

CORRIGENDA

In the article by K. W. BROMAN (GENETICS 169: 1133–1146) entitled “The Genomes of Recombinant Inbred Lines,” on page 1137 (right column, center), the two-locus joint probabilities for the autosomes of eight-way RILs by sib mating should read:

$$\Pr(g_1 = i, g_2 = j) = \begin{cases} \frac{1-r}{8(1+6r)} & \text{if } i = j \\ \frac{r}{8(1+6r)} & \text{if } i \neq j. \end{cases}$$

In addition, on page 1138 (left column, end of first full paragraph), the map expansion for the X chromosome in 2^n -way RIL by sib mating should read:

$$\frac{2}{3}(n + 4).$$

In the article by C. M. McDONALD, K. F. COOPER and E. WINTER (GENETICS 171: 901–911) entitled “The Amal-Directed Anaphase-Promoting Complex Regulates the Smk1 Mitogen-Activated Protein Kinase During Meiosis in Yeast” the authors reported in Figures 3 and 4 that a *cdc28-4/cdc28-4 ama1-Δ/ama1-Δ* strain could form spores. Subsequent analysis showed that the strain referred to as *cdc28-4/cdc28-4 ama1/ama1* (CMY37 in Table 1) contains an intact copy of *AMA1*. The authors have generated new *cdc28-4/cdc28-4 ama1/ama1* strains and have found that the strains do not form spores. Therefore, the conclusion that *cdc28-4* can suppress the sporulation defect of strains lacking *AMA1* is in error. This change does not affect the other conclusions of this article.

In the article by Y. NAKASE, K. FUKUDA, Y. CHIKASHIGE, C. TSUTSUMI, D. MORITA, S. KAWAMOTO, M. OHNUKI, Y. HIRAKO and T. MATSUMOTO (GENETICS 173: 569–578) entitled “A Defect in Protein Farnesylation Suppresses a Loss of *Schizosaccharomyces pombe tsc2⁺*, a Homolog of the Human Gene Predisposing to Tuberous Sclerosis Complex,” on page 571 (left column), the first sentence of the *Microarray analysis* section should have read:

Details have been described in GEO (<http://www.ncbi.nlm.nih.gov/geo/index.cgi>) under the accession no. GSE4449.

This change does not affect the conclusions of the article.