

**GENOTYPING BY PCR PROTOCOL  
MUTANT MOUSE REGIONAL RESOURCE CENTER: UC DAVIS**

[mmrrc@ucdavis.edu](mailto:mmrrc@ucdavis.edu)  
530-754-MMRRC

Please provide the following information required for genetic analysis of your mutant mice.

*Note to MAC users:* to ensure your graphic can be viewed on a PC please follow the steps below when inserting the graphic into this document. **DO NOT** drag and drop or copy/paste the graphic into this document.

- Open the original graphic in the program that created it
- Choose File, Save As
- Select No Compression in the save options.
- Save as JPG or PNG or similar format that's compatible with both PC and Mac Word versions.
- Switch to Word, choose Insert, Picture, From File and choose the newly saved picture.

*These instructions are very generic. The menu options for your graphics program may be different.*

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Institution <b>NIH/NIDDK</b>		
Address		
City	State	Zip
Lab Contact		
Email		
Telephone	FAX	
Strain Name <b>Ugcg</b>		MMRRC Stock Number <b>043532-UCD</b>

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NAME OF PCR: GlcT Lox MMRRC: 043532-UCD

Protocol: *(PCR protocol provided by Donating Investigator)*

Reagent/Constituent	Volume (µL)
Water	28.75
10x Buffer	5
MgCl <sub>2</sub> (stock concentration is 25mM)	5
Betaine (stock concentration is 5M) <i>Optional</i>	
dNTPs (stock concentration is 10mM)	5
DMSO <i>Optional</i>	
Primer 1. (stock concentration is 20µM)	0.5
Primer 2. (stock concentration is 20µM)	0.5
Primer 3. (stock concentration is 20µM)	
Primer 4. (stock concentration is 20µM)	
Taq Polymerase 5Units/µL	0.25
DNA (50-200ng/ µL) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	5
<b>TOTAL VOLUME OF REACTION:</b>	
<b>50.000 µL</b>	

*The total volume is auto-calculated based on volumes entered, right click the total and update field to show/recalculate the total volume.*

Comments on protocol:

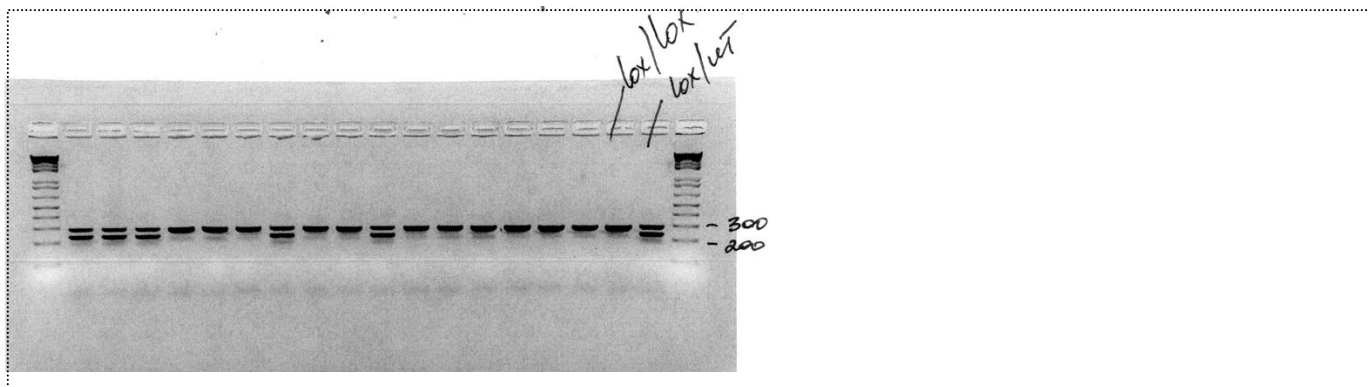
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Strategy:

Steps	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting <span style="float: right;">HOT START? <input type="checkbox"/></span>	94	5:00	1
2. Denaturation	94	1:00	
3. Annealing <span style="float: right;">steps 2-3-4 cycle in sequence</span>	68	1:00	45x
4. Elongation	72	1:00	
5. Amplification	72	3:00	1
6. Finish	4	∞	n/a

Primers:

Name	Nucleotide Sequence (5' - 3')	Argarose: 2.0% V: 90		
1. L-Ex6F	ATGTGCTAGATCAGGCAGGAGGGCTCATAG	Estimated Running Time: 45 min.		
2. L-int6-4	CCAACAGATATTGAATGCGAATGCTCTGCC	Primer Combination	Band (bp)	Genotype
3.		1 and 2	250	wt
4.		1 and 2	300	lox
5.				
6.				
7.				



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**Protocol / Gel Comments:**

**Gel pictures:**

*Please size gel images and comments  
to fit within this space*