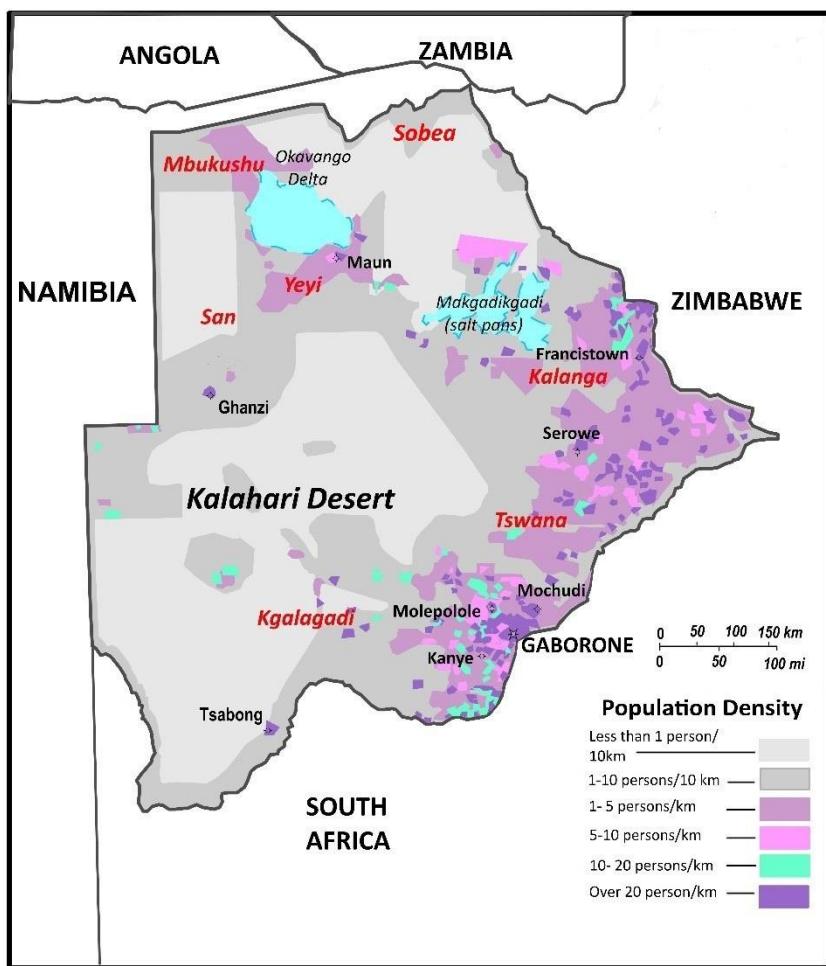


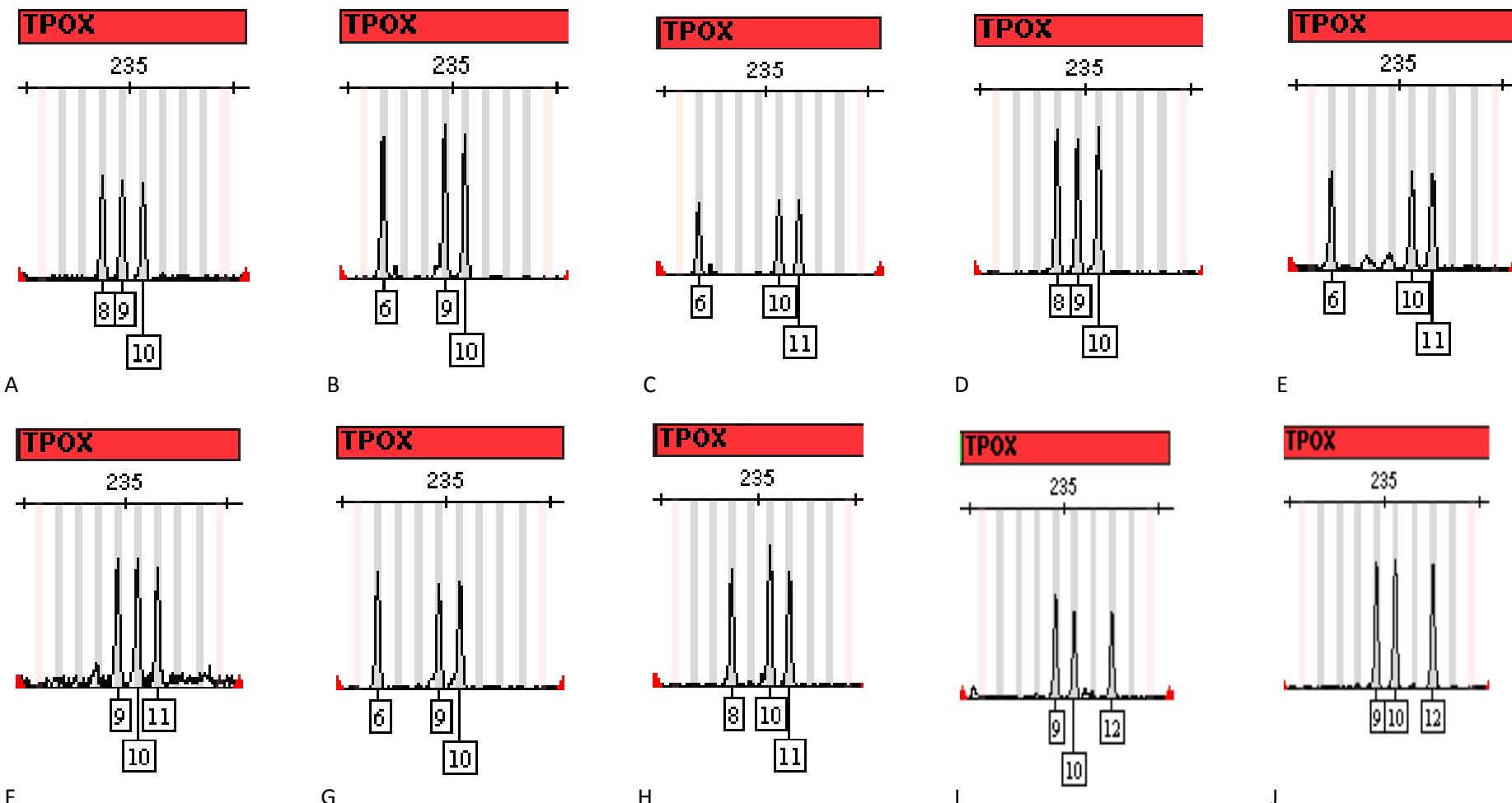
Genetic variation and population structure of Botswana populations as identified with AmpFLSTR Identifiler short tandem repeat (STR) loci

