

## DOG COAT COLOR / NATURAL BOBTAIL TEST REPORT

<p><i>Provided Information:</i></p> <p><i>Name:</i>           <b>GREEN BOY- PY</b></p> <p><i>Registration:</i></p>	<p><i>Case:</i>               <b>NCD151274</b></p> <p><i>Date Received:</i>    20-Apr-2021</p> <p><i>Report Issue Date:</i> 06-May-2021</p> <p><i>Report ID:</i>         2206-9483-5817-5182</p> <p style="text-align: center; font-size: small;">Verify report at <a href="http://www.vgl.ucdavis.edu/verify">www.vgl.ucdavis.edu/verify</a></p>
<p><i>DOB:</i> <b>04/05/2021</b>   <i>Sex:</i> <b>Male</b>   <i>Breed:</i> <b>French Bulldog</b></p>	
<p><i>Sire:</i>   YOLO</p> <p><i>Reg:</i></p> <p><i>Microchip:</i></p>	<p><i>Dam:</i>   PHYLLIS</p> <p><i>Reg:</i></p> <p><i>Microchip:</i></p>

**RESULT**

**INTERPRETATION**

MC1R (E LOCUS)	e <sup>1</sup> /e <sup>1</sup>	2 copies of red/yellow/cream
<b>BROWN (B LOCUS)</b>	<b>B/b</b>	1 copy of brown present - carrier
<b>DILUTE (D LOCUS)</b>	<b>d<sup>1</sup>/d<sup>1</sup></b>	Dilute. 2 copies of the dilution variants.
<b>DOMINANT BLACK (K LOCUS)</b>	<b>N/N</b>	Dog does not have the dominant black mutation
<b>AGOUTI (A LOCUS)</b>	<b>a<sup>t</sup>/a<sup>t</sup></b>	Homozygous for black-and-tan
<b>MERLE</b>	<b>N/N</b>	No copies of the merle associated SINE insertion.
<b>PIEBALD (S LOCUS)</b>	<b>N/N</b>	Dog has no copies of piebald.
<b>HARLEQUIN (GREAT DANE)</b>		Not requested.
<b>NATURAL BOBTAIL</b>		Not requested.
<b>DOBERMAN OCA</b>		Not requested.
<b>GERMAN SHEPHERD PANDA SPOTTING</b>		Not requested.
<b>INTENSITY DILUTION</b>	<b>In/In</b>	2 copies of intensity dilution. Red pigment is likely to be diluted to cream or white.

## DOG COAT COLOR / NATURAL BOBTAIL TEST REPORT

<i>Client/Owner/Agent Information:</i> MANDY PILGRIM	<i>Case:</i> <b>NCD151274</b> <i>Date Received:</i> 20-Apr-2021 <i>Report Issue Date:</i> 06-May-2021 <i>Report ID:</i> 2206-9483-5817-5182  Verify report at <a href="http://www.vgl.ucdavis.edu/verify">www.vgl.ucdavis.edu/verify</a>
<i>Name:</i> <b>GREEN BOY- PY</b>	

### Additional Information

If testing for a disease or a disorder was performed and results indicate the animal is affected or at risk, we recommend contacting your veterinarian for further clinical evaluation and for additional information on disease and management.

For more detailed information on Coat Color test results, please visit our website at:  
[www.vgl.ucdavis.edu/services/coatcolordog.php](http://www.vgl.ucdavis.edu/services/coatcolordog.php)

For terms and conditions of testing, please see [www.vgl.ucdavis.edu/about/terms-and-conditions](http://www.vgl.ucdavis.edu/about/terms-and-conditions)

