

CORRECTION

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Cite this: *Anal. Methods*, 2017, 9, 5557

# Correction: Normalization strategy for the LC-MS bioanalysis of protein kinetics assays *via* internal proteolytic analyte utilized as control standard: application in studies of HIV-1 protease cleavage of HIV-1 Gag polyprotein in HIV maturation inhibition research

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Correction for 'Normalization strategy for the LC-MS bioanalysis of protein kinetics assays *via* internal proteolytic analyte utilized as control standard: application in studies of HIV-1 protease cleavage of HIV-1 Gag polyprotein in HIV maturation inhibition research' by Joseph L. Cantone *et al.*, *Anal. Methods*, 2017, DOI: 10.1039/c7ay01666b.

DOI: 10.1039/c7ay90125a

[www.rsc.org/methods](http://www.rsc.org/methods)

Ref. 1, 2, 4 and 6 are incorrect, the correct references are listed below:

1. UNAIDS Fact Sheet, <http://www.unaids.org/en/resources/fact-sheet>.
  2. AVERT, <http://www.avert.org/global-hiv-and-aids-statistics>.
  4. AIDSinfo, <https://aidsinfo.nih.gov/understanding-hiv-aids/fact-sheets/21/58/fda-approved-hiv-medicines/>.
  6. WHO HIV Drug Resistance Report 2012, [http://apps.who.int/iris/bitstream/10665/75183/1/9789241503938\\_eng.pdf?ua=1](http://apps.who.int/iris/bitstream/10665/75183/1/9789241503938_eng.pdf?ua=1).
- The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

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