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

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530-754-MMRRC

NAME OF PCR: C57BL/6-Itgb2^{m1Btlr}/Mmcd, (Joker) MMRRC # 016138-UCD

Protocol:

| Reagent/ Constituent | Volume (µL) |
|--|-------------|
| Water | 12.675 |
| 10x Buffer (contains 15mM MgCl ₂) | 2.5 |
| Betaine (stock concentration is 5M) Optional | 6.5 |
| dNTPs (stock concentration is 25mM) | 0.5 |
| DMSO Optional | 0.325 |
| Primer 1 (stock concentration is 20µM) Joker PCR F2 | 0.5 |
| Primer 2 (stock concentration is 20µM) Joker PCR R2 | 0.5 |
| Taq Polymerase | 0.5 |
| DNA sample extracted with ☐ NaOH ☐ Proteinase K ☐ Other: Any | 1.0 |
| TOTAL VOLUME OF REACTION: | 25µL |

Comments on protocol:

- PCR products are verified to contain the correct amplicon size by running ~10µl of the reaction on a gel and the remaining 15µl purified via column based PCR purification method for sequencing.
- 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 and DMSO have been standardized due to high GC content. Protocol may be tested without. Also, may adjust MgCl₂ to increase reaction or decrease non specific amplifications.

Strategy:

| Steps | Temp (°C) | Time (m:ss) | # of Cycles |
|---|-----------------------|-------------|-------------|
| 1. Initiation/Melting HOT START? | 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 | J |
| 5. Denaturation | 94 | 0:15 | |
| 6. Annealing steps 5-6-7 will cycle in sequence | 55 | 0:30 | 30x |
| 7. Elongation | 72 | 0:40 | J |
| 8. Amplification | 72 | 5:00 | 1 |
| 9. Finish | 15 | ∞ | n/a |

Primers:

| Name | Nucleotide Sequence (5' - 3') |
|-----------------|-----------------------------------|
| 1: Joker PCR F2 | CGT ATC TGG GGG GTG TCT GTT TGT T |
| 2: Joker PCR R2 | CCA GTT TGC CAT CAC CAG CAA |
| 3: Joker_Seq3 | GGA AGC CAT CGT CTG TGG CAA |

Electrophoresis Protocol:

Agarose: 2% mV: 80 Estimated Running Time: 90 min

| Primer Combination | Band | Genotype | | |
|--|--------|----------|--|--|
| 1 and 2 | 407 bp | joker | | |
| SNP found at position ~ 42 of sequencing | | | | |

Mutation site (red) and flanking sequence:

WT ttcccAggagga
joker ttcccTggagga