

FOR THE RECORD

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Genetic Diversity of Sixteen STRs in the Peruvian Mesa Redonda Lima Population

POPULATION: Mesa Redonda (Lima-Peru)

KEYWORDS: forensic science, DNA typing, identifiler, Peru, Lima, Mesa Redonda, population genetics, multiplex PCR, STR

Blood samples were collected from relatives of the missing persons of the fire on December 29, 2001 at Mesa Redonda, in the area of Lima, Peru. The biological samples used in this study consisted of blood stored on FTA[®] cards (Whatman Bioscience) from 151 unrelated individuals (87 females, 64 males). DNA was purified of the FTA[®] according to manufacturer's protocols.

Sixteen locus were amplified and typed using the AmpFℓSTR Identifiler Kit and ABI 310 Genetic Analyzer (PE Applied Biosystems) with GeneScan 3.7 and Genotyper 3.7 NT software according to manufacturer's protocol.

Allele frequencies were estimated by gene counting. Evaluation of Hardy-Weinberg expectations and other forensic statistical parameters was performed with the GDA software, version 1.1 (1) and PowerStats, Promega (2), respectively.

The complete dataset is available upon request from Dr. Beatriz Lizárraga, Laboratorio Biomolecular y Genética, Instituto de

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TABLE 1—Allele frequency distributions.