

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

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 530-754-MMRRC

NAME OF PCR: B6.129X1-*Ntrk2*<sup>tm1Lfr</sup>/Mmucd MMRRC # 000187-UCD

**Protocol:**

Reagent/ Constituent	Volume (μL)
Water	17.6
10x Buffer (contains 15mM MgCl <sub>2</sub> )	2.5
MgCl <sub>2</sub> (stock concentration is 25mM)	1.7
dNTPs (stock concentration is 10mM)	0.5
Primer 1 (stock concentration is 25μM) TrkB n2	0.5
Primer 2 (stock concentration is 25μM) TrkB c8	0.5
Primer 3 (stock concentration is 25μM) TrkB c7	0.5
Taq Polymerase 5Units/μL	0.2
DNA Sample	1.0
<b>TOTAL VOLUME OF REACTION:</b>	<b>25μL</b>

**Comments on protocol:**

*TrkB* is an older synonym of the gene *Ntrk2*. Use a simplex reaction with TrkB n2 and TrkB c7 to amplify only the transgenic allele.

**Strategy:**

Steps	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting <span style="float:right">HOT START? <input type="checkbox"/></span>	94	3:00	1
2. Denaturation	94	1:00	} 35x
3. Annealing } steps 2-3-4 will cycle in sequence	63.5	1:30	
4. Elongation	72	1:30	
5. Amplification	72	7:00	1
6. Finish	4	∞	n/a

**Primers:**

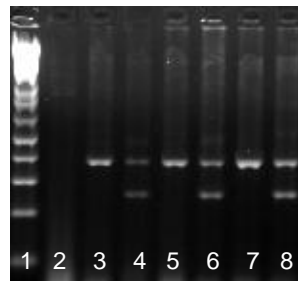
Primer Name	Nucleotide Sequence (5' - 3')
1: TrkB n2	ATG TCG CCC TGG CTG AAG TG
2: TrkB c8	ACT GAC ATC CGT AAG CCA GT
3: TrkB c7	GAT GAT TTC TAG CCT TTT CTG G

**Electrophoresis Protocol:**

% Agarose: 2 mV: 80

Estimated Running Time (min): 90

Expected Bands	Genotype
369 bp	WT +/+
245 bp	Tg/+



Lanes:  
 1. 1Kb+ ladder  
 2. H<sub>2</sub>O  
 3. Wild-type +/+ sample  
 4. Tg, hemizygous sample  
 5 & 7. WT  
 6 & 8. Tg