

EDITORIAL EXPRESSION OF CONCERN

Mutational analysis of the HIV-1 auxiliary protein Vif identifies independent domains important for the physical and functional interaction with HIV-1 reverse transcriptase

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The editors express a note of concern regarding the above article.

The editors wish to alert the readers that questions have been raised about the validity of some of the figures presented in this article. The editors note that Figures 2A, B, D, E, G, 3C, 4B and C have been inappropriately manipulated in order to enhance the presentation of the western blots. Although this indicates incorrect scientific practice, we believe, having scrutinized the original data, that these manipulations do not affect the conclusions of this study, which appear to be still valid. Nonetheless, we wish to alert our readership to these concerns.

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