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Supplementary Information



Supplementary Figure S1 Representation of the geographic locations of the different ethno-linguistic language groups used in the study. The Niger-Congo Bantoid branch (Bantu) being: Central S-Bantu-Kgalagadi, Tswana and Kalanga; Central K-Bantu- Yeyi; and Central R-Bantu- Mbukushu. The Khoisan (San) are also seen in the picture. Map adapted from http://www.lib.utexas.edu/maps/africa/botswana-population_density-2005.png (Courtesy of the University of Texas Libraries, The University of Texas at Austin) using Surfer 9 (<http://www.ssg-surfer.com>).



Supplementary Figure S2 Electropherograms for samples exhibiting tri-allelic patterns for TPOX. Male samples are D and E.

Supplementary Table S1 Rare “variant” and off-ladder variants observed in Botswana with the Identifiler kit. Off-ladder variants marked with an *. Allele counts in this report are shown along with NIST number of records and publications for African populations along with their reported frequencies in brackets. See Supplementary Table S2 for allele frequencies reported in the study.

Allele	Current Study						NIST STR variant database (http://www.cstl.nist.gov/strbase/var_tab.htm)	African Published data		
	Bantu Central K	Bantu Central R	Bantu Central S	Khoisan Khoe-Kwadi	Khoisan Kx'a	Khoisan Tuu		South African	Namibia	African
D21S11 - tetranucleotide repeat ([TCTA] [TCTG])										
24.3*			3	1			36 reports			
33.1*			10				86 reports	[2]Zulu (0.0102), [6] Bantu (0.0022 – 0.0120), [7] (Bantu 0.0039)		[4] Gabon (0.014), [5] Mozambique (0.004)
34.1*			6	3			21 reports	[1] Coloured (0.0100), [2]Xhosa (0.0042)		
35.1*	1	1	1	3	4		19 reports	[1] Coloured (0.0150), Bantu (0.0100), [2]Zulu (0.0051), Coloured (0.0090)		
36.1*			1				6 reports	[1] Coloured (0.0210)		
37.1*			1				1 report			
39			3		1		1 report	[1] Coloured (0.0050)		
D7S820 - Tetranucleotide repeat (GATA)										
5					3		1 report			
FGA - complex Tetranucleotide repeat ([TTTC]3TTTTTCT[CTTT]nCTCC[TTCC]2)										

16			2					8 reports			
18.2		2	8	1				10 reports	[2] Xhosa (0.0042)		[4] Gabon (0.005), [5] Mozambique (0.0292), [8] Mozambique (0.010), [9] Angolan (0.0146)
19.2			9	1				7 reports			[3] African (0.0050), [4] Gabon (0.005), [5] Mozambique (0.010), [8] Mozambique (0.0292), [9] Angolan (0.0010)
21.2			8	1				8 reports	[1] Coloured (0.0210), [2] Coloured (0.0045), Xhosa (0.0042), Zulu (0.0051)		
22.2	1		1					19 reports	[1] Coloured (0.0150) [2] Coloured (0.0045), Xhosa (0.0042), Zulu (0.0051)		[3] African (0.0100)
25.2				1				9 reports	[2] San (0.007)		
29.2			1					3 reports	[2] Xhosa (0.0042)		[5] Mozambique (0.003)
CSF1PO- Tetranucleotide repeat (AGAT)											
16					1			6 reports			
D16S539- Tetranucleotide repeat (GATA)											
17*					1			1 report			
D19S433- Tetranucleotide repeat (AAGG)(AAAG)(AAGG)(TAGG)[AAGG]ⁿ											
7		4	30	10	5		1	5 reports	[1] Coloured (0.0520), Bantu (0.0500), [2] (Coloured (0.0225)	[1] Khoe (0.0190)	
D18S51- Tetranucleotide repeat (GAAA)											
15.2			7	10	2	1		22 reports	[1] Coloured (0.0310), Bantu (0.0100), [2] Xhosa (0.0126), Zulu (0.0051)	[1] Khoe (0.0380)	[3] African (0.0030), [8] Mozambique (0.0033), [9] Angolan (0.0021)
18.2			1					5 reports			[3] African (0.0050)

19.2	1		3				3 reports	[2] Zulu (0.0051)		[3] African (0.0030)
21.1			1				3 reports			
21.2			1				13 reports	[2] Zulu (0.0153)		[9] Angolan (0.0010)

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Supplementary Table S2 Allele frequencies and Forensic summary statistics for each of the 15 Identifier STR loci found in the Botswana samples (n=990).

