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Author Correction: Infectious DNAs derived from insect-specific flavivirus genomes enable identification of pre- and post-entry host restrictions in vertebrate cells

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In this Article, the legend of Figure 4 is incorrect:

“Viral replication analyses in WT and IFNAR^{-/-} MEF cells. (a) Cells infected at an MOI of 1 or transfected CPEC constructs of either WNV_{KUN} or PaRV with a CMV promoter. (b) Cells transfected with genomic RNA of either PaRV, WNV_{KUN}, PCV, PCV/WNV_{KUN}-prME or WNV_{KUN}/PCV-prME. Monolayers were fixed 72 hrs post-infection. IFA analysis was performed by probing with anti-PaRV (7D11) and anti-WNV E (3.91D) mouse monoclonal antibodies. The nucleus of each cell was stained with Hoechst 33342. Images were taken at ×40 magnification”.

should read:

“Viral replication analyses in WT and IFNAR^{-/-} MEF cells. (a) Cells infected at an MOI of 1 with either WT WNV_{KUN} or WT PaRV. (b) Cells transfected with CPEC constructs of either WNV_{KUN} or PaRV with a CMV promoter. Monolayers were fixed 72 hrs post-infection. IFA analysis was performed by probing with anti-PaRV (7D11) or anti-WNV E (3.91D) mouse monoclonal antibodies. The nucleus of each cell was stained with Hoechst 33342. Images were taken at ×40 magnification”.



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