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Allelic- and parental-specific expression of the KCNJ11 gene in cattle

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Allelic variation and allelic parental origin effects on expression of genes of commercial interest in cattle can be caused by genetic and epigenetic mechanisms. The knowledge of these mechanisms may contribute to infer about gene effects on the next generation in animal selection programs. Considering this scope, we analysed the expression mechanism of KCNJ11 (potassium inwardly-rectifying channel, subfamily J, member 11; Gene Bank ID 532060), a candidate gene for meat tenderness identified previously by our research group. The two SNPs c.1526C>T (Gene Bank ID ss# 537718973) and c.2342T>C (ss# 537718995) in the KCNJ11 gene were used to investigate allelic-specific and parental origin expression in Nelore muscle tissue. Differential allelic expression for the SNP c.2342T>C was found and it was influenced by parental origin and genotypes of the SNP c.1526C>T. KCNJ11 total mRNA abundance showed different patterns according to parental origin of alleles. Eight miRNA target sites were found for the SNP c.2342T>C, two of them being specific for the sequence of allele C and two for the allele T. It was evidenced that bovine KCNJ11 gene shows differential allelic and parent-of-origin expression. This knowledge may contribute to marker assisted selection, since it will allow for more precise prediction of additive effects on quantitative traits. Further studies are needed to identify the mechanisms that control its differential and parental allelic expression identified in this study.