

Table S1. Numbers and genotype frequencies of embryos recovered from Sox9-IRES-Cre;*Efnb2*^{+/*LacZ*}
X *Efnb2*^{fl^{ox}/fl^{ox}} matings

Gestational day at collection	No. of litters	<i>Efnb2</i> ^{+/<i>fl^{ox}</i>}	<i>Efnb2</i> ^{<i>LacZ</i>/<i>fl^{ox}</i>}	Sox9-IRES-Cre ⁺ ; <i>Efnb2</i> ^{+/<i>fl^{ox}</i>}	Sox9-IRES-Cre ⁺ ; <i>Efnb2</i> ^{<i>LacZ</i>/<i>fl^{ox}</i>}	Totals
E9.5/10.5/11.5	14	41 (.30)	30 (.22)	27 (.20)	37 (.27)	135
E12.5/13.5/14.5	33	72 (.29)	75 (.30)	71 (.28)	33 * (.13)	251
E15.5/16.5/19	35	81 (.27)	95 (.32)	83 (.28)	42 * (.14)	301
Totals	82	194 (.28)	200 (.29)	181 (.26)	112 ** (.16)	687

Absolute numbers and genotype frequencies (in parentheses) are shown. P values for Chi-square tests are: * P < 0.01; and **P < 0.001.

