primer	Sequence(5'-3')	Size	target gene	accession
rUnc5h3-15	CAGATCAGCTCTGCTTTCCA	238bp	Unc5h	NM_199407.1
rUnc5h3−16	ATTTTCAAATTGGGCAGTGG	Zoobb		
comment				

PCR condition	าร								
Taq polymeras	se	BIOTAQTM HS	OTAQTM HS DNA Polymerase (Bioline Reagents Ltd., UK)						
Thermal cycle	r	PC-808(ASTEC)							
PCR buffer		Ampdirect Plus							
		first denature	denature	anneal	extension	cycle	final extension		
PCR	°C	94	94	55	72	35	72		
	min	3	0.5	1	0.75		3		
①PCR method; Blood was applied to the FTA®card (GE Healthcare UK Ltd.,UK) and dried. 1.5mm disc was removed from the bloody-stanined region on FTA®card. Untreated sample discs were placed directly in 15 μ L PCR mixture containig 1 × Ampdirect®Plus, 0.2 μ M each primer and 0.4 units of BIOTAQTM HS DNA Polymerase. ②RFLP of PCR products;Unc5c cvd mutaion causes BstEII cutting site in the Unc5c +/+ genome.									