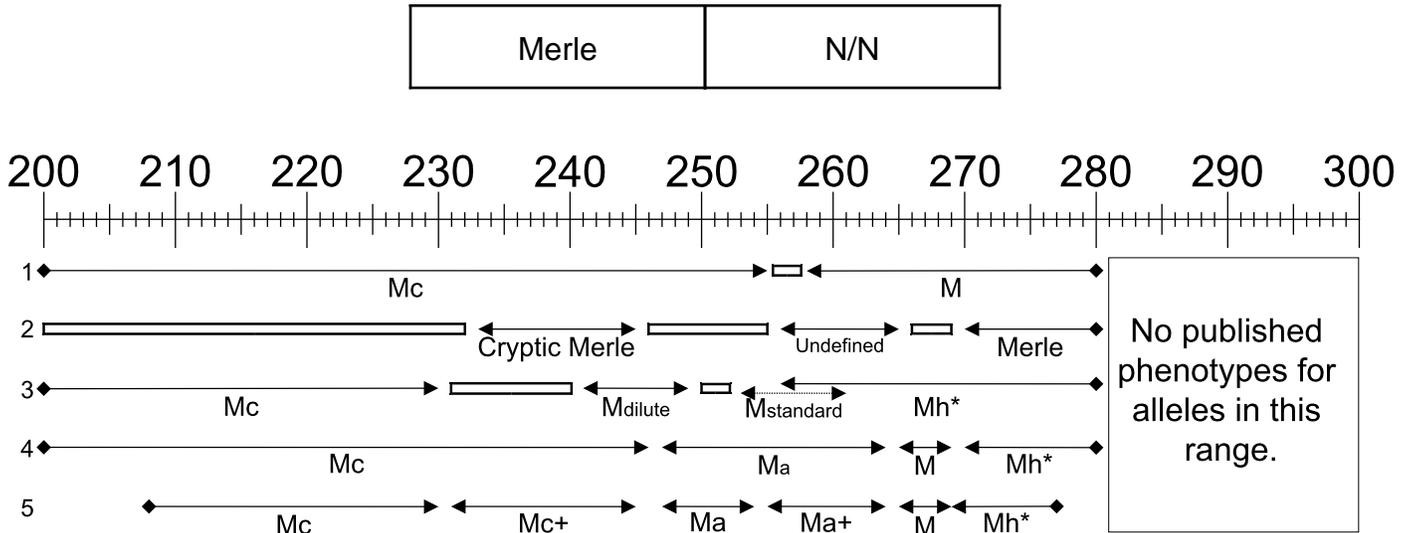
Results are determined using PCR-based methods. The results relate only to the sample tested as identified by the submitter (for example, identity and/or breed).

**Report authorized by Dr. Rebecca Bellone, VGL Director**

**ADDITIONAL INFORMATION FOR  
 MERLE RESULTS**

<b>Provided Information:</b>		<b>Case:</b> <b>NCD151274</b>
<b>Name:</b> <b>GREEN BOY- PY</b>		<b>Date Received:</b> 20-Apr-2021
<b>Registration:</b>		<b>Report Issue Date:</b> 06-May-2021
		<b>Report ID:</b> 2206-9483-5817-5182
Verify report at <a href="http://www.vgl.ucdavis.edu/verify">www.vgl.ucdavis.edu/verify</a>		
<b>DOB: 04/05/2021 Sex: Male Breed: French Bulldog</b>		
<b>Sire:</b> YOLO		<b>Dam:</b> PHYLLIS
<b>Reg:</b>		<b>Reg:</b>
<b>Microchip:</b>		<b>Microchip:</b>

Several interpretations and nomenclatures for the Merle variant have been proposed. Below is a graphical display of the merle alleles detected and the publications that define these nomenclatures.



Open boxes represent unassigned size variants within a specific naming system.

<sup>1</sup>Previous merle pattern result reported by the VGL.

Mc=200-255, M=258-280

<sup>2</sup>Merle pattern nomenclature defined by Clark et al. 2006.

<sup>3</sup>Merle pattern nomenclature defined by Murphy et al. 2018.

Mc=200-230, Mdilute=241-249, Mstandard=253-261, Mh=256-280

<sup>4</sup>Merle pattern nomenclature defined by Ballif et al. 2018.

Mc=200-246, Ma=247-264, M=265-269, Mh=270-280

<sup>5</sup>Merle pattern nomenclature defined by Langevin et al. 2018.

Mc=208-230, Mc+=231-245, Ma=247-254, Ma+=255-264, M=265-269, Mh=269-277

\* Mh “harlequin” is not the true Great Dane Harlequin (H) identified by Clark et al. 2008.

