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Corrections

Solution NMR Evidence That the HIV-1 Protease Catalytic Aspartyl Groups Have Different Ionization States in the Complex Formed with the Asymmetric Drug KNI-272, by Yun-Xing Wang, Darón I. Freedberg, Toshimasa Yamazaki, Paul T. Wingfield, Stephen J. Stahl, Joshua D. Kaufman, Yoshiaki Kiso, and Dennis A. Torchia*, Volume 35, Number 31, August 6, 1996, Pages 9945–9950.

Page 9946. In the caption to Figure 1, H₁₀ should read H₁. H₁ is the methylthioalanine H_α of KNI-272.

Page 9947. In Figure 3, H₁₀ should read H₁ in the spectrum, and H₂₃ should read H₂₂ in the caption.

Page 9948. In Figure 4, H₁₆, H₂₀, and H₂₁ should read H₁, H₁₈, and H₁₉, respectively, *T*₁₂/*T*₁₁₂ should be in italic letters, methyl should read thiomethyl, the D29–R87 link should read D30–R87, and the D129–R187 link should be omitted.

Page 9950. Tables S2 and S3 (in Supporting Information) have also been corrected and are available as new Supporting Information.

The Asp assignments in the paper are correct, as are the results and conclusions regarding the Asp p*K*_a values and ionization states.

SUPPORTING INFORMATION AVAILABLE

Proton chemical shifts of inhibitor KNI-272 (Table S2) and chemical shift assignments of the ¹H, ¹³C, and ¹⁵N signals of the HIV-1 protease/KNI-272 complex (Table S3) (7 pages). Ordering information is given on any current masthead page.

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