Allele	D8S1179	D21S11	D7S820	CSF1PO	D3S1358	TH01	D13S317	D16S539	D2S1338	D19S433	vWA	TPOX	D18S51	D5S818	FGA
5			0.002					0.005							
6				0.066		0.077						0.051		0.001	
7			0.007	0.053		0.309				0.025		0.026		0.001	
8			0.210	0.048		0.386	0.024	0.028				0.311		0.069	
9			0.136	0.048		0.169	0.009	0.186		0.002		0.274	0.002	0.035	
9,3						0.037									
10			0.308	0.233		0.020	0.050	0.146		0.018		0.107	0.011	0.045	
10,2														0.006	
11	0.027		0.218	0.197		0.002	0.283	0.356		0.057	0.005	0.226	0.001	0.193	
12	0.120		0.102	0.291	0.002		0.408	0.170		0.122		0.004	0.037	0.370	
12,2										0.034					
13	0.192		0.019	0.054			0.157	0.101		0.299	0.010		0.042	0.270	
13,2										0.053				0.003	
14	0.343			0.008	0.102		0.066	0.007		0.206	0.075		0.040	0.015	
14,2										0.067				0.004	
15	0.218			0.001	0.335		0.003	0.002	0.018	0.053	0.191		0.122	0.002	
15,2										0.031				0.011	
16	0.090			0.001	0.330				0.063	0.021	0.264		0.176		0.001
16,2										0.012					
17	0.009				0.190			0.001	0.068	0.001	0.203		0.164		0.001
18	0.001				0.041				0.072	0.001	0.147		0.134		0.010
18,2														0.001	0.006
19					0.001				0.160		0.077		0.121		0.054
19,2										0.088		0.020		0.068	0.005
20															0.057

21									0.175		0.005		0.026		0.091
21,1												0.001			
21,2												0.001		0.005	
22									0.158		0.002		0.017		0.169
22,2														0.002	
23									0.076		0.001		0.005		0.181
24									0.045			0.001		0.185	
24,2	0.002								0.001						
24,3	0.002														
25	0.001								0.052			0.004		0.102	
25,2														0.001	
26	0.003								0.020			0.002		0.068	
27	0.064								0.005			0.001		0.019	
28	0.231								0.001					0.016	
29	0.132													0.009	
29,2	0.001													0.001	
30	0.138													0.002	
30,2	0.008													0.004	
31	0.077														
31,2	0.072													0.005	
32	0.020														
32,2	0.082													0.002	
33	0.017														
33,1	0.006														
33,2	0.034														
33,3	0.001														
34	0.024														
34,1	0.005														
34,2	0.008														

35		0.034															
35,1		0.006															
35,2		0.002															
36		0.011															
36,1		0.001															
37		0.012															
37,1		0.001															
38		0.008															
39		0.002															
42																0.001	
42,2																0.007	
43,2																0.002	

Supplementary Table S3 Genetic diversity estimators for the Botswana populations: overall (A), according to the Bantu (B-D) and Khoisan (E-G) ethno-linguistic groups, and according to districts (H-P). The Observed heterozygosity (Hobs), expected heterozygosity (Hexp) and probability of Hardy-Weinberg equilibrium (HWE P). Hobs= observed heterozygosity. Significant p values ($P<0,0033$) are in bold. Values out of HWE that show Heterozygote deficiency indicated by an asterisk (*).

	A			B			C			D		
Locus	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P
D8S1179	0.7727	0.7738	0.0967	0.7441	0.7447	0.2290	0.75862	0.7658	0.9885	0.7879	0.7673	0.0688
D21S11	0.8549	0.8854	0.1890	0.8837	0.8738	0.1684	0.81034	0.8574	0.6888	0.8529	0.8717	0.6748
D7S820	0.7779	0.7870	0.1144	0.7441	0.7647	0.7829	0.7069	0.7643	0.7171	0.7832	0.7768	0.3912
CSF1PO	0.7804	0.8076	0.0007*	0.7209	0.7865	0.5515	0.7931	0.8115	0.4016	0.7724	0.7959	0.6330
D3S1358	0.7201	0.7293	0.8766	0.6976	0.7398	0.2769	0.7069	0.7117	0.7010	0.7275	0.7319	0.4994
TH01	0.7111	0.7188	0.2138	0.6976	0.7275	0.2921	0.6724	0.7163	0.1131	0.7182	0.7119	0.1035
D13S317	0.7175	0.7264	0.0973	0.5116	0.5641	0.5553	0.5000	0.6699	0.0014	0.7414	0.7289	0.6245
D16S539	0.7522	0.7718	0.3057	0.7209	0.7326	0.2737	0.77586	0.7823	0.6910	0.7631	0.7616	0.9347
D2S1338	0.8575	0.8859	0.1093	0.8604	0.8640	0.6568	0.91379	0.9039	0.7171	0.8498	0.8769	0.2828
D19S433	0.8023	0.8399	0.0006*	0.7907	0.829	0.7670	0.91379	0.8523	0.0062	0.8235	0.8311	0.2338
vWA	0.8048	0.8209	0.7287	0.8837	0.8459	0.9779	0.77586	0.8231	0.4979	0.8003	0.8178	0.2902
TPOX	0.7509	0.7591	0.1434	0.8139	0.7734	0.3943	0.77586	0.7733	0.1884	0.7631	0.7654	0.9037
D18S51	0.8716	0.8831	0.7349	0.8837	0.8730	0.1972	0.7931	0.8729	0.2608	0.8823	0.8761	0.9091
D5S818	0.7471	0.7479	0.0076	0.7209	0.7507	0.9813	0.58621	0.6735	0.0253	0.7445	0.7399	0.0800
FGA	0.8446	0.8752	0.1581	0.8604	0.8741	0.6342	0.8965	0.8694	0.6888	0.8513	0.8731	0.0695

