Supplementary Table S1. Primers used to amplify the CARD14 regions to genotype the selected SNPs.

No.	Primer	Sequence	Tm	Length	Hits	Product
1	Forward	CTGGATTTGCTGAAGACTCG	56.5	20	1	
	Reverse	ATCCAACCCCAATACAGTGA	56.1	20	1	310
2–10	Forward	CCTGCCCACCTATTACCTC	57.1	19	1	
	Reverse	GGCAGGGAAGTGTTAGACAA	57.8	20	1	504
	Reverse	ATTGCTATAGTGCAGCGAGA	56.4	20	1	330
11	Forward	ATCAAGGTCCCAAGCATTG	58.5	19	1	
	Reverse	ATAGGACACTCACCCTCTGC	56.7	20	1	392
	Reverse	GATGAAGATGCCCGTGAG	57.6	18	1	674
	Reverse	GACCCGGTGGATGAAGAT	58.2	18	1	683
12–15	Forward	GCAGAGGGTGAGTGTCCTAT	56.7	20	1	
	Reverse	ATGAGCAGAACGCTTTACG	56.7	19	1	549
	Reverse	ACGGCACGACTCTCATCTAT	57.3	20	1	527
16	Forward	GCAGAGGCTCGTATCTGTG	57.0	19	1	263
	Forward	GACGGTACACATACCACTCC	55.3	20		525
	Reverse	GGCAGTATCCTTCATGGTG	56.4	20	2	
17	Forward	GCCAATCATCTCCCCTGA	59.5	18	1	
	Reverse	GCCTAAATGAGTGGCATCC	58.1	19	1	304
18	Forward	CTATGGAACTGGGAGTGAGTC	56.3	21	1	
	Reverse	CACAGTTTCTCGCTCAGGAT	58.0	20	1	305