Supplementary Data

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A141R      GATGCTGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
PM4808F    GATGCTGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A141F      GATGCTGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
PM4913R    GATGCTGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
PM4913F    GATGCTGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A291R      GATGCTGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A291F      GATGCTGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A66F       GATGCCGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A1013F     GATGCCGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A185R      GATGCNGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A139R      GATGCNGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
A139F      GATGCNGAGACAGTGATGAGCCAGACTGTGAGTCCCCCCACTGTGACCTGAGCCCCCGAG 60
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A141R      CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
PM4808F    CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A141F      CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
PM4913R    CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
PM4913F    CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A291R      CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A291F      CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A66F       CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A1013F     CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A185R      CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A139R      CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
A139F      CCCACCCCAGCCCCGGGTCCACTCTCTTCACCTGCTCCTGTGGCACTGACTTCTGCAATG 120
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A141R      CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
PM4808F    CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A141F      CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
PM4913R    CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
PM4913F    CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A291R      CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A291F      CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A66F       CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A1013F     CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A185R      CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A139R      CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
A139F      CCAATTACAGCCATCTGCCTCCTCCAGGGAACCCTGGGACCCCTGGCTCCCAGGGTCCCC 180
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A141R      AAGCTATCCCAG 192
PM4808F    AAGCTATCCCAG 192
A141F      AAGCTATCCCAG 192
PM4913R    AAGCTATCCCAG 192
PM4913F    AAGCTATCCCAG 192
A291R      AAGCTATCCCAG 192
A291F      AAGCTATCCCAG 192
A66F       AAGCTATCCCAG 192
A1013F     AAGCTATCCCAG 192
A185R      AAGCTATCCCAG 192
A139R      AAGCTATCCCAG 192
A139F      AAGCTATCCCAG 192
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Figure1: Multiple sequence alignment of *MISRII* exon3 of animals in Figure 2. The PMDS mutation is highlighted. N represents the CT heterozygous condition. F and R represent forward and reverse reads respectively.