E

F

G

Locus	Khoisan Khoe-kwadi (n=133)			Khoisan Kx'a (n=58)			Khoisan Tuu (n=32)		
	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P
D8S1179	0.7519	0.7976	0.7088	0.7586	0.8093	0.6632	0.8438	0.7813	0.8805
D21S11	0.9023	0.9071	0.4263	0.9483	0.9041	0.2102	0.8438	0.8909	0.0652
D7S820	0.7820	0.8135	0.0427	0.8103	0.7939	0.1783	0.7188	0.7589	0.0570
CSF1PO	0.8045	0.8075	0.7385	0.7414	0.8000	0.1201	0.7813	0.8036	0.1402
D3S1358	0.7143	0.7212	0.7289	0.7241	0.7199	0.2188	0.6563	0.5947	0.8124
TH01	0.7368	0.7313	0.6719	0.6897	0.7003	0.2799	0.6875	0.6319	0.5724
D13S317	0.6993	0.7271	0.0562	0.7414	0.6891	0.4463	0.6250	0.7574	0.1643
D16S539	0.7293	0.8059	0.3992	0.7414	0.7924	0.5114	0.6250	0.8155	0.0584
D2S1338	0.8496	0.8861	0.3649	0.9655	0.9039	0.4063	0.9063	0.8958	0.0517
D19S433	0.8120	0.8593	0.1875	0.6379	0.7892	0.0609	0.5000	0.8115	0.0013*
vWA	0.8120	0.8206	0.9994	0.6724	0.7712	0.0885	0.6875	0.7753	0.7238
TPOX	0.6842	0.7053	0.2157	0.8103	0.7490	0.0490	0.7500	0.6701	0.0018
D18S51	0.8421	0.8946	0.5419	0.9138	0.8903	0.9802	0.9063	0.9023	0.9808
D5S818	0.7444	0.7755	0.0810	0.7241	0.7555	0.0406	0.7813	0.7173	0.6173
FGA	0.8196	0.8762	0.1149	0.8966	0.8906	0.0275	0.8750	0.8487	0.1045

	H			I			J			K			L		
	Central District (n=264)			Ghanzi District (n=168)			Kgalagadi District (n=20)			Kgatleng District (n=21)			Kweneng District (n=106)		
Locus	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P
D8S1179	0.7917	0.7563	0.1558	0.7738	0.7791	0.8184	0.8500	0.7923	0.9916	0.8095	0.8060	0.4429	0.8208	0.7786	0.2719
D21S11	0.8296	0.8677	0.1074	0.8869	0.8984	0.0375	0.8500	0.8885	0.7854	0.8571	0.8966	0.3985	0.8113	0.8683	0.6147
D7S820	0.7576	0.7711	0.2422	0.7321	0.8109	0.0318	0.9000	0.8064	0.1632	0.9048	0.7979	0.1585	0.8113	0.7752	0.7307
CSF1PO	0.7765	0.7898	0.7569	0.8214	0.8252	0.0023	0.6500	0.8013	0.0076	0.8571	0.8211	0.3982	0.8491	0.8068	0.4976
D3S1358	0.7689	0.7403	0.6279	0.6667	0.6784	0.5516	0.7500	0.7397	0.6453	0.7619	0.7549	0.6121	0.7359	0.7089	0.5400
TH01	0.7349	0.7014	0.1629	0.7024	0.7183	0.5698	0.6000	0.6180	0.8197	0.7143	0.6771	0.9085	0.6415	0.7206	0.2787
D13S317	0.7424	0.7415	0.4299	0.6786	0.6992	0.3339	0.7000	0.7769	0.2006	0.6191	0.6481	0.3524	0.7925	0.7314	0.4064
D16S539	0.7538	0.7641	0.4543	0.7202	0.8055	0.0129	0.7500	0.7756	0.7256	0.7143	0.7456	0.2767	0.7642	0.7359	0.3399
D2S1338	0.8182	0.8784	0.3588	0.8810	0.8917	0.2036	1.0000	0.8808	0.8813	0.8571	0.8862	0.3985	0.8868	0.8813	0.0073
D19S433	0.8220	0.8314	0.6908	0.7262	0.8402	0.0007*	0.7500	0.8397	0.3050	0.8571	0.9001	0.1621	0.7642	0.8154	0.5287
vWA	0.8447	0.8279	0.8909	0.8095	0.8010	0.9790	0.8500	0.8295	0.1221	0.8571	0.8420	0.5947	0.7642	0.8170	0.0466
TPOX	0.7424	0.7720	0.7687	0.7202	0.6952	0.0410	0.8000	0.7295	0.3789	0.6191	0.7096	0.1475	0.7736	0.7473	0.1849
D18S51	0.8864	0.8748	0.5201	0.8571	0.9000	0.4641	0.8500	0.9051	0.2641	0.9524	0.8432	0.7488	0.8868	0.8774	0.6548
D5S818	0.7500	0.7294	0.7622	0.7500	0.7824	0.0486	0.6000	0.7013	0.0730	0.7619	0.7689	0.9127	0.7453	0.7290	0.5017
FGA	0.8485	0.8741	0.0710	0.8155	0.8690	0.0018	0.8500	0.8782	0.4449	0.9048	0.9013	0.7433	0.8679	0.8789	0.1010

