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SAPAP3 Genotyping Protocol

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Protocol status: Working

We use this protocol and it's working



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Abstract

Protocol for SAPAP3 KO Mouse Model Genotyping from Dr. Lisa Gunaydin's Lab, University of California, San Francisco

1 **Cut the tails.**

To cut the tails and mark the animals, keep in mind to:

- cut as little amount of the tail as possible;
- always clean the scissors properly in between animals (EtOH and bleach);
- tattoo the animals at the same time as cutting the tails;
- use iso if needed.

2 **Digest the tails.**

Incubate the tails O/N at 55 C (water bath) in 200 ul 1x PBND buffer with Proteinase K at 0.25 mg/ml. Vortex before.

$200\text{ul} * (54+1) = 11000\text{ul} \rightarrow 11\text{ml PBND}$

$0.25\text{mg/ml} * 11\text{ml} = 2.75\text{mg} \rightarrow 0.00275\text{g Proteinase K}$

PBND (250 ml 1x, stored at 4 C)

2.5 ml 1M TrisHCl pH 8.3

1.125 ml IGEPAL

1.125 ml Tween 20

0.935 g KCl

0.125 g $\text{MgCl}_2 \cdot 6\text{H}_2\text{O}$

0.025 g Gelatin

Fill remainder with H_2O

*Gelatin will dissolve only after heated

3 **Inactivate Proteinase K.**

After taking the tails out of the bath, see if they are digested. If so, vortex and boil them for 10 min. Vortex again and centrifuge at max speed for 5 min (table top centrifuge, something like 14,000 rpm).

4 **PCR reaction.**

We use the KAPABiosystems HotStart ReadyMix (2x) which contains DNA Polymerase, rxn buffer, dNTPs and MgCl_2 .

Pipette 24 ul of Master Mix into each PCR tube

Add 1 ul of DNA from each sample into each tube

Sap3 KO F1, Forward primer, 5' - 3': ATTGGTAGGCAATACCAACAGG

Sap3 KO R1, Reverse Primer 5' - 3': GCAAAGGCTCTTCATATTGTTGG

Extra Primer, TKF2: CTTTGTGGTTCTAAGTACTGTGG



	A	B	C	D	E
		1 rxn	MM SAPAP (__ rxns)	Primer name	Total Rxns
	Forward primer	1		Sap3 KO F1	
	Reverse primer	1		Sap3 KO R1	
	Extra primer	1		TKF2	
	Rxn mix	12.5			
	H2O	8.5			
	DNA	1.0			
	PCR program		Sap3		

PCR Program for SAPAP3 Genotyping:

1. 95C for 4 minutes
2. 30 times:
 - 94C for 30 seconds
 - 62C for 30 seconds
 - 72C for 60 seconds
3. 72C for 5 minutes
4. 4C hold

Expected band lengths:

WT: one band at ~147 bp

Het: one band at ~147 bp and one band at ~222 bp

KO: one band at ~222 bp

5 Run the 1% agarose gel

200 ml TAE

2g agarose

2ul sybr safe gel stain

24ul DNA per well. 5ul DNA ladder before each new line