In the article, "Peptide Shifter: Enhancing Separation Reproducibility Using Correlated Expression Profiles" by Dmitri Sitnikov, Joanna M. Hunter, Clive Hayward, Kevin Eng, Isabelle Migneault, Sylvain Tessier, Gregory J. Opiteck and Paul Kearney, which was published in the September issue, Vol. 18, No. 9, pages 1638-1645, a printer error caused Figure 6 to print in black and white, when it should have been in color. The corrected figure is shown below, with the corresponding figure legend.

Figure 6. Expression profile correlation heatmap for all pairs of peptides from Fraction 2. Each point in the heatmap represents the Spearman correlation between the expression profiles of two peptides. Green is high correlation and red is low correlation. Peptides are highly correlated to themselves—thus, the diagonal green line. Hierarchical clustering has been applied to group highly correlated peptides. Framed in blue are the peptides corresponding to the two most frequently occurring expression profiles in Fraction 2, which appear in Figure 5. The upper left blue box corresponds to peptides shifted into Fraction 2, whereas the lower right blue box corresponds to peptides shifted out of Fraction 2. As expected, these two sets of peptides are highly negatively correlated (two yellow boxes).