

Table S1. Distribution of the genotypes from *Bicc1*^{+/-} × *Bicc1*^{+/-} crosses

<i>Bicc1</i> genotype	E12.5 and earlier (%)	E14.5 (%)	E15.5 (%)	E16.5 (%)	E18.5 (%)	P0/P1 (%)	P2 and older (%)
+/+	23	35	33	23	28	34	32
+/-	55	40	48	57	57	56	60
-/-	23	25	19	20	15	10	8
<i>n</i>	22	20	217	70	146	105	191