	M			N			O			P		
	Northeast District (n=45)			Northwest District (n=260)			Southeast District (n=32)			Southern District (n=74)		
Locus	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P	Hobs	Hexp	HWE P
D8S1179	0.7111	0.7738	0.5358	0.7462	0.7845	0.2919	0.8784	0.7550	0.2963	0.7973	0.7793	0.2805
D21S11	0.8444	0.8652	0.6275	0.9039	0.8958	0.2975	0.9063	0.8720	0.6853	0.8784	0.8590	0.6166
D7S820	0.8444	0.7758	0.0172	0.7846	0.7758	0.5823	0.7500	0.7674	0.9807	0.7838	0.7917	0.6176
CSF1PO	0.7556	0.7800	0.9794	0.7269	0.7942	0.1754	0.7500	0.8199	0.5721	0.7297	0.8121	0.1355
D3S1358	0.6222	0.7239	0.5922	0.7231	0.7358	0.5184	0.6875	0.7282	0.0856	0.6757	0.7411	0.1860
TH01	0.7333	0.7086	0.9956	0.7077	0.7258	0.0801	0.6563	0.7406	0.5800	0.8514	0.7289	0.0773
D13S317	0.6444	0.7316	0.0586	0.6615	0.6989	0.0142	0.6875	0.7064	0.4648	0.7568	0.7199	0.7431
D16S539	0.7111	0.7328	0.9665	0.7731	0.7731	0.1642	0.8125	0.7818	0.4018	0.7162	0.7911	0.2414
D2S1338	0.8444	0.8674	0.1625	0.8846	0.8904	0.8162	0.8438	0.8557	0.7053	0.8514	0.8689	0.4514
D19S433	0.8444	0.8315	0.5433	0.8346	0.8444	0.5295	0.8750	0.8581	0.5565	0.8108	0.8038	0.2050
vWA	0.8667	0.8162	0.9352	0.7269	0.8173	0.0619	0.8438	0.7986	0.9537	0.7162	0.8014	0.1353
TPOX	0.7333	0.7718	0.7762	0.7885	0.7768	0.0456	0.6563	0.7634	0.6332	0.7973	0.7570	0.7061
D18S51	0.8000	0.8582	0.2877	0.8654	0.8841	0.7004	0.8438	0.8765	0.0024	0.9324	0.8865	0.4182
D5S818	0.7556	0.7636	0.3835	0.7077	0.7258	0.3855	0.7813	0.7361	0.4285	0.6622	0.7605	0.1607
FGA	0.7556	0.8527	0.1670	0.8692	0.8751	0.4001	0.7813	0.8447	0.6009	0.9324	0.8800	0.4232

Supplementary Table S4 Maximum Likelihood (ML) Silent allele frequencies for the Botswana populations: overall (A). according to the Bantu (B-D) and Khoisan (E-G) ethno-linguistic groups. and according to districts (H-P).

	A	B	C	D	E	F	G
Locus	Overall population (n=990)	Bantu Central K (n=43)	Bantu Central R (n=58)	Bantu Central S (n=646)	Khoisan Khoe-kwadi (n=133)	Khoisan Kx'a (n=58)	Khoisan Tuu (n=32)
D8S1179	0.006	0	0	0.002	0.027	0.017	0
D21S11	0.012	0	0.031	0.012	0	0	0.025
D7S820	0.006	0	0.023	0	0.009	0	0.044
CSF1PO	0.014	0.022	0	0.005	0	0.024	0.031
D3S1358	0.002	0.023	0	0	0	0.015	0
TH01	0	0.02	0.013	0	0	0	0
D13S317	0.01	0.008	0.083	0	0.022	0	0.083
D16S539	0.009	0	0	0	0.032	0.012	0.09
D2S1338	0.013	0	0	0.017	0.017	0	0
D19S433	0.024	0.027	0	0.006	0.034	0.078	0.17
vWA	0.01	0	0.019	0.001	0	0.052	0.038
TPOX	0.005	0	0	0	0.015	0	0.009
D18S51	0.009	0.013	0.045	0	0.018	0	0
D5S818	0.013	0	0.031	0.004	0.018	0.023	0
FGA	0.008	0	0	0.01	0.019	0	0.004

H I J K L M N O P

Locus	Central District (n=264)	Ghanzi District (n=168)	Kgalagadi District (n=20)	Kgatleng District (n=21)	Kweneng District (n=106)	Northeast District (n=45)	Northwest District (n=260)	Southeast District (n=32)	Southern District (74)
D8S1179	0	0.001	0	0	0	0.041	0.021	0.02	0
D21S11	0.023	0.004	0	0	0.037	0	0.001	0	0
D7S820	0.004	0.036	0	0	0	0.012	0	0	0
CSF1PO	0.007	0.011	0.051	0	0	0	0.019	0.01	0.009
D3S1358	0	0	0	0	0	0.042	0.009	0.026	0.032
TH01	0	0.013	0	0	0.02	0	0	0.014	0
D13S317	0	0.016	0.018	0	0	0.039	0.024	0	0
D16S539	0	0.041	0.017	0.02	0	0.009	0	0	0.044
D2S1338	0.037	0.003	0	0.021	0.007	0.004	0	0	0
D19S433	0.012	0.07	0.032	0	0.017	0	0.005	0	0
vWA	0	0	0	0.006	0.024	0	0.04	0	0.045
TPOX	0.012	0	0	0.063	0	0.014	0	0.038	0
D18S51	0	0.013	0.036	0	0.003	0.022	0.015	0.004	0
D5S818	0	0.024	0.023	0	0	0	0.014	0	0.048
FGA	0.01	0.018	0.02	0	0.003	0.037	0	0	0

Supplementary Table S5 Pairwise genetic distance (F_{ST}) (below diagonal) and P-values (above diagonal) of the different Bantu and Khoisan ethno-linguistic groups in Botswana. Significant values ($P<0.003$) are indicated by an asterisk *.

	Central K	Central R	Central S	Khoe	Kx'a	Tuu
Central K	-	0.21226	0.00158	0.00000	0.00000	0.00000
Central R	0.00174	-	0.00020	0.00000	0.00000	0.00000
Central S	0.00402*	0.00377*	-	0.00000	0.00000	0.00000
Khoe	0.01363*	0.00921*	0.01024*	-	0.00000	0.00000
Kx'a	0.02312*	0.01905*	0.02026*	0.00656*	-	0.00000
Tuu	0.03069*	0.02819*	0.03009*	0.01622*	0.02183*	-

Supplementary Table S6 Pairwise genetic distance (F_{ST}) (below diagonal) and P-values (above diagonal) of the nine administrative districts in Botswana. Significant values ($P<0.0003$) are indicated by an asterisk *.

