

GENOTYPING PROTOCOL

MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

mmrrc@ucdavis.edu

530-754-MMRRC

Protocol Name: C57BL/6N-Dpp4em1(DPP4)Mbp/Mmucd

Stock: 068166-UCD

Protocol: GoTaq® G2 Colorless Master Mix(Promega)

Reagent/Constituent	Volume (µL)
Water	4.5
GoTaq® G2 Colorless Master Mix,2X	7.5
Primer 1. (stock concentration is 20µM) IVF	0.5
Primer 2. (stock concentration is 20µM) IVR	0.5
Primer 3. (stock concentration is 20µM) kiF	0.5
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.5
TOTAL VOLUME OF REACTION:	15.00 µL

Comments on protocol:

- Protocol may work with other DNA extraction methods.

Strategy:

Steps	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting HOT START? <input type="checkbox"/>	94	2:00	1x
2. Denaturation	94	0:10	
3. Annealing steps 2-3-4 cycle in sequence	65 (↓1°C/cycle)	0:30	10x
4. Elongation	68	2:00	
5. Denaturation	94	0:15	
6. Annealing steps 5-6-7 cycle in sequence	55	0:30	25x
7. Elongation	68	2:00 (↑20sec/cycle)	
8. Finish	4	∞	n/a

Primers:

Electrophoresis Protocol:

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5%	V: 90
1. C19-Dpp4-IVF	GCACTCTCATTTGGCATCAGACG	Estimated Running 90 min.	
2. C19-Dpp4-IVR	CCAGTCTCTTCAGCCATAACCCAG	Primer Combination	Band (bp)
3. C19_Dpp4-kiF	TAGGTGCTACAAAACACAGCAAGG	1 & 2	441
		1 & 3	542
			Genotype
			wildtype
			mutant

Allele Description: CRISPR targeted humanized DPP4 KI locus and mouse Dpp4 ([ENSMUSG0000035000.9](#)) KO. Human CDS consisting of exons 2*-26 was inserted into mouse exon 2 ([ENSMUSE0000062442](#)) beginning at the 11th amino acid into zygotes. The mouse exon 2 is replaced with human exon 2 CDS/h3'UTR (human 3' untranslated region) in the HR template to prevent potential downstream splicing or RNP re-cleavage (4 engineered silent mutations in human CDS). This production targeted directly into zygotes to produce a founder and backcross to N1 to create germline Het animals. Long range PCR products are fully sequenced.

