

**Supplemental Table 1****Genotyping of offspring from miR-499<sup>+/−</sup> intercrosses**

<u>Genotype</u>	<u>+/+</u>	<u>+/-</u>	<u>-/-</u>
Observed no:	30	41	26
Observed %:	31	42	27
<u>Expected %:</u>	<u>25</u>	<u>50</u>	<u>25</u>

**Supplemental Table 2****Genotyping of offspring from miR-208b<sup>+/−</sup> intercrosses**

<u>Genotype</u>	<u>+/+</u>	<u>+/-</u>	<u>-/-</u>
Observed no:	28	42	19
Observed %:	31	47	22
<u>Expected %:</u>	<u>25</u>	<u>50</u>	<u>25</u>

**Supplemental Table 3****Genotyping of offspring from miR-499<sup>+/−</sup>/miR-208b<sup>+/−</sup> intercrosses**

<u>Genotype</u>	<u>1</u>	<u>2</u>	<u>3</u>	<u>4</u>	<u>5</u>	<u>6</u>	<u>7</u>	<u>8</u>	<u>9</u>
miR-499	+/ <sup>+</sup>	+/ <sup>+</sup>	+/-	+/-	-/-	-/-	-/-	+/-	+/ <sup>+</sup>
miR-208b	+/ <sup>+</sup>	+/-	+/ <sup>+</sup>	+/-	+/ <sup>+</sup>	+/-	-/-	-/-	-/-
Observed no:	6	4	8	10	3	4	6	5	6
Observed %:	11.5	7.6	15.4	19.2	5.8	7.7	11.5	9.6	11.5
<u>Predicted %:</u>	<u>6.25</u>	<u>12.5</u>	<u>12.5</u>	<u>25.0</u>	<u>6.25</u>	<u>12.5</u>	<u>6.25</u>	<u>12.5</u>	<u>6.25</u>