	Central	Ghanzi	Kgalagadi	Kgatleng	Kweneng	Northeast	Northwest	Southeast	Southern
Central	-	0.00000	0.19444	0.78071	0.11118	0.88457	0.00000	0.18384	0.09395
Ghanzi	0.01456*	-	0.02792	0.00050	0.00000	0.00000	0.00000	0.00000	0.00000
Kgalagadi	0.00208	0.00571	-	0.75775	0.12771	0.23681	0.35393	0.44897	0.30344
Kgatleng	-0.00171	0.01017	-0.00298	-	0.65855	0.78052	0.50045	0.43996	0.57519
Kweneng	0.00071	0.01792*	0.00288	-0.00128	-	0.23235	0.00000	0.10593	0.06881
Northeast	-0.00125	0.01623*	0.00276	-0.00217	0.00101	-	0.50718	0.68290	0.07663
Northwest	0.00269*	0.01238*	0.00107	0.00011	0.00589*	0.00005	-	0.02505	0.00000
Southeast	0.00141	0.01190*	0.00058	0.00044	0.00231	-0.00091	0.00349	-	0.50431
Southern	0.00104	0.01226*	0.00144	-0.00061	0.00161	0.00254	0.00526*	-0.00004	-

Supplementary Table S7 Forensic statistics metric for each locus for the Botswana populations: the overall population (A), according to the Bantu (B-D) and Khoisan (E-G) ethno-linguistic groups, and according to districts (H-P). Discrimination capacity (DC), match probability (MP), combined random match probability, polymorphic information capacity (PIC), typical paternity index (TPI), combined paternity index, power of exclusion (PE), and combined power of exclusion (CPE).

A

Locus	Overall population (n=990)				
	PE	MP	PD	PIC	TPI
D8S1179	0.558	0.086	0.914	0.742	2.2
D21S11	0.585	0.053	0.947	0.805	2.4
D7S820	0.558	0.079	0.921	0.752	2.2
CSF1PO	0.549	0.061	0.939	0.782	2.2
D3S1358	0.459	0.119	0.881	0.684	1.7
TH01	0.452	0.124	0.876	0.673	1.7
D13S317	0.439	0.119	0.881	0.678	1.7
D16S539	0.511	0.079	0.921	0.747	2.0
D2S1338	0.72	0.024	0.976	0.876	3.6
D19S433	0.492	0.112	0.888	0.70	1.9
vWA	0.582	0.056	0.944	0.794	2.3
TPOX	0.516	0.097	0.903	0.723	2.0
D18S51	0.744	0.025	0.975	0.873	3.9
D5S818	0.477	0.107	0.893	0.705	1.8
FGA	0.698	0.029	0.971	0.862	3.3
Combined	0.999996541	6.28×10^{-19}			

	B					C					D				
Locus	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI
D8S1179	0.499	0.131	0.868	0.700	1.9	0.524	0.091	0.909	0.724	2.0	0.585	0.0934	0.906	0.732	2.4
D21S11	0.662	0.104	0.895	0.849	3.0	0.520	0.074	0.925	0.834	2.0	0.607	0.0568	0.943	0.7925	2.5
D7S820	0.163	0.287	0.712	0.718	0.9	0.240	0.226	0.773	0.718	1.1	0.387	0.2179	0.782	0.742	1.5
CSF1PO	0.336	0.142	0.857	0.745	1.3	0.391	0.136	0.863	0.779	1.5	0.382	0.1405	0.859	0.767	1.5
D3S1358	0.424	0.127	0.872	0.685	1.6	0.439	0.143	0.856	0.651	1.7	0.477	0.1215	0.878	0.685	1.8
TH01	0.424	0.137	0.862	0.671	1.6	0.421	0.143	0.856	0.656	2.2	0.457	0.1335	0.866	0.661	1.7
D13S317	0.197	0.254	0.745	0.493	1.0	0.179	0.177	0.822	0.608	0.9	0.468	0.1295	0.870	0.685	1.8
D16S539	0.336	0.214	0.785	0.683	1.3	0.450	0.172	0.827	0.740	1.7	0.404	0.1525	0.847	0.729	1.5
D2S1338	0.715	0.051	0.948	0.839	3.5	0.823	0.033	0.966	0.887	5.8	0.696	0.0283	0.971	0.864	3.3
D19S433	0.496	0.094	0.905	0.799	1.9	0.651	0.210	0.789	0.830	2.9	0.485	0.1116	0.888	0.697	1.8
vWA	0.7623	0.056	0.943	0.815	4.3	0.555	0.066	0.933	0.791	2.2	0.607	0.0584	0.941	0.793	2.5
TPOX	0.599	0.680	0.320	0.730	2.5	0.240	0.308	0.691	0.728	1.1	0.126	0.3990	0.601	0.726	0.8
D18S51	0.795	0.071	0.928	0.849	5.0	0.585	0.045	0.954	0.852	2.4	0.747	0.0312	0.968	0.863	4.0
D5S818	0.363	0.162	0.837	0.704	1.4	0.196	0.225	0.774	0.610	1.0	0.399	0.1568	0.843	0.697	1.5
FGA	0.702	0.050	0.949	0.849	3.4	0.772	0.053	0.946	0.848	4.5	0.680	0.0346	0.965	0.860	3.1
Combined	0.999991	5.91 x10 ⁻¹⁴				0.999976	1.05x10 ⁻¹⁴				0.999986	6.28 x10 ⁻¹⁶			

