

## Electronic Supplementary Material

# Phylogenetic analysis revealed the central roles of two African countries in the evolution and worldwide spread of Zika virus

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Table S1. Strains of discrepancies among tree topology by visual comparison

Strains	Locations	Time	Lineages		
			Full-length	<i>E</i> gene	<i>NS5</i> gene
ArD157995	Senegal	2001	A1	A1	A1
ArA1465	Cote d'Ivoire	1980	N/A	A2	A1

Notes: A1, African lineage 1; A2, African lineage 2; N/A, not applicable.