

Additional Files

Emergence and Potential for Spread of Chikungunya Virus in Brazil

Marcio Roberto Teixeira Nunes^{1*}, Nuno Rodrigues Faria^{2*}, Janaina Mota de Vasconcelos^{1*}, Nick Golding, Moritz U. G. Kraemer, Layanna Freitas de Oliveira, Raimunda do Socorro da Silva Azevedo, Daisy Elaine Andrade da Silva, Eliana Vieira Pinto da Silva, Sandro Patroca da Silva, Valéria Lima de Carvalho, Giovanini Evelin Coelho, Ana Cecília Ribeiro Cruz, Sueli Guerreiro Rodrigues, Joao Lídio da Silva Gonçalves Vianez Jr, Bruno Tardelli Diniz Nunes, Jedson Ferreira Cardoso, Robert B. Tesh, Simon I. Hay, Oliver G. Pybus*, and Pedro Fernando da Costa Vasconcelos*⁴.

‡Correspondence to:

Marcio Roberto Teixeira Nunes, Center for Technological Innovation,
Evandro Chagas Institute, Ministry of Health, Brazil.

Contents

	Page
1. Methodological details	
1.1. Viral Assembly Procedure	3
1.2. Accession numbers for the global CHIKV full genome dataset	3
1.3. Accession numbers for the global CHIKV envelope genome dataset	3
1.4. Predicting CHIKV importation and establishment risk	6
2. Supplementary Tables	
2.1 Table S1	7
2.2 Table S2	8
3. Supplementary Figures	
3.1 Figure S1. Full genome ML phylogeny of CHIKV	9
3.2 Figure S2. Envelope ML phylogeny of CHIKV	10
4. References	11

1. 1. Viral Assembly Procedure

Raw reads generated by the PGM device were used for a *de novo* assembly with Mira v4.0[1]. Relevant parameters used in the assembly were as follows: minimum read length of 100 bases (mrl=100), minimum overlap of 19 bases (mo=19), use of at least 5 reads per contig (mrpc=5) and 5 passes for the whole assembly (nop=5). The closed genomes were annotated using open reading frame predictions from Geneious v 7.0[2] and BLASTX searches[3]. Finished genomes were inspected to check if length, gene number, orientation and non-coding 5' and 3' regions corresponded to what was expected for genomes of the same family.

1.2. Accession numbers for the global CHIKV full genome dataset

Accession numbers for the representative global dataset are as follows. ECSA genotype: HM045792, HM045795, HM045805, AF490259, HM045821, HM045806, HM045809, HM045822, JQ067624, HM045812, HM045784 and HM045823. Asian genotype: HM045803, EF027140, HM045813, HM045788, EF027141, HM045810, HM045814, HM045808, HM045789, HM045802, HM045796, HM045787, HM045797, HM045791, HM045800, HM045790, HE806461, FN295483, FN295484, EU703762, EU703759, EU703760, EU703761, FJ807897, KF318729, KJ451623, KJ451622, CNR20236, CNR20235, KJ451624. Indian Ocean Genotype (Because the Brazilian isolates consistently clustered outside the IOL genotype based on preliminary maximum-likelihood[4] and Bayesian phylogenetic analysis[5], we only kept a sample of the 102 IOL genotype full-length genomes): JF274082, EU372006, GU908223, FR717336, GQ428213, KJ796852, KJ796851, KJ796845, KC862329, FN295485, FN295487, GU189061, HQ456255, HQ456254, HQ456253, HQ456252, HQ456251. West African genotype: HM045816, HM045815, HM045798, HM045785, HM045786, HM045807, AY726732, HM045817, HM045820, HM045819, HM045818.

1.3. Accession numbers for the global CHIKV envelope genome dataset

Accession numbers for the global [envelope](#) dataset are as follows. [KP164567](#), [KP164568](#), [KP164569](#), [KP164570](#), [KP164571](#), [KP164572](#), [AB857841](#), [HM045823](#), [AB857833](#), [AB857825](#), [AB857817](#), [AB857809](#), [AB857801](#), [AB857793](#), [KC524772](#),

