Corrigendum

Recruitment of a chromatin remodelling complex by the Hog1 MAP kinase to stress genes

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Since the publication of this article, the authors have been alerted to an error in Figure 6 of this article.

During the assembly of Figure 6, a control panel (lane 6) was mistakenly re-used in lane 3 with the same control purpose. As these control lanes contain naked DNA from isogenic strains, the conclusions of the figure remain however unchanged.

The correct Figure 6 is shown here.

The authors apologize for any confusion these errors might have caused.

Figure 6 Hog1 mediates major changes in nucleosome positioning in response to high osmolarity. Chromatin and naked DNA from wild-type and *hog1Δ* cells treated with (+NaCl) or without (–NaCl) were digested with micrococcal nuclease (MN), digested with *EcoRI* restriction enzyme, resolved in agarose gels and hybridized with a DNA probe to map nucleosomes at chromosomal *STL1*. The depicted scheme shows the location of nucleosomes (ovals) at the *STL1* gene in the wild-type strain without stress.