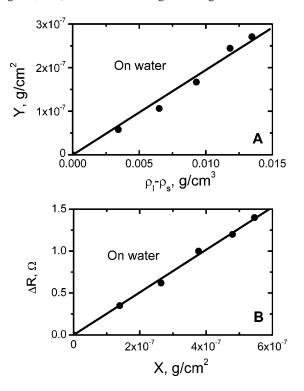
## Additions and Corrections

Adsorption of Bituminous Components at Oil/Water Interfaces Investigated by Quartz Crystal Microbalance: Implications to the Stability of Water-in-Oil Emulsions

Lamia Goual, Géza Horváth-Szabó,\* Jacob H. Masliyah, and Zhenghe Xu. *Langmuir* **2005**, *21*, 8278–8289.

The axes units in Figure 3 on page 8281 are erroneous, as kindly brought up by Dr. Kunal Karan (Queen's University, Kingston, ON). The corrected Figure 3 is given below. In this



Figure, we are not providing the data on gold because, contrary to our original assumption, the viscosity difference between H<sub>2</sub>O

and  $D_2O$  is not negligible. The nonlinear viscosity change of  $H_2O$  and  $D_2O$  mixtures should be used to obtain the precise h and  $C_r$  parameters of gold. Because the corrections applied at high bitumen concentrations are based on these parameters, the isotherm corresponding to the oil/gold interface at 10 wt % bitumen concentration in Figure 4 on page 8283 should not be considered.

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Amphiphilic 4-Helix Bundles Designed for Biomolecular Materials Applications

Shixin Ye, Joseph W. Strzalka, Bohdana M. Discher, Dror Noy, Songyan Zheng, P. Leslie Dutton, and J. Kent Blasie\* *Langmuir* **2004**, *20*, 5897–5904.

- (1) Materials and Methods Section, Synthesis. The peptide (NH<sub>2</sub>-¹EIWKLHE.EF¹¹0LKKFE.ELLKL²¹0HE.ERL-KKLL.L³¹0LALLQL.LLAL⁴¹0LQL.GGC-CONH₂) rather than (NH<sub>2</sub>-¹EIWKLHE.EF¹¹0LKKFE.ELLKL²¹0HE.ERLKKLL.L³¹0-QALLQL.LQAL⁴¹0LQL.GGC-CONH₂) was synthesized on Applied Biosystems' Pioneer continuous flow solid phase synthesizer using the standard Fmoc/tBu protection strategy on a Fmoc-PEG-PAL-PS resin (Applied Biosystems) on the 0.1 mmol scale.
- (2) Results Section, Design of the Amphiphilic Peptide AP0. The two C-terminal heptads (residues 29-42) are based on the design of synthetic proton channel LS $_2$  (H $_2$ N-(Leu-Ser-Leu-Leu-Leu-Ser-Leu) $_3$ -CONH $_2$ ). We replace only two serines in LS $_2$  with glutamine to incorporate hydrogen bonding in the interior of the hydrophobic domain and thereby induce proper assembly of the protein.

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