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ADDITION/CORRECTION

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A High-Throughput *O*-Glycopeptide Discovery Platform for Seromic Profiling [*Journal of Proteome Research* **2010**, *9* 5250–5261 DOI: 10.0121/pr1005229]. Ola Blixt, Emiliano Cló, Aaron S. Nudelman, Kasper Kildegaard Sørensen, Thomas Clausen, Hans H. Wandall, Philip O. Livingston, Henrik Clausen, and Knud J. Jensen

Page 5258. Figure 5 was incorrect in the original publication. The correct Figure 5 is listed below.

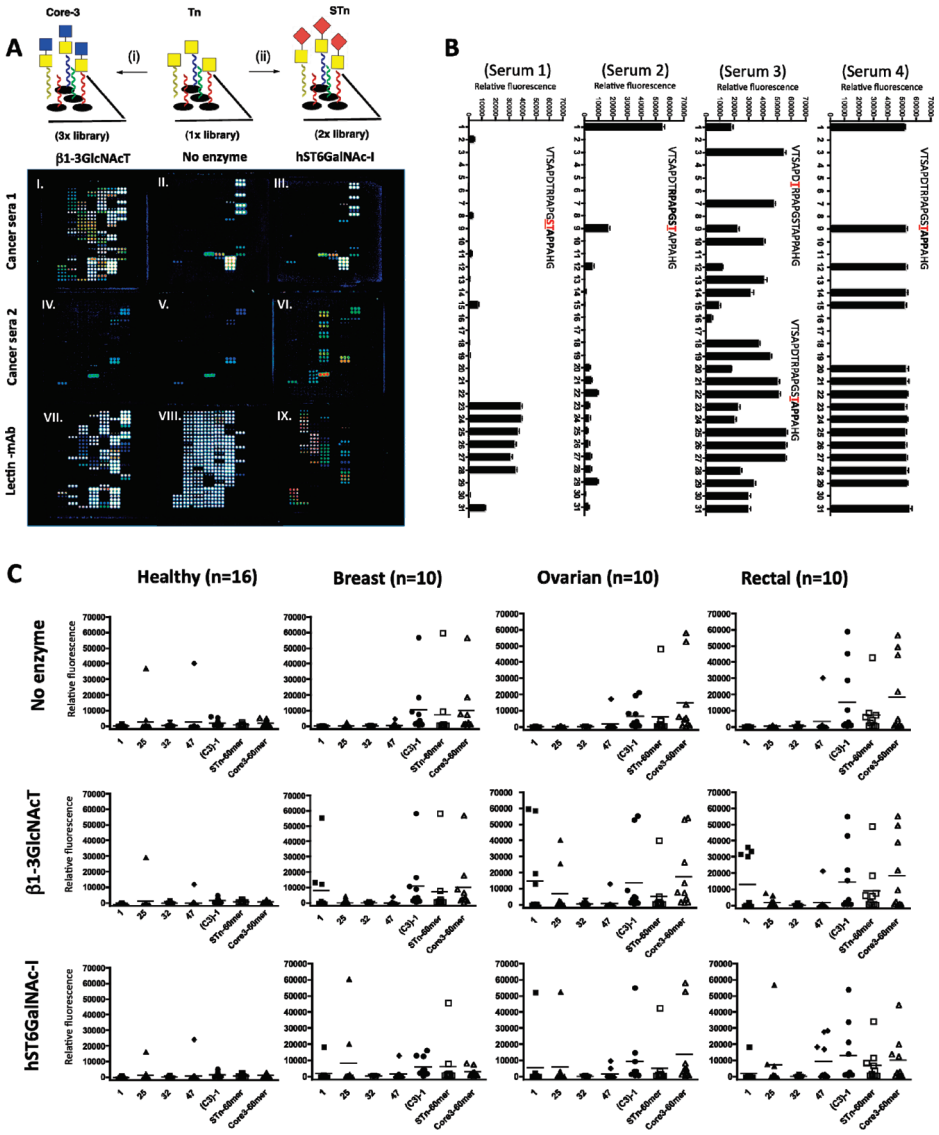

Figure 5. On-slide enzymatic glycosylation and detection of serum autoantibodies. (A) Image display after on-slide enzymatic glycosylation using core3 3GlcNAc-T6 (UDP-GlcNAc) (i) and ST6GalNAc-I (CMP-NeuAc) (ii) followed by subsequent staining with sera (1:20) and Cy3-labeled anti-human-IgG. (B) Glycoform-specific MUC1 epitope autoantibody reactivity in four different cancer sera. Predominant MUC1 epitope for each sera is depicted as a bold sequence in each serum panel. (C) Detection of cancer sera and healthy sera on a panel of selected MUC1 20mers and MUC1 60mer controls. Spot-to-spot variations for three replicates of each compound are represented by the error bars.

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