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D20S161 data for three ethnic populations and forensic validation

Amendment to the designation of the STR D20S161 alleles

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Erratum

Dear Sir,

We previously reported the allele distribution of D20S161 for various ethnic groups and then designated the alleles according to the number of repeat units (Hou et al. 1999; Kido et al. 2000). In the course of further population studies and sequence analysis, to our regret, we have found an error in counting the amplified fragment size and the number of repeat motifs. Therefore, we would like to correct the designation of D20S161 alleles as shown in Table 1.

Table 1 Correct designation of D20S161 alleles

Previous allele	Correct allele	Repeat number	Fragment size (bp)
14	13	13	166
16	15	15	174
17	16	16	178
18	17	17	182
19	18	18	186
20	19	19	190
21	20	20	194
22	21	21	198

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