E

F

G

Locus	Khoisan Khoe-kwadi (n=133)					Khoisan Kx'a (n=58)					Khoisan Tuu (n=32)				
	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI
D8S1179	0.5130	0.074	0.925	0.764	2.0	0.524	0.071	0.928	0.774	2.0	0.682	0.103	0.896	0.733	3.2
D21S11	0.6531	0.041	0.925	0.896	2.9	0.898	0.08	0.920	0.887	10	0.487	0.113	0.886	0.866	1.9
D7S820	0.4557	0.200	0.799	0.783	1.7	0.543	0.242	0.757	0.756	2.1	0.378	0.283	0.716	0.708	1.5
CSF1PO	0.3294	0.182	0.817	0.782	1.3	0.209	0.160	0.839	0.763	1.05	0.330	0.157	0.842	0.764	1.3
D3S1358	0.2409	0.308	0.691	0.675	1.1	0.466	0.152	0.847	0.667	1.8	0.363	0.212	0.787	0.552	1.4
TH01	0.4875	0.118	0.881	0.687	1.9	0.412	0.145	0.854	0.655	1.6	0.409	0.197	0.802	0.581	1.6
D13S317	0.4174	0.133	0.866	0.686	1.6	0.418	0.197	0.802	0.651	1.6	0.226	0.173	0.826	0.703	1.0
D16S539	0.3662	0.124	0.875	0.775	1.4	0.409	0.156	0.843	0.754	1.6	0.165	0.163	0.836	0.775	0.9
D2S1338	0.6941	0.030	0.969	0.871	3.3	0.930	0.037	0.962	0.887	14.5	0.808	0.064	0.935	0.870	5.3
D19S433	0.5214	0.103	0.896	0.843	2.0	0.232	0.142	0.857	0.763	1.1	0.202	0.094	0.905	0.769	1.0
vWA	0.6216	0.057	0.942	0.793	2.6	0.386	0.101	0.898	0.728	1.5	0.409	0.099	0.900	0.725	1.6
TPOX	0.4507	0.591	0.408	0.653	1.7	0.187	0.500	0.500	0.707	1	0	0.555	0.444	0.611	0.5
D18S51	0.6490	0.027	0.972	0.882	2.8	0.817	0.036	0.963	0.871	5.6	0.795	0.04	0.960	0.880	5
D5S818	0.4202	0.156	0.843	0.742	1.6	0.407	0.158	0.841	0.708	1.5	0.477	0.210	0.789	0.658	1.8
FGA	0.6051	0.038	0.961	0.861	2.5	0.754	0.062	0.937	0.872	4.2	0.736	0.0884	0.911	0.815	3.8
Combined	0.999978	1.21×10^{-15}				0.999999	1.52×10^{-14}				0.999958	6.28×10^{-19}			

H**I****J**

Locus	Central District (n=264)					Ghanzi District (n=168)					Kgalagadi District (n=20)				
	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI
D8S1179	0.584	0.103	0.897	0.719	2.4	0.551	0.086	0.914	0.742	2.2	0.695	0.105	0.895	0.737	3.3
D21S11	0.609	0.066	0.934	0.784	2.5	0.769	0.028	0.972	0.887	4.4	0.695	0.060	0.940	0.853	3.3
D7S820	0.523	0.088	0.912	0.735	2.1	0.480	0.066	0.934	0.781	1.9	0.795	0.140	0.860	0.752	5.0
CSF1PO	0.556	0.076	0.924	0.758	2.2	0.639	0.062	0.938	0.800	2.8	0.355	0.135	0.865	0.751	1.4
D3S1358	0.543	0.120	0.880	0.695	2.2	0.379	0.156	0.844	0.632	1.5	0.510	0.140	0.860	0.673	2.0
TH01	0.484	0.151	0.849	0.646	1.9	0.432	0.120	0.880	0.677	1.7	0.291	0.179	0.821	0.557	1.3
D13S317	0.497	0.108	0.892	0.699	1.9	0.396	0.138	0.862	0.656	1.6	0.428	0.130	0.870	0.720	1.7
D16S539	0.906	0.090	0.910	0.730	1.3	0.460	0.069	0.931	0.774	1.8	0.510	0.110	0.890	0.725	2.0
D2S1338	0.633	0.028	0.972	0.865	2.8	0.757	0.028	0.972	0.879	4.2	1.000	0.075	0.925	0.845	0
D19S433	0.640	0.050	0.950	0.811	2.8	0.470	0.049	0.951	0.820	1.8	0.510	0.085	0.915	0.798	2.0
vWA	0.684	0.056	0.944	0.803	3.2	0.617	0.071	0.929	0.770	2.6	0.695	0.125	0.875	0.782	3.3
TPOX	0.497	0.089	0.911	0.733	1.9	0.460	0.166	0.834	0.643	1.8	0.599	0.170	0.830	0.671	2.5
D18S51	0.768	0.033	0.967	0.860	4.4	0.709	0.022	0.978	0.889	3.5	0.695	0.065	0.935	0.872	3.3
D5S818	0.510	0.124	0.876	0.684	2.	0.510	0.088	0.912	0.748	2.	0.291	0.185	0.815	0.625	1.3
FGA	0.692	0.032	0.968	0.860	3.3	0.628	0.039	0.961	0.853	2.7	0.695	0.075	0.925	0.840	3.3
Combined	0.999999563	7.27×10^{-18}				0.99999684	2.14×10^{-18}				1.000000	5.21×10^{-15}			

