

N/N - No evidence for the altered sequence detected.

N/O - One copy of the altered sequence detected. If bred to another N/O horse, there is a 25% chance of producing a lethal white overo foal. The N/O type has been detected in Paints (including breeding stock), Pintos, Thoroughbreds, Miniatures, Quarter Horses and Tennessee Walking Horses.

O/O - Only the altered sequence in the EDNRB gene detected. This result has only been obtained with samples from lethal white overo foals.

Sabino 1

N/N - No evidence of altered sequence detected.

N/SB1 - One copy of the Sabino 1 gene detected. Horse typically may have 2 or more white legs, blaze, spots or roaning in the midsection and jagged margins around white areas.

SB1/SB1 - Two copies of the Sabino 1 gene detected. Complete or nearly complete white phenotype expected.

Champagne

 $N\!/N$ - No evidence of altered sequence detected.

N/Ch - One copy of the altered sequence detected. Chestnut color (red) is diluted to gold, bay to tan with brown points and black to darker tan with brown points.

Ch/Ch - Two copies of the altered sequence detected. All offspring are expected to be Champagne diluted.

<u>Gray</u>

 $N\!/N$ - No copies of the gray gene. Horse will not turn gray.

N/G - One copy of the gray gene. Horse will turn gray and approximately 50% of offspring will be gray.

G/G - Two copies of the gray gene. Horse will turn gray and all offspring will be gray.

Splashed White

SW-1

 $N\!/N$ - No copies of SW-1 mutation.

N/SW1 - Horse has one copy of the SW-1 mutation.

SW1/SW1 - Horse has two copies of the SW-1 mutation.

SW-2

N/N - No copies of SW-2 mutation.

 $N\!/SW2$ - Horse has one copy of the SW-2 mutation.

SW-3

N/N - No copies of SW-3 mutation.

N/SW3 - Horse has one copy of the SW-3 mutation.

Roan

Rn/Rn - 2 copies of Classic Roan variant are present. All offspring will inherit Classic Roan.

Rn/N - 1 copy of Classic Roan variant is present. 50% of offspring will inherit Classic Roan.

Rn*/Rn* - 2 copies of an alternative Roan variant are present. All offspring will inherit the alternative Roan variant.

 Rn/Rn^* - 1 copy of Classic Roan and 1 copy of an alternative Roan variant are present. All offspring will inherit either Classic Roan or the alternative Roan variant.

 Rn^*/N - 1 copy of an alternative Roan variant is present.

N/N - No copies of Roan are present.

The Roan Zygosity test is not a direct test for the Roan gene. The analysis is based on markers associated with Classic Roan in Belgian Draft Horses, Gypsy Cobs, Miniature Horses, Quarter Horses, Tennessee Walking Horses and Welsh Ponies.

Note: There are horses with a roaned appearance that test negative for the Classic Roan gene. The genetic basis for this pattern called roaning is not known at this time.