individuals including 55 cases and

the inhabitants of the

CATT dilution on plasma

after

Cameroonian population with 95.3% (FDR) chance of this locus being associated with HAT (Table 2).

For the three remaining SNPs where the association was not significant after Bonferroni correction, our results

Discussion

In this study we obtain good quality

association of rs8062041 with a CNV is the most plausible mechanism by which this SNP could be associated with protection against HAT. Our results reveal also that the association between

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