

Wellcome Trust Sanger Institute Wellcome Trust Genome Campus Hinxton Cambridge CB10 1SA, U.K.

MGPgenotyping@sanger.ac.uk www.sanger.ac.uk

Gle1 (MCXN; EPD0359\_2\_A04)

Allele: Gle1<sup>tm1a(EUCOMM)Wtsi</sup>

# **Genotyping Information**

These mice may be genotyped through a combination of separate PCR reactions that detect *LacZ*, the gene-specific wild type allele, and a mutant allele-specific short range PCR. Interpretation of the consolidated results produces the genotype of the mice.

For example: LacZ positive, mutant positive, wild type positive = heterozygous.

### PCRs primer pairs and expected size bands

PCR type	Forward primer	Reverse primer	Expected size band (bp)
Mutant PCR	Gle1I_51135_F	CAS_R1_Term x	117
Wild type PCR	Gle1I_51135_F	Gle1I_51135_R	459
LacZ PCR	LacZ_2_small_F	LacZ_2_small_R	108

#### Primer sequences

Primer name	Primer sequence (5' > 3')		
CAS_R1_Term	TCGTGGTATCGTTATGCGCC		
Gle1I_51135_F	CTGTGGGGGTGGTGACTATG		
Gle1I_51135_R	TGGTGCCTACTAATTTCCTGTTCTC		
LacZ_2_small_F	ATCACGACGCGCTGTATC		
LacZ_2_small_R	ACATCGGGCAAATAATATCG		

This technical data sheet and information ("Datasheet") is supplied by Genome Research Limited ("GRL").

Although reasonable care is taken in the preparation of this Datasheet, GRL gives no warranties express or implied for any use of the Datasheet or for the accuracy of the Datasheet. GRL assumes no responsibility or liability for any decisions based upon the Datasheet. Without limiting the foregoing the Datasheet was prepared for mice supplied directly from GRL and where copies of this Datasheet are available from third party repositories or distribution centres ("Third Parties") GRL shall not be liable for any inconsistency between the mouse strain supplied by the Third Party and the Datasheet howsoever arising.

Last updated: 08/08/2010



Wellcome Trust Sanger Institute Wellcome Trust Genome Campus Hinxton Cambridge CB10 1SA, U.K.

MGPgenotyping@sanger.ac.uk www.sanger.ac.uk

#### Reaction

Reagent	μΙ
DNA (~50-100 ng)	1.0
10x Buffer MgCl <sub>2</sub> (50 mM)	2.0 0.6
PtTaq (Platinum® Taq (Invitrogen)) dNTPs (100 mM)	0.2 0.2
Primer 1 (10 μM) Primer 2 (10 μM)	0.4 0.4
H <sub>2</sub> 0 <b>Total</b>	15.2 <b>20.0</b>

## Cycling conditions

Wild type and mutant PCRs

Whatype and mataner ons		Lucz I OR			
Cycle			Cycle		
1	94 °C	5 min	1	94 °C	5 min
2	94 °C	30 sec	2	94 °C	30 sec
3	58 °C	30 sec	3	60 °C	30 sec
4	72 °C	45 sec	4	72 °C	30 sec
5		' + 34 cycles	5		+ 34 cycles
6	72 °C	5 min	6	72 °C	5 min
7	12 °C	forever	7	12 °C	forever

Lac7 PCR

 $This \ technical \ data \ sheet \ and \ information \ ("Datasheet") \ is \ supplied \ by \ Genome \ Research \ Limited \ ("GRL").$ 

Although reasonable care is taken in the preparation of this Datasheet, GRL gives no warranties express or implied for any use of the Datasheet or for the accuracy of the Datasheet. GRL assumes no responsibility or liability for any decisions based upon the Datasheet. Without limiting the foregoing the Datasheet was prepared for mice supplied directly from GRL and where copies of this Datasheet are available from third party repositories or distribution centres ("Third Parties") GRL shall not be liable for any inconsistency between the mouse strain supplied by the Third Party and the Datasheet howsoever arising.

Last updated: 08/08/2010