

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

mmrrc@ucdavis.edu
530-754-MMRRC

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

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Strain Name		MMRRC Stock Number
Modvl4 Subcongenic 2		42091

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NAME OF PCR: Modv14 Sub2 genotyping using SSLP MIT markers MMRRC: 0-UCD

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

Reagent/Constituent	Volume (µL)
Water sddH2O	4
2x Buffer GoTaq Green (Promega); contains Taq DNA polymerase	10
MgCl ₂ (stock concentration is 3mM) included in Buffer	
Betaine (stock concentration is 5M) <i>Optional</i>	
dNTPs (stock concentration is 10mM) 2X GoTaq Buffer contains 400uM dNTPs	
DMSO <i>Optional</i>	
Primer 1. (stock concentration is 10µM)	2
Primer 2. (stock concentration is 10µM)	2
Primer 3. (stock concentration is 10µM)	
Primer 4. (stock concentration is 10µM)	
Taq Polymerase 5Units/µL included in Buffer	
DNA (50-200ng/ µL) extracted w/ "Promega Wizard SV Genomic DNA Purification System"	2
<i>The total volume is auto-calculated based on volumes entered, right click the total and update field to show/recalculate the total volume.</i>	
TOTAL VOLUME OF REACTION:	20.000 µL

Comments on protocol:

- One primer pair is used for each reaction; to genotype the whole Modv14 region, we used 14 SSLP MIT markers that are evenly distributed within the Modv14 55 Mb interval. Expected band size below indicate MOLF/Ei background in Modv14 region.
- Genotyping PCR was carried out using Promega GoTaq Green Master Mix; concentrations listed in the table above are approximations of Master Mix contents

Strategy:

Steps	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting HOT START? ☐	94	1:00	1
2. Denaturation	94	0:15	
3. Annealing steps 2-3-4 cycle in sequence	51	0:15	30x
4. Elongation	72	0:45	
5. Amplification	72	10:00	1
6. Finish	4	∞	n/a

Primers

Name	Nucleotide Sequence (5' - 3')	Polyacrylamide: 8% V: 150		
1. D15MIT51	ACCTTGTTTCAGAACGAATAAATG	Estimated Running Time: 120 min.		
2.	GATGCTCAGAGAGTGCACCA	Primer Combination	Band (bp)	Genotype
3. D15MIT81	ATGAAACACAAACGTAAAGTTATACA	D15MIT51	124	MOLF/MOLF
4.	ATCTTCTTTTATGCCTGTGTAGGC	D15MIT81	132	MOLF/MOLF
5. D15MIT130	CATATTTTGCAATTTTGTAGTATAGGC	D15MIT130	185	MOLF/MOLF
6.	CAACACAGAAATAAAAGTGAGAGAGG	D15MIT252	147	MOLF/MOLF
7. D15MIT252	CTTCAAACATGTTATCATTGTCACA	D15MIT267	310	MOLF/MOLF
8.	CTTCTGTATTCACAGGTGCTCG	D15MIT94	160	MOLF/MOLF
9. D15MIT267	CTGAAACTTCCAAGTCACTCTG	D15MIT204	188	MOLF/MOLF
10	CAAGCAGAAAAGCAAACCAG	D15MIT49	158	C3H/C3H
11 D15MIT94	CCACTTCTGACCTTCACATGT	D15MIT24	246	C3H/C3HF
12	TGCCTATGTATGTGGTATGTATGA	D15MIT86	196	C3H/C3H
13 D15MIT204	GACTTTGTCTTATGTGATATGGTGTG	D15MIT59	116	C3H/C3H
14	TTCTAGACTTGCTAACATGAAACA	D15MIT195	186	C3H/C3H
15 D15MIT49	TGTACTCATGTGTACAGACCCAC	D15MIT27	208	C3H/C3H
16	GGTGACCCTGTGAAAACGAT	D15MIT257	122	C3H/C3H
17 D15MIT24	TCAACAGCTTTATTAGGAACCTCC			
18	CTTACCATCCCTTGCCAAAA			
19 D15MIT86	TACACTTATCTGTTGGAACATCCC			

