Supplementary table S1：Primer set for genotyping

Primer name	Sequence (5’-3’)	
Snai1-flox/flox	Forward	CTGCCAGGTGGGAAGGACT
	Reverse	CAAGGACATGCGGGAGAAGGT
Lrat-Cre	Forward	TGAGCCAAGCACTTTGGCTTC
	Reverse	TCACATCCTCAGGTTCAGCAGG
		AACAGGATGTCAGGCATTAGATGG
Snai1-LacZ	Forward	GCAGCCTCTGTTCCACATACACTTCA
	Reverse	GTCTGTTGTACCTCAAAGAAGGTGGC