Supplementary Table S3. Mutational profile of ZIKV-*ICD* genome.

A position within ZIKV-*ICD* genome that contained single nucleotide polymorphism (SNP) at 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	1.37	
3									