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20	ATATAGATAAACACACGCACACCA			
21 D15MIT59	ATGGCTATGATTAAGAAGACAGCC			
22	ATACCATGAATGAGGGTTCACC			
23 D15MIT195	CAAGAGAGATTCACTGACCACG			
24	GATGCAAGAGCCTCACCAG			
25 D15MIT27	GTTAGGCCCTGTGGTTTTGA			
26	ATCCTTTCTAAACATGGACTTTGG			
27 D15MIT257	AAGAAGGTCTATGAAGTCTGCAGG			
28	CTTCACCAGATGCCATGCTA			

Protocol / Gel Comments:

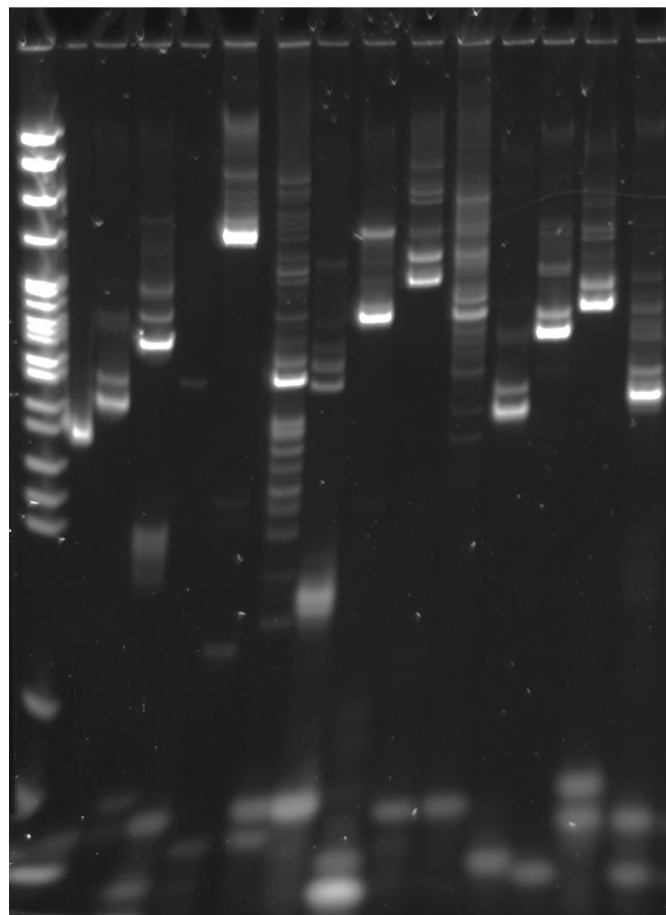
PCR products are resolved on a 8% polyacrylamide gel prepared as follows:

- H₂O: 13.25 mL
- 10X Tris-Borate-EDTA (TBE) buffer: 2.0 mL
- 40% stock acrylamide solution (Acrylamide:Bis-acrylamide 29:1): 4.0 mL
- 10% stock Ammonium persulfate (APS): 0.25 mL
- Tetramethylethylenediamine (TEMED): 0.025 mL

Molecular Weight Marker used is pBR322 MspI from New England Biolabs

Gel run: 150V, 75 min

Gel pictures:



Lane

- 1 – pBR322 MW marker
- 2 – D15MIT51
- 3 – D15MIT81
- 4 – D15MIT130
- 5 – D15MIT252
- 6 – D15MIT267
- 7 – D15MIT94
- 8 – D15MIT204
- 9 – D15MIT49
- 10 – D15MIT24
- 11 – D15MIT86
- 12 – D15MIT59
- 13 – D15MIT195
- 14 – D15MIT27
- 15 – D15MIT257