



Corrigendum: Heat Shock Improves Random Spore Analysis in Diverse Strains of Saccharomyces cerevisiae

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A Corrigendum on

Heat Shock Improves Random Spore Analysis in Diverse Strains of Saccharomyces cerevisiae by Burke, M. K., McHugh, K. M., and Kutch, I. C. (2020). Front. Genet. 11:597482. doi: 10.3389/fgene.2020.597482

In the original article, there was a mistake in **Figure 2** as published. **The originally published figure had a panel out of order.** The corrected **Figure 2** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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FIGURE 2 | Site frequency spectrum of SNPs identified in the 4-parent population initially, after six outcrossing events, after 12 outcrossing events, and after 12 outcrossing events + a heat shock of 55° C for 20 min. Color in included in the histograms to emphasize the bins with high counts (red) compared to those with low counts (green); notice the reduction in sites with frequency = 0 or 1 between cycle 12 and cycle 12 + heat shock.