line in figure 1 of (2). Chakraborty and Kidd refer to more than 2000 U.S. court cases that have employed DNA evidence [(2),reference 4]. This is misleading, as the majority are paternity cases in which the DNA types of mother, child, and putative father are usually available for comparison. Chakraborty and Kidd compare binning the variable number of tandom repeat (VNTR) sequences with the grouping of A_1 and A_2 blood type alleles. This comparison is inappropriate because the problem with binning VNTRs is not that alleles are grouped but that sometimes they are assigned to the wrong bin. Their statement that the "worst case" scenario is an equal mixture of Poles and Italians is incorrect-it is actually the best case for their argument. Their statement [(2), p. 1737] that the "arithmetic and underlying principles are identical" for linkage equilibrium and Hardy-Weinberg equilibrium is simply incorrect. In a quote from our article, they also did not include an important qualifier present in our original text [(2), reference 6].

Wills states in his letter that our arguments are based on "old" blood group data. Why Wills would disregard reliable blood group data is unclear, because even Chakraborty and Kidd concede their relevance. Wills also cites a high mutation rate among some VNTRs as the basis of a "mutational churning process" that is "a very large reason for the relative uniformity of allele frequencies from one human group to another...." This hypothesis "explains" what has not yet been shown to exist, as relative uniformity of allele frequencies is precisely the point in dispute. We see no evidence of VNTR "churning" in French and Israelis or in the South American Indian tribes studied by Kidd et al. (6) [(1), p. 1749]. Wills also makes much of the observation that one of our subheadings (1) is identical to the title of a 1902 pamphlet by V. I. Lenin. This coincidence has no more relevance to DNA typing than the fact that Wills' letter makes favorable reference to Japanese automobile companies.

Austad concedes the validity of our arguments, but points out that DNA typing is certainly better than polygraph results, so why all the fuss? His argument seems to be that new sources of scientific evidence should be held to a standard of reliability no greater than methods currently in use. We would make a fundamental distinction between the intrinsic limitations of a technology and limitations imposed by the use of false assumptions, particularly when simple alternatives are available. For example, although polygraph examinations have a high intrinsic error rate, we suppose that Austad would object to a test in which an electrical short in the machine produced additional



Nanosphere[™] Size Standards. Certified in billionths of a meter by Duke Scientific

Nanosphere Size Standards are calibrated in billionths of a meter (nanometers) and are available in 22 sizes from 21 to 900nm-all traceable to the National Bureau of Standards. Nanospheres are part of our complete line of spherical particles from 0.02 to 2000 micrometers in diameter. They are used as standards for instrument calibration, quality control, filter checking, and in numerous biotechnology applications. At Duke Scientific-established in 1971-we have the expertise and resources to meet any of your requirements for microspheres and particles. Call us today for information.



erratic readings. Our view is that erroneous

assumptions about genetic uniformity

among ethnic groups are no more necessary

to DNA typing than electrical shorts are

dispute is not about the use of DNA evidence

in the courtroom. DNA typing is a very pow-

erful procedure. We regard it as "possibly the

most powerful innovation in forensics since the

development of fingerprinting in the last part

of the 19th century" [(1), p. 1746]. All we ask

is a basic degree of candidness in reporting the

statistical significance of a match. With databas-

es as large as x = 10,000, why not use 1/x as a

conservative estimate of the probability [(1), p].

1749]? After all, 0.0001 is already a pretty

small number. Why invoke unsupported as-

sumptions in order to obtain a still smaller

probability that is exaggerated and unreliable?

Perhaps it is because the organizations whose

interests are served by numerical exaggeration

have also been in charge of choosing the statis-

Washington University School of Medicine,

RICHARD C. LEWONTIN

Harvard University,

DANIEL L. HARTL

Cambridge, MA 02138

Department of Genetics,

St. Louis, MO 63110

Museum of Comparative Zoology,

tical procedures.

We would finally like to emphasize that this

necessary to polygraph machines.

1135D San Antonio Road, Palo Alto, CA 94303, Toll Free (800) 334-3883, in CA (415) 962-1100, Fax (415) 962-0718



Circle No. 34 on Readers' Service Card

REFERENCES

- 1. R. C. Lewontin and D. E. Hartl, Science 254, 1745 (1991).
- R. Chakraborty and K. K. Kidd, *ibid.*, p. 1735.
 U.S. v. Yee, 129 Federal Rules Decisions 629 (Northern District of Ohio, 1990).
- (Northern District of Ohio, 1990).
 F. M. Salzano and S. M. Callegari-Jacques, South American Indians: A Case Study in Evolution (Oxford Univ. Press, New York, 1988).
- 5. R. Chakraborty et al., Am. J. Hum. Genet. 50, 145 (1992).
- 6. J. R. Kidd et al., Human Biol. 63, 775 (1991).

Erratum: In the abstract and in the text (line 35 in the middle column of page 185) of the report "Electrical resistivity and stoichiometry of K_xC_{60} films" by G. P. Kochanski *et al.* (10 Jan., p. 184), the minimum resistivity was given incorrectly as 2.2 microohm-cm. The correct value is 2.2 milliohm-cm.

Erratum: In the News & Comment article "Is homosexuality biological?" by Marcia Barinaga (30 Aug., p. 956), it was suggested incorrectly that the suprachiasmatic nucleus is not part of the hypothalamus.

Erratum: The Table of Contents for the issue of 31 January 1992 (p. 508) incorrectly listed a letter by J. Bello as appearing in the Letters section beginning on page 514. The letter appeared in the issue of 14 February on page 784.

Erratum: In figure 1 (p. 1509) of the Research Article "Radar images of Mars" by D. O. Muhleman *et al.* (27 Sept., p. 1508), the Mars longitude of the sub-Earth point was mislabeled in each of the six snapshot radar images of Mars. None of the labels should have contained a decimal point. The values of λ in the labels should have been 78, 92, 104, 120, 133, and 147, respectively.