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Spontaneous mutations in diploid *Saccharomyces cerevisiae*: another thousand cell generations – ERRATUM

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Last two columns of Table 3 should be deleted and Table 3 legend should read:

(A) ML estimates (and 95% confidence intervals) of the genome-wide, haploid mutation rate (U), the absolute value of the average effect of a mutation ($E(hs)$) and the proportion of mutations that are beneficial (P) at transfer 50 (151 MA lines) and transfer 100 (149 MA lines). Transfer 50 estimates are from Joseph & Hall (2004). Corrected estimates of the parameters, indicated by asterisks (*), were generated by performing the Otto & Orive (1995) correction for the effects of natural selection during colony growth (see text). (B) Bateman–Mukai estimates of the genome-wide, haploid mutation rate (U) and the absolute value of the average effect of a mutation ($E(hs)$).

Reference

Hall, D. W., Mahmoudizad, R., Hurd, A. W. & Joseph, S. B. (2008). Spontaneous mutations in diploid *Saccharomyces cerevisiae*: another thousand cell generations. *Genetics Research* **90**, 229–241.