

Erratum

In the January 2000 issue of the *Journal*, in the article “Microsatellite Polymorphism in the Heme Oxygenase-1 Gene Promoter Is Associated with Susceptibility to Emphysema,” by Yamada et al. (66:187–195), the repeat numbers of cloned alleles used as size markers were wrong. We reanalyzed the length of the (GT)_n repeats in smokers with or without chronic pulmonary emphysema (CPE), using new cloned alleles with repeat numbers sequenced with the ABI Prism Dye Terminator sequencing kit (Perkin-Elmer Applied Biosystems). The allele of each

sample from the subjects was also cloned, and the length of the (GT)_n repeats was confirmed by sequencing. The distribution of the numbers of (GT)_n repeats shown in figure 1b of the article needs to be corrected. However, the proportion of allele frequencies in classes L (≥33 repeats), M (27–32 repeats), and S (<27 repeats), as well as the proportion of genotypic frequencies in the group with class L alleles (L/L, L/M, and L/S), is the same as that originally reported. A corrected version of figure 1b appears below.