KC524771, KC524770, JN711138, JN711137, JN711136, JQ886094, JQ886093, JQ886092, JQ231211, JQ231210, JN711135, JN711134, JN711133, JN711132, JN711131, JN711130, JN711129, JN711128, JN711127, JN712437, JN712436, FJ998173, FJ513678, FJ513677, FJ513676, FJ513674, FJ513672, FJ513671, FJ513670, FJ513669, FJ513668, FJ513667, FJ513666, FJ513665, FJ513664, FJ513663, FJ513662, FJ513661, FJ513660, FJ513659, FJ513658, FJ513656, FJ513655, FJ513653, FJ513652, FJ513651, FJ513650, FJ513649, FJ513648, FJ513647, FJ513646, FJ513644, FJ513643, FJ513642, FJ513641, FJ513640, FJ513639, FJ513638, FJ513636, FJ513634, FJ513633, FJ513631, FJ513630, FJ445509, FJ445508, FJ445507, FJ445506, FJ445505, FJ445504, FJ445503, FJ445501, FJ445500, FJ445499, FJ445498, FJ445497, FJ445496, FJ445495, FJ445494, FJ445493, FJ445492, FJ445491, FJ445490, FJ445489, FJ445488, FJ445482, FJ445481, FJ445480, FJ445479, FJ445478, FJ445477, FJ445476, FJ445475, FJ445474, FJ445473, FJ445472, FJ445471, FJ445470, FJ445469, FJ445468, FJ445467, FJ445466, FJ445465, FJ445464, FJ445462, FJ445461, FJ445460, FJ445459, FJ445458, FJ445457, FJ445456, FJ445455, FJ445454, FJ445453, FJ445452, FJ445451, FJ445450, FJ445449, FJ445448, FJ445447, FJ445446, FJ445444, FJ445442, FJ445441, FJ445440, FJ445439, FJ445438, FJ445437, FJ445436, FJ445435, FJ445434, FJ445429, FJ445425, FJ445424, EF613344, EF613343, EF613342, FJ445486, JN661156, JN661155, JN661154, JN661153, JN661152, JN661151, JN661150, JN661149, JN661148, JF264892, JF264891, JF264890, GQ143750, GQ143749, GQ143748, GQ143747, GQ143746, GQ143745, GQ143743, GQ143742, FN295489, FJ617290, FJ617289, FJ617288, FJ617287, FJ617286, FJ617285, FJ617284, FJ617283, FJ617282, EU288003, EU288002, EU288001, EU287999, EU287998, EU287997, EU287995, EU287994, EU170527, EU170526, EU170525, EU170524, EU170523, EU244646, FN295495, FN295494, FN295493, FN295492, FN295491, FN295490, FN295488, FN295486, DQ489787, GQ996377, GQ996376, GQ996375, GQ996374, GQ996373, GQ996372, GQ996371, GQ996370, JX839809, JX839808, JX839807, JX839806, JX839805, JX839804, JX839803, JX839802, JX839824, JX839823, JX839822, JX839821, JX839820, JX839819, JX839818, GQ143744, JX839784, JX839783, JX839782, JX839781, JX839780, JX839779, JX839778, JX839777, JQ013707, JQ013706, JQ013705, JQ013704, JQ013703, JQ013702, JQ013701, JQ013700, JQ013699, JQ013693, JQ013692, JQ013691, JQ013690, JQ013689, JQ013687, AB678695, AB678694, AB678693, AB678692, AB678691, AB678690, AB678689, AB678688, AB678687, AB678686, AB678685, AB678684, AB678683, AB678682, AB678681, AB678680, AB678679, AB678678, AB678677, GQ229489, GQ229488, GQ229487, GQ229486, GU969242, FJ445487, FJ445485, FJ445483, EU441883, EU441882, KC731586, KC731585, KC731584, KC731583, KC731582, KC731581, JQ943720, JQ943719,

