**Supplementary table: Primers employed to identify genetic variants in *CTLA4*.**

|  |  |  |
| --- | --- | --- |
| Region | Forward Primer | Reverse Primer |
| *Exon 1* | AAGTCCTTGATTCTGTGTGGG | ATCACTGCCTTTGACTGCTG |
| *Exon 2* | CCATGAAGGAGCATGAGTTC | ACTGCAATGCAACAGGTGTC |
| *Exon 3* | GGCTACCCATGCAATTTAGG | TGGCCAGTTGATACCACTAAAG |
| *Exon 4* | ATGGTTAGAAGTGGCTTCCG | AGAATTGCCTCAGCTCTTGG |
| *rs231770* | TCAAGTAATCTGTGGGGTGG | AAAATGTCACAGGCCAAGC |

eTable 1. Sequence of each pair of primers used to sequence the four coding exons of *CTLA4* and rs231770 genetic variant.