

Supplementary Information

Simple reverse genetics systems for Asian and African Zika viruses

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Supplementary Table S1: Primers used for the amplification of the different DNA fragments

Strain	DNA fragment	Primer Forward	Position*	Primer Reverse	Position*
PF/MART	I	CACCCAAGTATCTTCAGCATCT	¶	GCAGCACCATTCTCGATCACC	3407-3428
	II	GAAACATGTGGAACAAGAGGACCA	3354-3377	GAAGCTCCAGCCAAGTAACTTC	7600-7621
	III	GTAAGTGAAGTCTCTACAGCCA	7553-7575	CTCAGGGTCAATGCCAGCGCTT	£
DAK	I	CACCCAAGTATCTTCAGCATCT	¶	GCAGCACCATTCTCTATGACC	3407-3428
	II	CACGTGGAGGAGACATGCGGAAC	3345-3367	GTGTAAATAAGGGAAGCCCTGC	7611-7633
	III	GTAAGTGAAGTCTCCACAGCCA	7553-7575	CTCAGGGTCAATGCCAGCGCTT	£

Primers located respectively at the 5' and 3' terminus of the pCMV (¶) and the HDR/SV40pA (£).

* All positions based on complete PF strain sequence.

Supplementary Note S1: Nucleotide sequence of the DNA fragments used for the ISA procedure

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CACCCAAGTATCTTCAGCATCTTCAATATTGGCCATTAGCCATATTATTCATTGGTTATATAGCATAAATCAATAT
TGGCTATTGGCCATTGCATACGTTGTATCTATATCATAAATATGTACATTTATATTGGCTCATGTCCAATATGACCGCC
ATGTTGGCATTGATTATTGACTAGTTATTAATAGTAATCAATTACGGGGTCATTAGTTCATAGCCCATATGGAGT
TCCCGTTACATAAAGTACGGTAAATGGCCCGCTGGCTGACCGCCCAACGACCCCGCCATTGACGTCAATAAAT
GACGTATGTTCCCATAGTAACGCCAATAGGGACTTTCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCC
CACTTGGCAGTACATCAAGTGTATCATATGCCAAGTCCGCCCTATTGACGTCAATGACGGTAAATGGCCCGCT
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GAATGGTGCTGC

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DAK-III

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MART-II

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