Supplementary Table S4. Mutational profile of ZIKV-*NS3m* genome.

A position within ZIKV-*NS3m* genome that contained single nucleotide polymorphism (SNP) at a frequency above 1% of quality-filtered reads, and corresponding amino acid substitution are indicated.

	A	B	C	D	E	F	G	H	I
1	Nucleotide	Base	Mutation	Gene	Amino acid	Reference amino acid	Mutant amino acid	% Mutants	
2	6861	C	A	protein 2K	6	Gln	Lys	1.27	
3									