

CORRECTION

Correction: Examining the Species-Specificity of Rhesus Macaque Cytomegalovirus (RhCMV) in Cynomolgus Macaques

The *PLOS ONE* Staff

There are multiple errors in [Fig 5](#) and its caption, “Sequence alignment comparing RhCMV-eGFP to RhCMV 68–1 BAC.” Please see the complete, corrected [Fig 5](#) here. The publisher apologizes for the error.



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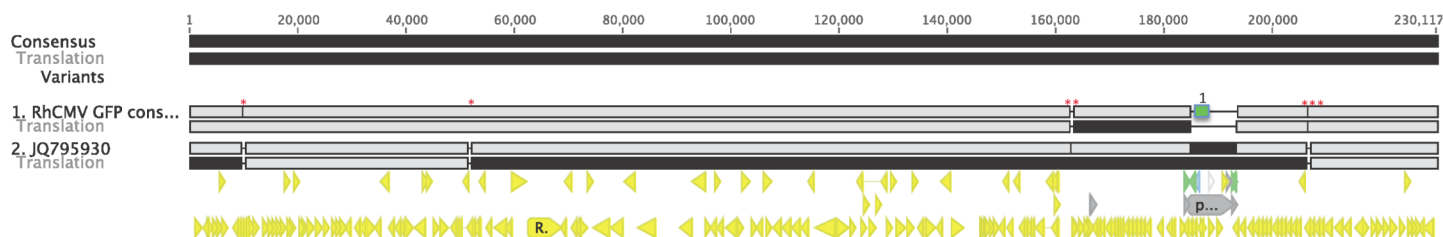


Fig 5. Sequence alignment comparing RhCMV-eGFP to RhCMV 68–1 BAC. Sequence of RhCMV-eGFP obtained from NGS analysis was aligned with that of RhCMV 68–1 BAC (JQ795930) using Geneious Pro 6.1.4 software. Whole genome comparisons yielded 96.2% similarity and the major difference can be attributed to the BAC cassette present in RhCMV68-1 BAC. “1” refers to eGFP reporter * sequencing error; ** 22 bp deletion; *** 56 bp insertion.

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Reference

1. Marsh AK, Ambagala AP, Perciani CT, Russell JNH, Chan JK, Janes M, et al. (2015) Examining the Species-Specificity of Rhesus Macaque Cytomegalovirus (RhCMV) in Cynomolgus Macaques. PLoS ONE 10(3): e0121339. doi:[10.1371/journal.pone.0121339](https://doi.org/10.1371/journal.pone.0121339) PMID: [25822981](https://pubmed.ncbi.nlm.nih.gov/25822981/)