

Does the viral circulation in slums have a global impact? The lesson learned from early SARS-CoV-2 circulation in *Complexo de Favelas da Maré*, Rio de Janeiro, Brazil

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SUPPLEMENTARY FIGURE

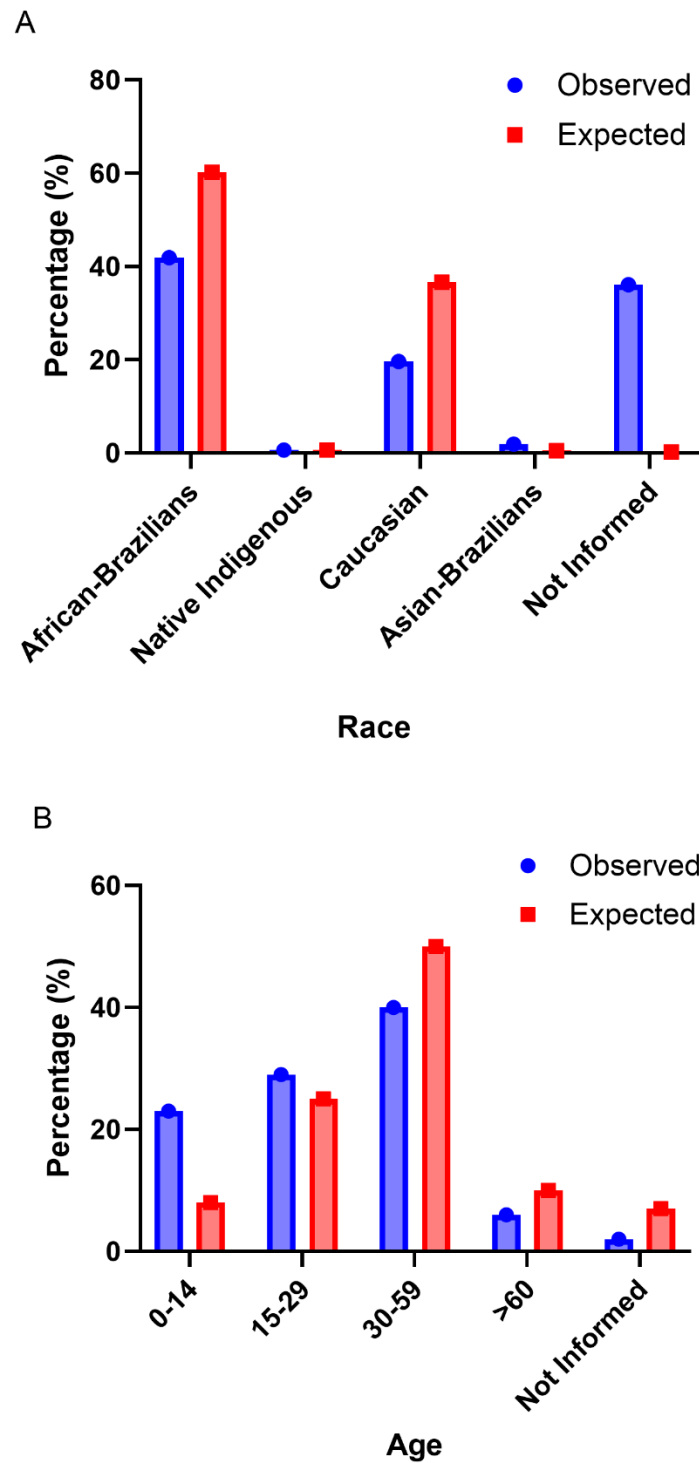


Figure 1. Descriptive comparison of the collected patients (observed) and censored individuals (expected) at *Maré*. Panel A represents the races and panel B the ages.

SUPPLEMENTARY TABLES

Supplementary Table 1 – SARS-CoV-2 genomes from Gamma VoC similar to those of strains circulating in *Complexo de Favela da Maré*

<i>Description</i>	<i>Total Score</i>	<i>Query Cover</i>	<i>E value</i>	<i>Acc. Len</i>	<i>Accession</i>
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/BRA/HCOR_A823000 661/2021</i>	54215	54215	100%	99.94	29903
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/BRA/HCOR_666488/ 2021</i>	54143	54143	100%	99.90	29903
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/BRA/RJ-NVBS648GENOV82755957296 9/2021</i>	54143	54143	100%	99.90	29831
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/BRA/1061/2021</i>	54143	54143	100%	99.90	29903
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/VSP2639/2021</i>	54141	54141	100%	99.89	29907
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-</i>	54137	54137	100%	99.89	29866

Supplementary Material

2/human/BRA/CD1739-
P4/2020

Severe acute respiratory syndrome coronavirus 2 isolate Switzerland/GE- 33242357/2021 genome assembly	54135	54135	100%	99.89	29865
Severe acute respiratory syndrome coronavirus 2 isolate Switzerland/GE- 33173470/2021 genome assembly	54135	54135	100%	99.89	29850
Severe acute respiratory syndrome coronavirus 2 isolate Switzerland/GE- 33173437/2021 genome assembly	54135	54135	100%	99.89	29863
Severe acute respiratory syndrome coronavirus 2 isolate Switzerland/GE- 33154798/2021 genome assembly	54135	54135	100%	99.89	29865
Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV- 2/human/USA/FAA4B8136EQ/ 2021	54131	54131	100%	99.89	29821
Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV- 2/human/USA/FA6C70F0FCE/ 2021	54131	54131	100%	99.89	29806

Supplementary Table 2 – SARS-CoV-2 genomes from Delta VoC strains similar to those of strains circulating in *Complexo de Favela da Maré*

<i>Description</i>	<i>Total Score</i>	<i>Query Cover</i>	<i>E value</i>	<i>Acc. Len</i>	<i>Accession</i>
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/IND/SRHU_58/2021</i>	<i>54200</i>	<i>54200</i>	<i>100%</i>	<i>99.93</i>	<i>29903</i>
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/CA-SEARCH-49941/2021</i>	<i>54194</i>	<i>54194</i>	<i>100%</i>	<i>99.92</i>	<i>29845</i>

Supplementary Material

<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/France/10029DT/2020</i>	54194	54194	100%	99.92	29903
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/NY-PRL-210217_00L24/2021</i>	54189	54189	100%	99.92	29752
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/ZAF/NHLS-UCT-GP-5276/2020</i>	54189	54189	100%	99.92	29823
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/ZAF/NHLS-UCT-GP-5326/2020</i>	54189	54189	100%	99.92	29820
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/ZAF/NHLS-UCT-GS-0087/2020</i>	54189	54189	100%	99.92	29820
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/CHN/SH-P94-B-3-Europe-UK/2020</i>	54189	54189	100%	99.92	29689
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/CHN/SH-P125-3-Asia-UAE/2020</i>	54189	54189	100%	99.92	29689
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/SWE/14_SE300_DE20-005545/2020</i>	54189	54189	100%	99.92	29865
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/IND/SRHU_54/2021</i>	54189	54189	100%	99.92	29903

Supplementary Material

<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/CMI-IGM-004/2021</i>	54187	54187	100%	99.91	29803
<i>Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/FL-BPHL-3645/2020</i>	54185	54185	100%	99.91	29714