JQ943718, JQ943717, JQ943716, JQ943715, JQ943714, JQ943713, JQ943712, JQ943711, JQ943710, JQ943709, JQ943708, JQ943707, JQ943706, JQ943705, JQ943704, JQ943703, JQ943702, JQ943701, JQ943700, JQ943699, JQ943698, JQ943696, JQ943695, JQ943694, JQ943693, JQ943692, GQ143741, JQ943691, JQ943690, JQ943689, JQ943688, HQ529778, JQ943687, HE578162, JQ943697, KC879578, KC879577, KC879576, KC879575, KC879574, KC879573, KC879572, KC879571, KC879570, KC879569, KC879568, KC879567, KC879566, KC879565, KC879564, KC879563, KC879562, KC879561, KC879560, KC879559, KC969208, KC969207, JX839829, JX839828, JX839827, JX839826, JX839825, JX839817, JX839816, JX839815, JX839814, JX839813, JX839812, JX839811, JX839810, JX839801, JX839800, JX839799, JX839798, JX839797, JX839796, JX839795, JX839794, JX839793, JX839792, JX839791, JX839790, JX839789, JX839788, JX839787, JX839786, JX839785, JX839776, JX839775, JX839774, JX839773, JX839772, JX839771, JX839770, JX839769, JX839768, JX839767, JX839766, GQ996379, GQ996378, EF555200, EF555199, EF555198, EF555197, EF555196, FJ432665, AF339485, FJ705371, FJ705370, FJ705369, CNR20236, CNR20235, HM045820, HM045819, HM045818, HM045817, HM045816, HM045807, HM045804, HM045798, HM045815, HM045785, HM045786, AY726732, HM045784, JF274082, EU037962, KF318729, JN558836, JN558835, JN558834, JX088705, DRDE-07, HM045822, HM045821, HM045814, HM045813, HM045812, HM045810, HM045809, HM045808, HM045806, HM045805, HM045803, HM045802, HM045801, HM045800, HM045799, HM045797, HM045796, SAH2123, HM045794, HM045793, HM045792, HM045791, HM045790, HM045789, HM045788, HM045787, AF490259, GU301781, GU301780, GU301779, GU908223, GQ905863, FJ445511, FJ445510, FJ445502, FJ445484, FJ445463, FJ445445, FJ445443, FJ445433, FJ445432, FJ445431, FJ445430, FJ445428, FJ445427, FJ445426, RGCB356, RGCB355, RGCB120, GQ428212, GQ428211, GQ428210, FJ000069, FJ000068, FJ000067, FJ000066, FJ000065, FJ000064, FJ000063, FJ000062, EF210157, KJ796852, KJ796851, KJ796850, KJ796849, KJ796848, KJ796847, KJ796846, KJ796845, KJ796844, KJ451624, KJ451623, KJ451622, KC862329, FN295483, FN295485, FN295487, FN295484, JQ067624, GU189061, HQ456255, HQ456254, HQ456253, HQ456252, HQ456251, GU199353, GU199352, GU199351, GU199350, FJ513679, FJ513675, FJ513673, FJ513657, FJ513654, FJ513645, FJ513637, FJ513635, FJ513632, FJ513629, FJ513628, FJ807899, FJ807898, FJ807897, FJ807896, FJ959103, EU244823, EU564335, EU564334, EF012359, EF027141, EF027140, EF027138, EF027137, EF027136, EF027135, EF027134, DQ443544, HE806461, FR717337, FR717336, EU703762, EU703761, EU703760, EU703759, AB455494, AB455493, HM159390, HM159389, HM159388, HM159387, HM159386, HM159385, HM159384.

1.4. Predicting importation and establishment risk

Under this model M_{ij} , the number of people moving from region i to j is given by:

$$M_{ij} = p \frac{\left(\frac{1-\lambda(n_i + s_{ij} + 1)}{n_i + s_{ij} + 1} - \frac{1-\lambda(n_i + n_j + s_{ij} + 1)}{n_i + n_j + s_{ij} + 1} \right)}{\frac{1-\lambda(n_i + 1)}{n_i + 1}},$$

where n_i and n_j are the populations at regions i and j respectively, s_{ij} is the total population living in all regions no further from region i than region j is, and p and λ are free parameters bounded on the unit interval which respectively govern the fraction of the population moving and the shape of the resulting distribution.

2.1 Supplementary Table S1.

Epidemiological and clinical characteristics of CHIKV confirmed cases.

