



Article

Population data for 21 autosomal short tandem repeat markers in the Arabic population of the United Arab Emirates

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Supplementary Table 1: Population differentiation exact test between the two UAE population datasets for each locus

Locus	P-Value
D3S1358	0.27590±0.07084
vWA	0.51635±0.05404
D16S539	0.46340±0.05499
CSF1PO	0.27370±0.6450
TPOX	0.02595±0.01628*
D8S1179	0.31850±0.04799
D21S11	0.50080±0.04294
D18S51	0.05430±0.01989
D2S441	0.09855±0.04905
D19S443	0.44905±0.06866
TH01	0.99075±0.00272
FGA	0.00945±0.00366*
D22S1045	0.0560±0.0057*
D5S818	0.10040±0.03256
D13S317	0.52585±0.08496
D7S820	0.20525±0.05905
SE33	0.44045±0.08680
D10S1248	0.77245±0.04410
D1S1656	0.22980±0.05511
D12S391	0.36060±0.06192
D2S1338	0.14675±0.04035

Note: Statistically significant p-value ($P < 0.05$) indicated in bold (20,000 Markov steps done)

*Before the Bonferroni Correction