Corrigendum

In the article by J. Kaur and A. K. Bachhawat (GENETICS **176**: 877–890) entitled "Yct1p, a Novel, High-Affinity, Cysteine-Specific Transporter from the Yeast *Saccharomyces cerevisiae*," on page 883, in the control methionine panel of Figure 4A, two duplications have been identified. This oversight, which has no bearing on the results or conclusions of the article, was brought to the attention of the authors through public discussions on PubPeer.

The authors could access neither the original high-resolution images nor the replicates for the methionine panel in Figure 4A. However, since all the strains were still available, the authors have redone the growth analysis using the same serial dilution spotting experiment and prepared a fresh image for the Figure 4A methionine panel (presented below).

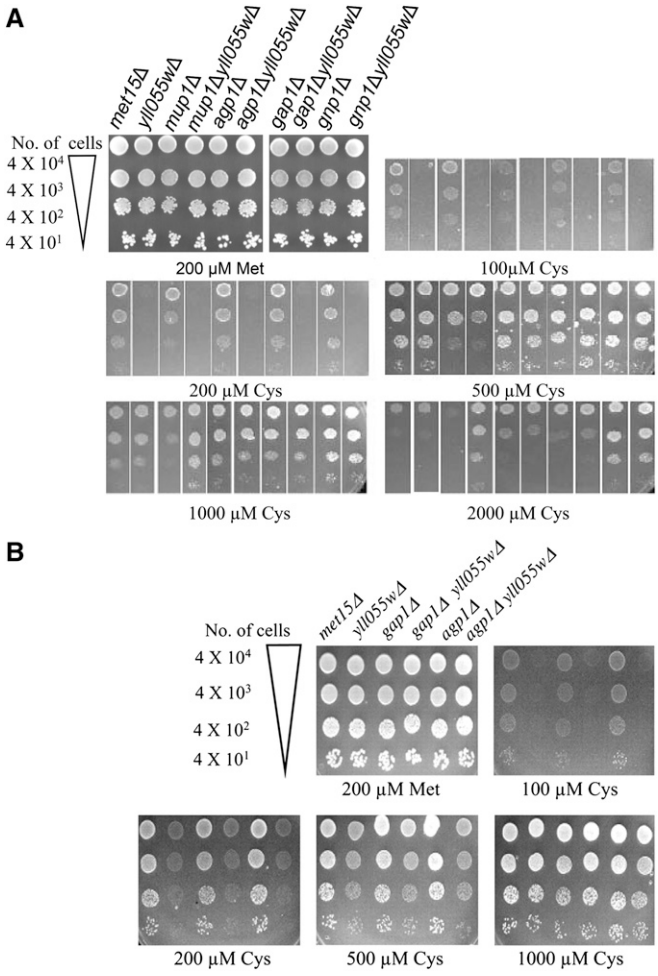

Figure 4 Effect of *YLL055w* disruption on utilization of cysteine as the sole sulfur source in different strain backgrounds deleted in the major cysteine permeases in the presence of different nitrogen sources. (A) Strains *met15***Δ**, *met15***Δ** *yll055w***Δ**, *met15***Δ** *mup1***Δ**, *met15***Δ** *mup1***Δ** *yll055w***Δ**, *met15***Δ** *agp1***Δ**, *met15***Δ** *agp1***Δ** *yll055w***Δ**, *met15***Δ** *gap1***Δ**, *met15***Δ** *gap1***Δ** *yll055w***Δ**, *met15***Δ** *gnp1***Δ**, and *met15***Δ** *gnp1***Δ** *yll055w***Δ** were used for the dilution spotting on ammonia as the nitrogen source in minimal media. (B) Strains*met15***Δ**, *met15***Δ** *yll055w***Δ**, *met15***Δ** *gap1***Δ**, *met15***Δ** *gap1***Δ** *yll055w***Δ**, *met15***Δ** *agp1***Δ**, and *met15***Δ** *agp1***Δ** *yll055w***Δ** were analyzed for growth on minimal proline medium by dilution spotting.