Patient code (Acc. No.)	Age, Y (I)	Municipality of residence, Z (State, 3)	Simplified Country of origin	Diagnostic exam	Date of symptoms onset (SD)	Date of sample collection (SO)	SO-SC (d)	DENV serology	Location of travel	Date of travel (d)
P1	30-39	Povo Alegre (Rio Grande do Sul) (D)	Imported	ELISA	08-Aug	24-Jun	77	-	Brazil (Paraná and Rio Grande do Sul)	early April
P2	30-39	Povo Alegre (Rio Grande do Sul) (D)	Imported	ELISA	04-May	30-May	26	Positive	Haiti	early May
P3	20-29	Nova Friburgo (Rio de Janeiro)	Imported	ELISA	12-May	02-Jun	19	-	Haiti	-
P4	30-39	Monte Azul Paulista (São Paulo)	Imported	ELISA	14-May	03-Jun	19	-	Haiti	-
P5	40-49	Aratiba do Carmo (Rio de Janeiro)	Imported	ELISA	15-May	18-Jun	18	Negative	Haiti	-
P6	40-49	Aratiba do Carmo (Rio de Janeiro)	Imported	ELISA	31-May	05-Jun	18	Negative	Haiti	-
P7	30-39	Marzagão (Paraná) (C)	Imported	ELISA	01-Jun	08-Jun	17	Negative	Haiti	-
P8	30-39	Itui (São Paulo)	Imported	ELISA	01-Jun	08-Jun	17	Negative	Haiti	-
P9	30-39	Marzagão (Paraná) (C)	Imported	ELISA	01-Jun	08-Jun	17	Negative	Haiti	-
P10	20-29	São Vicente (São Paulo)	Imported	ELISA	03-Jun	05-Jun	2	Negative	Haiti	-
P11	40-49	Taubaté (São Paulo)	Imported	ELISA	03-Jun	05-Jun	2	Negative	Haiti	-
P12	20-29	Horizontina (São Paulo)	Imported	ELISA	03-Jun	05-Jun	2	Negative	Haiti	-
P13	20-29	São Vicente (São Paulo)	Imported	ELISA	04-Jun	05-Jun	2	Negative	Haiti	-
P14	20-29	Campana (São Paulo)	Imported	ELISA	04-Jun	05-Jun	2	Negative	Haiti	-
P15	20-29	São Vicente (São Paulo)	Imported	ELISA	04-Jun	05-Jun	2	Negative	Haiti	-
P16	20-29	Campana (São Paulo)	Imported	ELISA	04-Jun	05-Jun	2	Negative	Haiti	-
P17	20-29	Campana (São Paulo)	Imported	ELISA	07-Jun	07-Jun	0	Negative	Haiti	-
P18	20-29	Itaó Paulo (São Paulo)	Imported	ELISA	22-Jun	03-Jul	20	-	Dominican Rep.	-
P19	20-29	Itaó Paulo (São Paulo)	Imported	ELISA	22-Jun	03-Jul	20	-	Dominican Rep.	-
P20	20-29	Itaó Paulo (São Paulo)	Imported	ELISA	25-Jun	04-Jul	22	-	Dominican Rep.	-
P21	50-59	Brasília (Distrito Federal)	Imported	ELISA	27-Jun	04-Jul	27	-	Dominican Rep.	-
P22	40-49	Brasília (Distrito Federal)	Imported	ELISA	01-Jul	04-Jul	3	Negative	Dominican Rep.	-
P23	40-49	Brasília (Distrito Federal)	Imported	ELISA	02-Jul	04-Jul	3	Negative	Dominican Rep.	-
P24	40-49	Brasília (Distrito Federal)	Imported	ELISA	02-Jul	04-Jul	3	Negative	Dominican Rep.	-
P25 (KP164571)	30-39	Riachão (Pernambuco)	Imported	ELISA	02-Jul	03-Jul	1	-	Dominican Rep.	early June
P26	30-39	Fortaleza (Ceará)	Imported	ELISA	08-Jul	18-Jul	11	-	Dominican Rep.	-
P27	50-59	Fortaleza (Ceará)	Imported	ELISA	08-Jul	18-Jul	11	-	Dominican Rep.	-
P28	50-59	Fortaleza (Ceará)	Imported	ELISA	13-Jul	21-Jul	8	-	Dominican Rep.	-
P29	20-29	Fortaleza (Ceará)	Imported	ELISA	13-Jul	21-Jul	8	-	Dominican Rep.	-
P30	60-69	Aracajuana (São Paulo)	Imported	ELISA	23-Jul	31-Jul	4	-	Dominican Rep. and Haiti	-
P31	10-19	Santana (Amazon)	Imported	ELISA	23-Jul	31-Jul	4	-	Did not travel outside Brazil	-
P32	10-19	Santana (Amazon)	Imported	ELISA	31-Jul	31-Jul	0	-	Caribbean region	-
P33	50-59	Colônia (Cuiabá)	Imported	ELISA	09-Aug	21-Aug	3	-	Guadeloupe	-
