



# ZIBRA Project

The Zibra project aimed to sequence a thousand Zika genomes from Brazil to provide important epidemiological information about the spread of this disease. At the time, there was a paucity of complete genome sequences for this virus, in part due to difficulties in transporting material outside of Brazil for sequencing, hindering attempts to determine virus origins, epidemiology and any genomic basis to microcephaly. The project employed a fully-equipped sequencing trailer during a two-month road trip.

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Suitable	No: RNA; infectious disease; sample availability
Organism	Mosquitoes/Zika virus
Substrate	RNA
Full Genome	Yes
Genome size	10 Kb
Reference available	No



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