Genotyping of Mouse Tail DNA via PCR

I. Mouse tailing

[Pups are tailed (for DNA) and toed (for identification) between 8-14 days of age.]

- A. Remove tail sample of approximately 0.25 inches by pinching the tip of the tail to expel blood and cutting with scissors.
- B. Place tail sample in 1.5 mcf tube for digestion.
- C. Scissors is washed liberally with 75% EtOH between cuts to prevent cross contamination.

II. Tail lysis and sample preparation

Preparation of tail lysis buffer

Final concentration [100mM Tris-HCl, pH8.8; 5mM EDTA, pH8.0; 0.2%SDS; 200mMNaCl]

- Just prior to use, add proteinase K (Stock conc= 20mg/mL) to the buffer to achieve a final concentration of 100 ug/mL.
- A. Add 500 uL of tail lysis buffer to each tube and vortex briefly to mix.
- B. Tails are incubated o/n at 55C with agitation (250rpm on a heated shaker).
- C. The following morning, tubes are vortexed and centrifuged [5 min x 14000 rpm@RT] to pellet debris.
- D. Dilute lysis 1:11 in sterile H₂0 for use in PCR analysis.

III. PCR

A. Primer sequences and product size

GENE	Primer name	Primer sequence	Amplicon size
nfi-a	I2B	TGCTGTGTTCTGGTCAGTCAAG	405 bp
	I2CC	CAAAGCAAATCTCCATGCTCGG	
neo	neo57	GGAGAGGCTATTCGGCTATGAC	315 bp
	neo371R	CGCATTGCATCAGCCATGATGG	
sry (sex)	sry1	AACAACTGGGCTTTGCACATTG	166, 146 bp
	sry2	GTTTATCAGGGTTTCTCTCTAGC	(doublet)

B. PCR Reaction mixture

	COMPONENT		[STOCK]	[FINAL RXN]
1.	10X Taq Buffer		10X	1X
		Tris-HCl	100mM	10mM
		MgCl ₂	15mM	1.5mM
		KCl	500mM	50mM
		BSA	0.2mg/mL	0.02mg/mL
2.	Multiplex primer mix		10X	1X
	•	I2B	2 uM	200 nM
		I2CC	2 uM	200 nM
		neo57	2 uM	200 nM
		neo371R	2 uM	200 nM
		sry1	2 uM	200 nM
		sry2	2 uM	200 nM
3.	dNTP mix		1 mM each	200 uM each
	(dA,dT,dG,dC)			
4.	taq polymerase		5 U/uL	0.5 Units
	(Life Technologies			
	Cat# 10966026)			
	q.s. to final volume			19.0 uL
	With H ₂ 0			
	DNA (diluted lysate)			1.0 uL
				20.0 uL rxn

C. PCR Parameters

- 1. 94C for 4 min (initial melt)
- 2. 94C, 60C, 72C x 33 cycles (amplification)
- 3. 72C for 9 min (final extension)

