Supplementary

Table S1 Sequences of the primers used for the amplification and sequencing of rs576016632

Gene	Locus/Version	SNP	Alleles	Primer Information
SPEG	NM_005876.5	rs576016632	G>A,C,T	GGTAACCTGACTCCCTGCAGTGGGTCGTGTGGGGGCCAGGCACACT ACGGAGGGGGAAAGCCTGGAACAAATACCGAGGGACTCCCTTAAGCCG GGCCGGCGATGGGGGCTCCTGGAGGGAGAGAAGGAGCCAAGTGGA GTCAAGTCCCTCCCCTGCTGCCCCCTCCCT*CCACGGCTCCCTCGCA* *ACCC*GAGCCGGGGGGGCCTAAAAATAGCCCCCAGGCGCAATCGCCGG CCGCCCCGGTGACCTTCTGGGTAGCACAGGCCGAAGGCGGGCGGG CAGCAGGAAGGCAGGCCGCCGGCCCCCCAGACTTGTCTCCTAGGGC ACCGTCCCGCGGGTGCCCCCGTGGCCGCCCAGTTCCGGCGTCCCC CCAGCCCAGCTCTCAGTGGCCATGCAGAAAGCCCG*GGGCACGCGA GGCGAGGATGCGGGCACGAGGGCACCCCCCAGCCCCGGAGTGCCC CCGAAAAGGGCCAAGGTGGGGGCCGGCGGCGGGGCTCCTGTGGCC GTGGCCGGGGCGCCAGTCTTCCTGCGGCCCCTGAAGAACGCGGCG GTGTGCGCGGGCAGCGACGTGCGGCTGCGGGTGGTGGTGAGCGGG ACGCCCCAGCCCAGCCTCCGCTGGTTCCGGGATGGGCAGCTCCTGC CCGCGCCGGCCCCCGAGCCCAGCTGCCTGTGGCTGCGGCGCTGCG GGGCGCAGGACGCCGGCGTGTACAGCTGCATGGCCCAGAACGAGC GGGGCCGGGCCTCCTGCGAGGCGGTGCTCACAGTGCTGGAGGTCG GAGGTAAAGGGCAGGTGGGGGCCGCGCCC

>hg19_dna range = chr2:220299313-220300113 5'pad =400 3'pad =400 strand = + repeatMasking = none Ex1. The underline sequence represents the amplification primer; the italics sequence represents the sequencing primer; *, represents the target site. NM, Nucleotide Molecular; SNP single nucleotide polymorphism.

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