

Table I: Mendelian distribution of *Icap-1* progeny.

Genotype Crossing	Wild-type	Heterozygous	<i>Icap-1</i> null
(+/-) x (+/-)			
Expected (%)	25	50	25
P0 (%) (n=128)	27	49	24
P28 (%) (n=213)	27	53	20
(+/-) x (-/-)			
Expected (%)	-	50	50
P28 (%) (n=34)	-	60	40

P, postnatal day