

# Genotyping protocol

## General information:

Strain name	CD1.129-Sigmar1<tm1Lmon>/Cnbc
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## **Primers:**

Name	Sequence	Primer type
SKO1	CTTACGGCGGTGATTTTGGCGATAC	
SKO2	CAACTGTTTACCTTGTGGAGCGACATC	

In case more than two primers are introduced, please indicate how they should be combined:

	Forward primer	Reverse primer
KO	SKO1	SKO2

## Reaction mix:

ddH <sub>2</sub> O	18,5	μl
PCR Buffer 15mM MgCl <sub>2</sub>	2,5	μl
dNTPs 2,5mM	2,5	μl
Primer 100μM	0,125	μl
Primer 100μM	0,125	μl
Taq Polymerase 5U/μl	0,250	μl
DNA	1	μl
Final volume	25	μl

## PCR program KO:

94	°C	2	min	X35
94	°C	30	sec	
60	°C	60	sec	
72	°C	60	sec	
72	°C	10	min	

## Expected fragment size:

KO	~300	bp
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## çComments/Additonal information:

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