P34 (KP164572)	40-49	Belém (Pará)	Imported	ELISA	18-Aug	26-Aug	3	-	Guadeloupe	-
P35 (KP164580)	60-69	Feira de Santana (Bahia)	Autochthonous	ELISA	25-Aug	26-Aug	1	-	None	-
P36 (KP164581)	30-39	Feira de Santana (Bahia)	Autochthonous	ELISA	25-Aug	26-Aug	1	-	None	-
P37 (KP164587)	30-39	Feira de Santana (Bahia)	Autochthonous	ELISA	25-Aug	26-Aug	1	-	None	-
P38 (KP164570)	20-29	Feira de Santana (Bahia)	Autochthonous	ELISA	25-Aug	26-Aug	1	-	None	-
P39 (KP164589)	20-29	Feira de Santana (Bahia)	Autochthonous	ELISA	25-Aug	26-Aug	1	-	None	-
P40	20-29	Feira de Santana (Bahia)	Autochthonous	ELISA	27-Aug	28-Aug	1	-	None	-
P41	20-29	Feira de Santana (Bahia)	Autochthonous	ELISA	27-Aug	28-Aug	1	-	None	-
P42	60-69	Feira de Santana (Bahia)	Autochthonous	ELISA	27-Aug	28-Aug	1	-	None	-
P43	50-59	Feira de Santana (Bahia)	Autochthonous	ELISA	27-Aug	28-Aug	1	-	None	-
P44	50-59	Feira de Santana (Bahia)	Autochthonous	ELISA	27-Aug	28-Aug	1	-	None	-
P45	20-29	Feira de Santana (Bahia)	Autochthonous	ELISA	28-Aug	29-Aug	1	-	None	-
P46	30-39	Feira de Santana (Bahia)	Autochthonous	ELISA	28-Aug	29-Aug	1	-	None	-
P47	0-9	Capitão (Amazon)	Autochthonous	ELISA	28-Aug	29-Aug	1	-	None	-
P48	70-79	Feira de Santana (Bahia)	Autochthonous	ELISA	30-Aug	04-Sep	5	-	None	-
P49	30-39	Feira de Santana (Bahia)	Autochthonous	ELISA	30-Aug	04-Sep	5	-	None	-
P50	30-39	Feira de Santana (Bahia)	Autochthonous	ELISA	30-Aug	04-Sep	5	-	None	-
P51	0-9	Boa Vista (Roraima)	Imported	ELISA	02-Sep	08-Sep	7	-	Venezuela	early September
P52	30-39	Olápoque (Amapá)	Autochthonous	ELISA	03-Sep	15-Sep	11	-	None (São Isabel, Piqui)	-
P53	30-39	Olápoque (Amapá)	Autochthonous	ELISA	03-Sep	15-Sep	11	-	None (São Isabel, Piqui)	-
P54	0-9	Boa Vista (Roraima)	Imported	ELISA	06-Sep	17-Sep	12	-	Venezuela	late August
P55	20-29	Olápoque (Amapá)	Autochthonous	ELISA	06-Sep	17-Sep	12	-	Venezuela	late August
P56	20-29	Olápoque (Amapá)	Autochthonous	ELISA	06-Sep	17-Sep	12	-	Venezuela	late August
P57	30-39	Olápoque (Amapá)	Autochthonous	ELISA	08-Sep	15-Sep	6	-	None	-
P58	40-49	Olápoque (Amapá)	Autochthonous	ELISA	08-Sep	15-Sep	6	-	None	-
P59	40-49	Olápoque (Amapá)	Autochthonous	ELISA	08-Sep	15-Sep	6	-	None	-
P60	40-49	Olápoque (Amapá)	Autochthonous	ELISA	08-Sep	15-Sep	6	-	None	-
P61	40-49	Olápoque (Amapá)	Autochthonous	ELISA	11-Sep	12-Sep	1	-	Brazil (Pernambuco, Ceará, Maranhão, São Paulo)	late August
P62	20-29	Olápoque (Amapá)	Autochthonous	ELISA	11-Sep	12-Sep	1	-	Brazil (Pernambuco, Ceará, Maranhão, São Paulo)	late August
P63	20-29	Olápoque (Amapá)	Autochthonous	ELISA	11-Sep	12-Sep	1	-	Brazil (Pernambuco, Ceará, Maranhão, São Paulo)	late August
P64	0-9	Marzagão (Paraná)	Imported	ELISA	15-Sep	22-Sep	7	-	Brazil (Ceará)	early September
P65	20-29	Planalto (Rio de Janeiro)	Imported	ELISA	15-Sep	22-Sep	7	-	Brazil (Ceará)	early September
P66	30-39	Planalto (Rio de Janeiro)	Imported	ELISA	15-Sep	22-Sep	7	-	Brazil (Ceará)	early September
P67	20-29	Campana (São Paulo)	Imported	ELISA	15-Sep	22-Sep	7	-	Brazil (Ceará)	early September
P68	0-9	Boa Vista (Roraima)	Imported	ELISA	20-Sep	06-Oct	16	-	Brazil (Guiana)	-

