

Protocol for genotyping LE-Tg(CX3CR1-Cre-ERT2)3Ottc transgenic rats January 08, 2019

Genomic DNA Preparation by Macherey-Nagel Tissue Spin Columns

Using this kit according to the manufacturer’s protocol is the preferred way for preparing genomic DNA at OTTC when it is intended for ddPCR (i.e. copy-number quantification). Typically, 10 to 90 ng of genomic DNA are used in an 11-25uL PCR reaction.

Genomic DNA Preparation by the “Hot Shot” method

This method was adapted by NIDA-OTTC from the protocol originally described in *Biotechniques* 29(1), 52-54 (2000). It has been used successfully with ear punches and tail clips of rats. This method is quicker and more cost-effective when compared to the MN Tissue Spin columns, but in our experience, it does not produce templates that are reliable for ddPCR and quantitative analysis. Typically, 1uL of template is sufficient for an 11-25uL PCR reaction.

- Place the biopsy sample in 1.5ml microfuge tube.
- Add 300 microliters of 50mM NaOH
- Incubate tubes at 95C for 60 minutes.
- Vortex tubes on medium power setting for 5 seconds.
- Quick spin the tubes to bring down the condensation.
- Neutralize each sample by adding 30 microliters of 1M Tris-HCl (pH 8).
- Vortex tubes on medium power setting for 5 seconds.
- Quick spin the tubes to bring down the condensation.

Debris (the “undigested” remnant of sample) may remain visible at the bottom of the tube. This is OK, but be sure to take only from the supernatant when setting the PCR reaction.

Use 1uL of this supernatant in a PCR reaction (11uL – 25uL final volume).

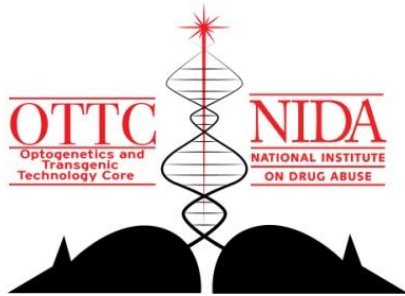
General PCR reaction setup:

- 11.0 uL 2x OneTaq master mix (New England Biolabs)
- 11.0 uL 2x specific oligos (0.5 uM Forward + 0.5uM Reverse; in water)
- 3.0 uL genomic DNA
- 25.0 uL PCR reaction

PCR Program CR2193		
Line	Temp	Time
Step 1	94oC	HOLD (hot start)
Step 2	94oC	2 min
Step 3	94oC	30 sec
Step 4	68oC	120 sec
Step 5	Go to Step 2	Repeat 34x
Step 6	68oC	5 min
Step 7	12oC	HOLD

This protocol was updated on 01-08-2019 by CR.

Any questions regarding protocol, contact nidatransgenicprojects@mail.nih.gov.



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<u>Primer Name</u>	<u>Primer Sequence (5' to 3')</u>
CX3CR1 F103289	TCAGGGTGGCCCATAACCAC
CreERT2 R713	AGGTAGTTATTCGGATCATCAGCTACAC

These oligos produce a 1017 bp amplicon spanning the 5' end of the CX3CR1-Cre-ERT2 using with OneTaq polymerase with 68oC annealing.

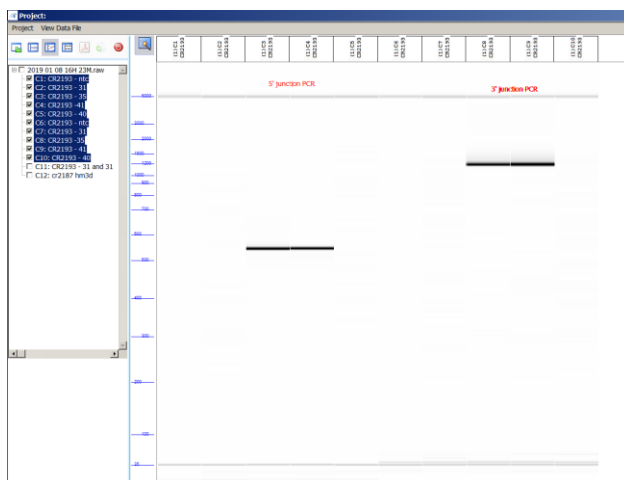
<u>Primer Name</u>	<u>Primer Sequence (5' to 3')</u>
CX3CR1 F103289	TCAGGGTGGCCCATAACCAC
CREERT2 R246	ACCGCGGCCTGAAGATATAG

These oligos produce a 550 bp amplicon spanning the 5' end of the CX3CR1-Cre-ERT2 using with OneTaq polymerase with 68oC annealing.

<u>Primer Name</u>	<u>Primer Sequence (5' to 3')</u>
CreERT2 F1206	CAGTGAAGCTTCGATGATGGGC
CX3CR1 R104157	GCCAGATTCCTGCAGGACCT

These oligos produce a 1553 bp amplicon spanning the 3' end of the CX3CR1-Cre-ERT2 using with OneTaq polymerase with 68oC annealing.

Gel image from CR2193 validation of 5' (R246) and 3' junctions.



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