## DOG COAT TYPE TEST REPORT

<p><i>Provided Information:</i></p> <p><i>Name:</i>           <b>GREEN BOY- PY</b></p> <p><i>Registration:</i></p>	<p><i>Case:</i>               <b>NCD151274</b></p> <p><i>Date Received:</i>    20-Apr-2021</p> <p><i>Report Issue Date:</i> 03-May-2021</p> <p><i>Report ID:</i>         7692-5441-4405-7111</p> <p style="text-align: center; font-size: small;">Verify report at <a href="http://www.vgl.ucdavis.edu/verify">www.vgl.ucdavis.edu/verify</a></p>
<p><i>DOB:</i> <b>04/05/2021</b>   <i>Sex:</i> <b>Male</b>   <i>Breed:</i> <b>French Bulldog</b></p>	
<p><i>Sire:</i>   YOLO</p> <p><i>Reg:</i></p> <p><i>Microchip:</i></p>	<p><i>Dam:</i>   PHYLLIS</p> <p><i>Reg:</i></p> <p><i>Microchip:</i></p>

**RESULT**

**INTERPRETATION**

COAT LENGTH	S/L4	
<b>CURL</b>		Not requested.
<b>FURNISHINGS</b>		Not requested.
<b>IMPROPER COAT</b>		Not requested.

## DOG COAT TYPE TEST REPORT

<i>Client/Owner/Agent Information:</i> MANDY PILGRIM	<i>Case:</i> <b>NCD151274</b> <i>Date Received:</i> 20-Apr-2021 <i>Report Issue Date:</i> 03-May-2021 <i>Report ID:</i> 7692-5441-4405-7111  Verify report at <a href="http://www.vgl.ucdavis.edu/verify">www.vgl.ucdavis.edu/verify</a>
<i>Name:</i> <b>GREEN BOY- PY</b>	

### Additional Information

If testing for a disease or a disorder was performed and results indicate the animal is affected or at risk, we recommend contacting your veterinarian for further clinical evaluation and for additional information on disease and management.

For more detailed information on Dog Coat Type test results, please visit our website at:  
[www.vgl.ucdavis.edu/services/DogCoatLengthCurlandFurnishings.php](http://www.vgl.ucdavis.edu/services/DogCoatLengthCurlandFurnishings.php)

For terms and conditions of testing, please see [www.vgl.ucdavis.edu/about/terms-and-conditions](http://www.vgl.ucdavis.edu/about/terms-and-conditions)

Results are determined using PCR-based methods. The results relate only to the sample tested as identified by the submitter (for example, identity and/or breed).

**Report authorized by Dr. Rebecca Bellone, VGL Director**

## COCOA TEST REPORT

<i>Provided Information:</i>	<i>Case:</i> <b>NCD151274</b>
<i>Name:</i> <b>GREEN BOY- PY</b>	<i>Date Received:</i> 20-Apr-2021
<i>Registration:</i>	<i>Report Issue Date:</i> 28-Apr-2021
	<i>Report ID:</i> 8735-8238-0810-7124
Verify report at <a href="http://www.vgl.ucdavis.edu/verify">www.vgl.ucdavis.edu/verify</a>	
<i>DOB:</i> <b>04/05/2021</b> <i>Sex:</i> <b>Male</b> <i>Breed:</i> <b>French Bulldog</b>	
<i>Sire:</i> YOLO	<i>Dam:</i> PHYLLIS
<i>Reg:</i>	<i>Reg:</i>
<i>Microchip:</i>	<i>Microchip:</i>

### RESULT

### INTERPRETATION

<b>COCOA</b>	<b>co/co</b>
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2 copies of the cocoa variant.

## COCOA TEST REPORT

<i>Client/Owner/Agent Information:</i> MANDY PILGRIM	<i>Case:</i> <b>NCD151274</b> <i>Date Received:</i> 20-Apr-2021 <i>Report Issue Date:</i> 28-Apr-2021 <i>Report ID:</i> 8735-8238-0810-7124  Verify report at <a href="http://www.vgl.ucdavis.edu/verify">www.vgl.ucdavis.edu/verify</a>
<i>Name:</i> <b>GREEN BOY- PY</b>	

### Additional Information

If testing for a disease or a disorder was performed and results indicate the animal is affected or at risk, we recommend contacting your veterinarian for further clinical evaluation and for additional information on disease and management.

For more detailed information on Cocoa test results, please visit our website at:  
[www.vgl.ucdavis.edu/test/cocoa-dog](http://www.vgl.ucdavis.edu/test/cocoa-dog)

This test is specific for the autosomal recessive variant causing cocoa in French Bulldogs and is distinct from the other known variants resulting in a brown phenotype

For terms and conditions of testing, please see [www.vgl.ucdavis.edu/about/terms-and-conditions](http://www.vgl.ucdavis.edu/about/terms-and-conditions)

Results are determined using PCR-based methods. The results relate only to the sample tested as identified by the submitter (for example, identity and/or breed).

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