

**GENOTYPING BY PCR PROTOCOL**  
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 530-754-MMRRC

NAME OF PCR: B6;129S4-*Neurod1*<sup>tm1Jle</sup>/Mmucd MMRRC # 000367-UCD

**Protocol:**

Reagent/ Constituent	Volume ( $\mu$ L)
Water	10.775
10x Buffer (contains / without 15mM MgCl <sub>2</sub> )	2.5
MgCl <sub>2</sub> (stock concentration is 25mM)	1.7
Betaine (stock concentration is 5M) <i>Optional</i>	6.5
dNTPs (stock concentration is 10mM)	0.5
DMSO <i>Optional</i>	0.325
Primer 1 (stock concentration is 20 $\mu$ M) NDW-1	0.5
Primer 2 (stock concentration is 20 $\mu$ M) NDW-2 (JL-25)	0.5
Primer 3 (stock concentration is 20 $\mu$ M) NDW-3	0.5
Taq Polymerase 5Units/ $\mu$ L	0.2
DNA extracted with <input checked="" type="checkbox"/> NaOH <input type="checkbox"/> Proteinase K <input type="checkbox"/> Other:	1.0
<b>TOTAL VOLUME OF REACTION:</b>	<b>25<math>\mu</math>L</b>

**Comments on protocol:**

- Use Touch-Down cycling protocol-first 10 cycles anneal at 65° C decreasing in temperature by 1.0 ° C; next 30 cycles anneal at 55° C.
- Betaine/DMSO is standardized due to high GC content in promoter regions and protocol may be tested without. Also, may adjust MgCl<sub>2</sub> to increase reaction or decrease non specific amplifications.

**Strategy:**

Steps	HOT START? <input type="checkbox"/>	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting		94	5:00	1
2. Denaturation		94	0:15	
3. Annealing	} steps 2-3-4 will cycle in sequence	65 to 55 (↓1°C/cycle)	0:30	10x
4. Elongation		72	0:40	
5. Denaturation		94	0:15	
6. Annealing	} steps 5-6-7 will cycle in sequence	55	0:30	30x
7. Elongation		72	0:40	
8. Amplification		72	5:00	1
9. Finish		15	∞	n/a

**Primers:**

Name	Nucleotide Sequence (5' - 3')
1: NDW-1	ACC ATG CAC TCT GTA CGC ATT
2: NDW-2 (JL-25)	GAG AAC TGA GAC ACT CAT CTG
3: NDW-3	AAA CGC CGA GTT AAA GCC ATC

**Electrophoresis Protocol:**

Agarose: 2% V: 100 Estimated Running Time: 60 min

Expected Bands	Genotype
433 bp	WT +/-
643 / 433 bp	HET +/-
643 bp	KO -/-