

Erratum

In "Genotypes of Alcohol-metabolizing Enzymes in Japanese with Alcohol Liver Diseases: A Strong Association of the Usual Caucasian-Type Aldehyde Dehydrogenase Gene ($ALDH_2^1$) with the Disease," (*AJHG* 43:774-748, 1988), the figure legend should be as follows:

Figure 1 Hybridization of human genomic DNAs with the oligonucleotide probes for the ADH_2 locus. DNA samples were digested with *EcoRI* and *PvuII*, electrophoresed in 0.9% agarose gel, and hybridized with oligonucleotide probe-1, which is complementary to the usual ADH_2^1 allele (A), and were washed out and rehybridized with oligonucleotide probe-2, which is complementary to the atypical ADH_2^2 allele (B). Probe 1: 3' CCT·TAG·ACA·GCG·TGT·CTA·CTG 5'. Probe 2: 5' GCA·ATC·TGT·CAC·ACT·GAT·GAC 3'. (Position of base substitution is underlined). Sample 11 is identified to be homozygous usual 1-1; samples 2 and 10 are identified to be heterozygous atypical 1-2; and samples 1 and 3-9 are identified to be homozygous atypical 2-2.

Figure 2 Hybridization of human genomic DNAs with the oligonucleotide probes for the $ALDH_2$ locus. DNA samples were digested with *PstI*, electrophoresed in 0.9% agarose gel, and hybridized with oligonucleotide probe-1, which is complementary to the usual $ALDH_2^1$ allele (A), and were washed out and rehybridized with oligonucleotide probe-2, which is complementary to the atypical $ALDH_2^2$ allele (B). Probe 1: 3' C·CGT·ATG·TGA·CTT·CAC·TTT·TG 5'. Probe 2: 5' G·GCA·TAC·ACT·AAA·GTG·AAA·AC 3'. (Position of the base substitution is underlined). Samples 1, 4, and 8 are identified to be homozygous usual 1-1; samples 2, 3, 5-7 and 9 are identified to be heterozygous atypical 1-2; and samples 10 and 11 are identified to be homozygous atypical 2-2.

In table 1, the title should read "Genotypes of the ADH_2 Locus in Patients with Alcoholic Liver Disease and Controls." In table 2, the gene frequency headings should read " $ALDH_2^1$ " and " $ALDH_2^2$ ".

NOTICE

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