Notes:
 1. Ages have been grouped into decades to preserve anonymity.
 2. A, B, C and D reside in the same street or address and are presumed clusters of CHIKV infection.
 3. Federal state of notification.
 4. Dates of travel have been summarized to the nearest fortnight to ensure anonymity.

2.2 Supplementary Table S2.

The 35 municipalities at higher risk of CHIKV importation from both Oiapoque and Feira de Santana municipalities.

Municipality	Federal State	Country Region
Brasilia	Distrito Federal	Central-West
Goiania	Goias	Central-West
Manaus	Amazonas	North
Belem	Para	North
Maceio	Alagoas	Northeast
Feira de Santana	Bahia	Northeast
Salvador	Bahia	Northeast
Caucaia	Ceara	Northeast
Crato	Ceara	Northeast
Fortaleza	Ceara	Northeast
Sao Luis	Maranhao	Northeast
Bayeux	Paraiba	Northeast
Campina Grande	Paraiba	Northeast
Joao Pessoa	Paraiba	Northeast
Santa Rita	Paraiba	Northeast
Abreu E Lima	Pernambuco	Northeast
Camaragibe	Pernambuco	Northeast
Jaboatao Dos Guararapes	Pernambuco	Northeast
Paulista	Pernambuco	Northeast
Petrolina	Pernambuco	Northeast
Recife	Pernambuco	Northeast
Teresina	Piaui	Northeast
Natal	Rio Grande Do Norte	Northeast
Aracaju	Sergipe	Northeast
Belo Horizonte	Minas Gerais	Southeast
Contagem	Minas Gerais	Southeast
Duque de Caxias	Rio de Janeiro	Southeast
Nilopolis	Rio de Janeiro	Southeast
Rio de Janeiro	Rio de Janeiro	Southeast
Carapicuiaba	Sao Paulo	Southeast
Diadema	Sao Paulo	Southeast
Guarulhos	Sao Paulo	Southeast
Maua	Sao Paulo	Southeast
Sao Caetano do Sul	Sao Paulo	Southeast
Sao Paulo	Sao Paulo	Southeast

