

**Table S2** Allele size adjustments used to make the African data set comparable to the combined HGDP-CEPH, Native American, Latino, Jewish, Asian Indian, CGP, and Pacific Islander data set

ID in combined data set	ID in African data set	Amount added to genotypes in the African data set (c*)
D10S1412	ATA31G11P	-7
D11S1304	UT2095M	-1
D11S1392	GATA6B09P	-7
D11S1984	GGAA17G05P	-7
D11S1998	GATA23E06L	-3
D11S1999	GATA23F06L	-3
D11S2000	GATA28D01M	-1
D12S1042	ATA27A06P	-7
D12S1045	ATA29A06P	-7
D12S1052	GATA26D02M	-1
D12S1064	GATA63D12P	-7
D12S1300	GATA85A04M	-1
D12S1301	GATA91H06M	-1
D12S2070	ATA25F09M	-1
D13S1807	GATA11C08P	-7
D13S787	GATA23C03P	-7
D13S796	GATA51B02M	-1
D14S608	GATA43H01M	-1
D14S617	GGAA21G11L	-2
D15S1515	GATA197B10P	-7
D15S816	GATA73F01M	-1
D16S2624	GATA81D12M	-1
D16S3253	GATA22F09P	-7
D17S1298	GAAT2C03P	-7
D17S2180	ATC6A06M	-1
D17S2195	ATA58A02P	-7
D18S1376 <sup>a</sup>	GATA185C06Z	-17
D19S589	GATA29B01L	1
D19S591	GATA44F10P	-7
D1S1596	GATA26G09P	-7
D1S1612	GGAA3A07M	-1
D1S1627	ATA25E07M	-1
D1S3669	GATA29A05P	-7
D20S477	GATA29F06Z	-1
D22S686	GGAA10F06M	-1
D2S1352	ATA27D04P	-7
D2S1384	GATA52A04M	-1
D2S1391	GATA65C03M	-1
D2S1394	GATA69E12M	-1
D2S1400	GGAA20G10M	-1
D2S2944	GATA30E06P	-7
D2S2968	GATA178G09M	-1
D3S1744	GATA3C02M	-1
D3S1768	GATA8B05M	-1
D3S2427	GATA22F11NZ	58

D3S2432	GATA27C08P	-7
D3S4529	GATA128C02M	-1
D4S1627	GATA7D01ZP	37
D4S1652	GATA5B02M	-1
D4S2366	GATA22G05M	-1
D4S2397	ATA27C07P	-7
D4S2417	GATA42H02P	-8
D5S1456	GATA11A11P	-7
D5S1462	GATA3H06M	-1
D5S1470	GATA7C06M	-1
D5S1480	ATA23A10M	-1
D5S1725 <sup>b</sup>	GATA89G08Z	31
D5S2488	ATA20G07M	-1
D6S1017	GGAT3H10M	-1
D6S1027	ATA22G07P	-7
D6S2436	GATA165G02M	-1
D7S1802	GATA41G07M	-1
D7S1818	GATA24D12P	-7
D7S2204	GATA73D10L	4
D7S3070	GATA189C06M	-1
D8S1048	UT7129L	1
D8S1110	GATA8G10M	-1
D8S1132	GATA26E03M	-1
D8S1477	GGAA20C10M	-1
D8S373	UT721M	-1
D8S592	GATA6B02P	-7
D9S1120	GATA81C04M	-1
D9S2169	GATA62F03M	-1
D9S910	ATA18A07M	-1
NA.D18S.1	GATA178F11Z	67
NA.D1S.3	GATA133A08Q	-4
TPO.D2S	SRAP	-7

<sup>a</sup>This locus was not present in the list of adjusted loci reported by Tishkoff *et al.* [3].

<sup>b</sup>Tishkoff *et al.* [3] used an adjustment of 27 nt.