

Addition/Correction

**Microarray Platform for Profiling Enzyme Activities in Complex
Proteomes [*J. Am. Chem. Soc.* 2004, 126, 15640–15641].**

Stephan A. Sieber, Tony S. Mondala, Steven R. Head, and Benjamin F. Cravatt

J. Am. Chem. Soc., **2005**, 127 (11), 4114-4114 • DOI: 10.1021/ja0506915 • Publication Date (Web): 26 February 2005

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Stepwise Hydration, and Multi-Body Deprotonation with Steep Negative Temperature Dependence, in the Benzene⁺–Water System [*J. Am. Chem. Soc.* **2004**, *126*, 12766–12767]. Yehia Ibrahim, Edreese Alsharaeh, Keith Dias, Michael Meot-Ner (Mautner),* and M. Samy El-Shall*

In Supporting Information for this contribution, an error occurred in plotting Figure 2. The corrected Figure 2 is below. The actual Figure 2 in the Supporting Information at <http://pubs.acs.org> has also been corrected.

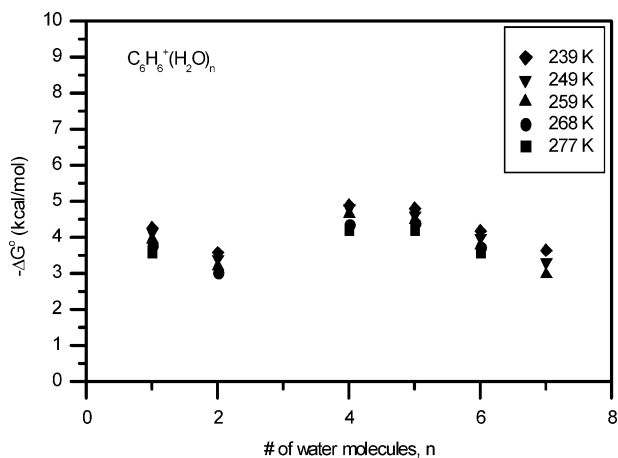


Figure 2.

JA059903O

10.1021/ja059903o

Published on Web 02/23/2005

Microarray Platform for Profiling Enzyme Activities in Complex Proteomes [*J. Am. Chem. Soc.* **2004**, *126*, 15640–15641]. Stephan A. Sieber, Tony S. Mondala, Steven R. Head, and Benjamin F. Cravatt*

Page 15641. In the text and the legend to Figure 3, the units for the sensitivity limits of ABPP methods were mistakenly inverted. For the text, the correct units for gel- and microarray-based ABPP are 125–250 and 2–8 ng/mL, respectively. Conversely, in the legend for Figure 3, the correct units for gel- and microarray-based ABPP are 0.12–0.25 and 0.002–0.008 μg/mL, respectively.

JA0506915

10.1021/ja0506915

Published on Web 02/26/2005