

## Mouse Genotyping by PCR / Tail DNA Extraction

### **I. Protocol for Processing Blood for Flow Cytometry**

1. Place mouse in isoflurane chamber in order to briefly anesthetize.
2. Snip end of tail, and tag each mouse appropriately.
3. Gently squeeze blood from tail, and collect 2-3 drops of blood.
4. Place blood into 50  $\mu$ L of heparin (100 USP Heparin units/mL).
5. After collection, stop bleeding by applying Kwik Stop styptic powder using a moistened cotton applicator to end of tail.
6. After collection of blood samples is completed, wash heparin by adding 150  $\mu$ L of 1X PBS, and spin at 1500 rpm for 5 min.
7. Discard supernatant.
8. Resuspend in 200  $\mu$ L of Red Blood Cell Lysis buffer ( $\text{NH}_4\text{Cl}$  (155 mM) and  $\text{NaHCO}_3$  (10 mM), pH 7.2).
9. Spin and discard supernatant.
10. Resuspend in 200  $\mu$ L of 1X PBS.
11. Spin and discard supernatant
12. Resuspend in 50  $\mu$ L of a 1:50 dilution of Fc block in FACS buffer (0.1% sodium azide in 1X PBS).
13. Incubate on ice for 15-20 min.
14. Wash by adding 150  $\mu$ L of FACS buffer.
15. Spin and discard supernatant.
16. Resuspend in 50  $\mu$ L of the appropriate antibody mix for each strain.
17. Incubate on ice for 45- 60 min.
18. Wash by adding 150  $\mu$ L of FACS buffer.
19. Spin and discard supernatant.
20. Resuspend in 200  $\mu$ L of FACS buffer.

Samples are now ready for flow cytometry.

## II. Protocol for Tail Extraction

1. Snip 0.5 cm of tail and place in 1.5 mL clean labeled eppendorf tube.
  - (Store in -20°C if necessary)
2. Add 500 µL of tail cocktail (2% SDS, 50 mM TRIS pH 8.0, 20 mM EDTA pH 8.0, Water) and 50 µL of Proteinase K to each tube.
3. Incubate tubes in 60°C circulating waterbath overnight.
4. Place tubes at 4°C until ready to extract DNA.
5. Add 200 µL of saturated NaCl to each tube and invert to mix.
6. Spin at 4°C for 20 min at 13,000 rpm.
7. Transfer supernatants into clean, labeled eppendorf tubes.
8. Add cold 100% ethanol to the top of each tube and invert to mix until DNA strands precipitate out.
9. Spin at 4°C for 10 min at 13,000 rpm.
10. Discard supernatant and resuspend pellets in 300 µL ddH<sub>2</sub>O and briefly vortex.
11. Incubate tubes in 60°C water bath for 1 hour to resolubilize DNA.
12. Store at -20°C.

## III. Protocol for PCR Amplification

1. Mix PCR reaction components together as described in protocol for given gene.
2. Visualize DNA bands with agarose gel electrophoresis
  - a. Add solid agarose to 1X TBE buffer to desired concentration.
  - b. Microwave to melt the agarose in the buffer, and add 10 µL of 1% ethidium bromide solution per 100 mL and pour into gel box.
  - c. Let the gel solidify and load samples with appropriate base pair markers and loading dye.
  - d. Run at approximately 100V for 60 min.
  - e. Image and record results.

## PCR Protocol for STAT<sup>fl</sup> PCR

### Master Mix

Reagent	uL/ sample
5X PCR Buffer	5.0
25 mM MgCl <sub>2</sub>	2.0
10 mM Dntp	0.5
10 μM Forward Primer 5'-GCA TGA CAT GAT CAG CAT TGC- 3'	1.0
10 μM Reverse Primer 5'- ACT GAC GTC AAC CAA GCC TG- 3'	1.0
Platinum Taq DNA polymerase (Invitrogen)	0.2
Water	38.3
DNA	2.0

### Cycling Parameters:

Step 1: 94°C          2 min

Step 2: 94°C          20 sec

        65°C          15 sec          For 10 cycles

        68°C          10 sec

Step 3: 94°C          20 sec

        60°C          15 sec          For 28 cycles

        72°C          10 sec

Step 4: 72°C          2 min

Step 5: 4°C          hold

## PCR Protocol for Lambda Receptor<sup>fl</sup> PCR

### Master Mix

Reagent	uL/ sample
5X PCR Buffer	5.0
25 mM MgCl <sub>2</sub>	2.0
10 mM Dntp	0.5
20 μM Forward Primer 5'- GGC CGC CCA AAT ATA AAC C - 3'	1.0
20 μM Reverse Primer 5'- TCT GAC ATC CGC TCA GCA CCA A - 3'	1.0
Platinum Taq DNA polymerase (Invitrogen)	0.2
Water	38.3
DNA	2.0

### Cycling Parameters:

Step 1: 94°C          5 min

Step 2: 94°C          30 sec  
          48°C          30 sec          For 35 cycles  
          72°C          30 sec

Step 3: 72°C          10 min

Step 4: 4°C          hold