subsequent interpretation; for example, when only synonymous polymorphisms are analyzed for the haplogroup L mtDNA sequences, the correlation coefficient is 0.002, by use of the r^2 measure of LD. Finally, the following minor corrections to this article should be noted: (1) In figure 1, the African mtDNA network should include a transition at nucleotide position 5262 in sequences 180 and 242. This polymorphism is specific to this L3e clade. (We thank Dr. Hans-Jürgen Bandelt [Universität Hamburg] for bringing this error to our attention.) (2) A transition at nucleotide position 921 is present in all of the L3d haplotypes, and it should have been included in the African mtDNA network (fig. 1). (3) The transversion at nucleotide position 3796 is present exclusively in haplogroup L1c mtDNA sequences 194 and 207 (fig. 1)—that is, this polymorphism is not present in a haplogroup H mtDNA sequence, and the 3796 transversion should not have been listed in table 2; instead, the *transition* at nucleotide position 3796 is present in four (not three, as we reported) mtDNA sequences from haplogroup H and in one mtDNA sequence from haplogroup B (table 2). The authors regret these errors.

Reference

Elson JL, Andrews RM, Chinnery PF, Lightowlers RN, Turnbull DM, Howell N (2001) Analysis of European mtDNAs for recombination. Am J Hum Genet 68:145–153

In the July 2002 issue of the *Journal,* in the article, "Analysis of the *RNASEL* Gene in Familial and Sporadic Prostate Cancer," by Wang et al. (71:116–123), the following two errors occur in the text: (1) In the 15th line in the left-hand column on page 118, "ratio 19:1" should be "ratio 99:1" and (2) in the 26th line in the left-hand column on page 119, "exon 5" should be "exon 6." The authors regret these errors.