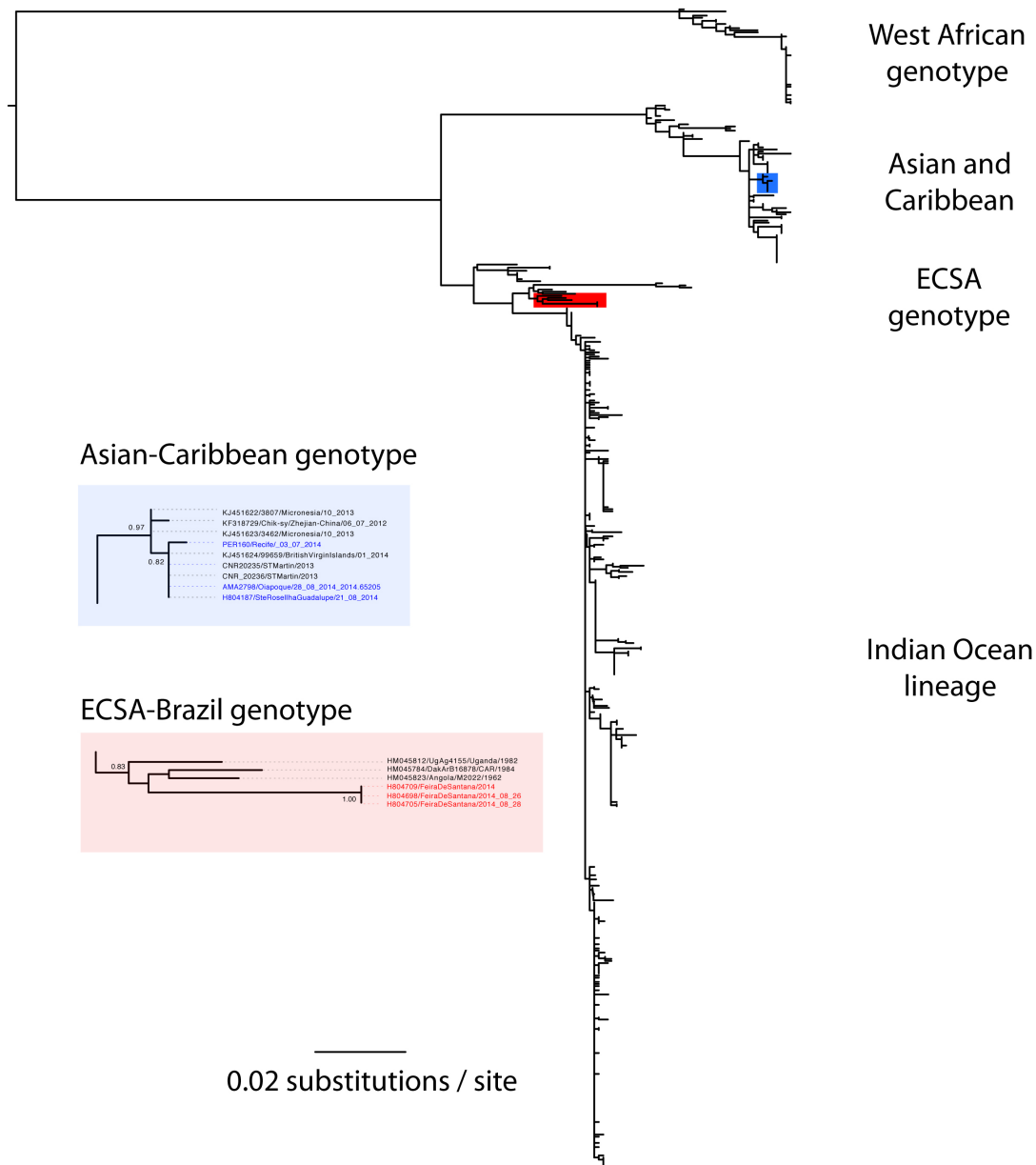
3.1. Supplementary Figure S1.

Maximum-likelihood phylogenetic reconstruction of 76 CHIKV genomes, including the Brazilian strains. The strain names in red correspond to Brazilian isolates belonging to the ECSA genotype and the strain names in blue indicate Brazilian isolates belonging to the Asian genotype. Numbers from 0 to 1 indicate phylogenetic branch support obtained using an approximate likelihood ratio test (SH-like) in PhyML [6]. Branch lengths in units of substitutions per site.



3.2. Supplementary Figure S2.

Maximum-likelihood phylogenetic reconstruction of 554 CHIKV envelope sequences, including the Brazilian strains. The strain names in red correspond to Brazilian isolates belonging to the ECSA genotype and the strain names in blue indicate Brazilian isolates belonging to the Asian genotype. Numbers from 0 to 1 indicate phylogenetic branch support obtained using an approximate likelihood ratio test (SH-like) in PhyML [6].



4. References

1. Chevreux B, Pfisterer, T., Drescher, B., Driesel, A. J., Muller, W. E. G., Wetter, T., Suhai, S. (2004) Using the miraEST Assembler for Reliable and Automated mRNA Transcript Assembly and SNP Detection in Sequenced ESTs. *Genome Research*: 1147-1159.
2. Kearse M, Moir R, Wilson A, Stones-Havas S, Cheung M, et al. (2012) Geneious Basic: an integrated and extendable desktop software platform for the organization and analysis of sequence data. *Bioinformatics* 28: 1647-1649.
3. Altschul SF, Gish W, Miller W, Myers EW, Lipman DJ (1990) Basic local alignment search tool. *J Mol Biol* 215: 403-410.
4. Guindon S, Dufayard JF, Lefort V, Anisimova M, Hordijk W, et al. (2010) New algorithms and methods to estimate maximum-likelihood phylogenies: assessing the performance of PhyML 3.0. *Syst Biol* 59: 307-321.
5. Drummond AJ, Suchard MA, Xie D, Rambaut A (2012) Bayesian phylogenetics with BEAUti and the BEAST 1.7. *Mol Biol Evol* 29: 1969-1973.
6. Gouy M, Guindon S, Gascuel O (2010) SeaView version 4: A multiplatform graphical user interface for sequence alignment and phylogenetic tree building. *Mol Biol Evol* 27: 221-224.