K

L

M

	Kgatleng District (n=21)					Kweneng District (n=106)					Northeast District (n=45)				
Locus	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI
D8S1179	0.617	0.116	0.884	0.754	2.6	0.638	0.098	0.902	0.740	2.8	0.446	0.094	0.906	0.730	1.7
D21S11	0.709	0.061	0.939	0.864	3.5	0.620	0.035	0.965	0.851	2.7	0.684	0.049	0.951	0.842	3.2
D7S820	0.805	0.152	0.848	0.746	5.3	0.620	0.094	0.906	0.737	2.7	0.684	0.136	0.864	0.730	3.2
CSF1PO	0.777	0.135	0.865	0.775	3.3	0.693	0.081	0.919	0.776	3.3	0.519	0.083	0.917	0.738	2.0
D3S1358	0.530	0.143	0.857	0.691	2.1	0.486	0.147	0.853	0.654	1.9	0.318	0.123	0.877	0.665	1.3
TH01	0.451	0.179	0.821	0.601	1.8	0.344	0.121	0.879	0.669	1.4	0.482	0.144	0.856	0.647	1.9
D13S317	0.314	0.202	0.798	0.580	1.3	0.585	0.126	0.874	0.684	2.4	0.348	0.128	0.872	0.681	1.4
D16S539	0.451	0.138	0.862	0.697	1.8	0.534	0.107	0.893	0.701	2.1	0.446	0.119	0.881	0.688	1.7
D2S1338	0.709	0.075	0.925	0.851	3.5	0.769	0.040	0.960	0.865	4.4	0.684	0.056	0.944	0.842	3.2
D19S433	0.709	0.070	0.930	0.868	3.5	0.534	0.059	0.941	0.790	2.1	0.684	0.064	0.936	0.802	3.2
vWA	0.709	0.088	0.912	0.798	3.5	0.534	0.073	0.927	0.789	2.1	0.728	0.075	0.925	0.780	3.8
TPOX	0.314	0.170	0.830	0.645	1.3	0.551	0.124	0.876	0.700	2.2	0.096	0.095	0.905	0.725	4.4
D18S51	0.903	0.093	0.907	0.800	10.5	0.769	0.037	0.963	0.860	4.4	0.599	0.053	0.947	0.832	2.5
D5S818	0.530	0.111	0.889	0.714	2.1	0.502	0.131	0.869	0.680	2.0	0.519	0.105	0.895	0.716	2.0
FGA	0.805	0.061	0.939	0.868	5.3	0.730	0.037	0.963	0.862	3.8	0.519	0.057	0.943	0.826	2.0
Combined	0.999999929	5.12×10^{-15}				0.99999927	5.12×10^{-15}				0.99999227	1.14×10^{-16}			

N**O****P**

Locus	Northwest District (n=260)					Southeast District (n=32)					Southern District (74)				
	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI	PE	MP	PD	PIC	TPI
D8S1179	0.503	0.079	0.921	0.752	2.0	0.565	0.129	0.871	0.705	2.3	0.594	0.098	0.902	0.740	2.5
D21S11	0.803	0.024	0.976	0.885	5.2	0.808	0.057	0.943	0.844	5.3	0.751	0.042	0.958	0.839	4.1
D7S820	0.571	0.086	0.914	0.739	2.3	0.510	0.098	0.902	0.716	2.	0.569	0.084	0.916	0.753	2.3
CSF1PO	0.471	0.068	0.932	0.763	1.8	0.510	0.074	0.926	0.782	2.0	0.476	0.068	0.932	0.782	1.9
D3S1358	0.465	0.119	0.881	0.688	1.8	0.409	0.162	0.838	0.669	1.6	0.392	0.123	0.877	0.690	1.5
TH01	0.440	0.122	0.878	0.678	1.7	0.364	0.117	0.883	0.688	1.5	0.698	0.165	0.835	0.677	3.4
D13S317	0.371	0.136	0.864	0.653	1.5	0.409	0.154	0.846	0.654	1.6	0.521	0.140	0.860	0.664	2.1
D16S539	0.550	0.088	0.912	0.739	2.2	0.622	0.119	0.881	0.734	2.7	0.454	0.088	0.912	0.753	1.8
D2S1338	0.764	0.024	0.976	0.879	4.3	0.683	0.059	0.941	0.825	3.2	0.698	0.044	0.956	0.849	3.4
D19S433	0.665	0.043	0.957	0.827	3.0	0.745	0.066	0.934	0.827	4.0	0.510	0.112	0.888	0.707	2.0
vWA	0.471	0.057	0.943	0.791	1.8	0.683	0.096	0.904	0.756	3.2	0.454	0.078	0.922	0.767	1.8
TPOX	0.578	0.091	0.909	0.740	2.4	0.364	0.102	0.898	0.709	1.5	0.594	0.112	0.888	0.711	2.5
D18S51	0.725	0.026	0.974	0.871	3.7	0.683	0.086	0.914	0.848	3.2	0.862	0.040	0.960	0.869	7.4
D5S818	0.440	0.119	0.881	0.683	1.7	0.565	0.150	0.850	0.675	2.3	0.372	0.098	0.902	0.716	1.5
FGA	0.733	0.032	0.968	0.861	3.8	0.565	0.061	0.939	0.811	2.3	0.862	0.043	0.957	0.861	7.4
Combined	0.99999863	1.02x10 ⁻¹⁸				0.99999835	5.56x10 ⁻¹⁶				0.999999666	4.25x10 ⁻¹⁷			

Supplementary Table S8 Population samples according to district and self-declared ethno-linguistic population group used in the study (n=990).

District	n	Khoisan	Bantu
Central	264	9	255
Ghanzi	168	156	12
Kgalagadi	20	6	14
Kgatleng	21	-	21
Kweneng	106	1	105
Northeast	45	-	45
Northwest	260	66	194
Southeast	32	-	32
Southern	74	-	74
total	990	238	752

Supplementary Table S9 Population samples according to their self-declared Bantu and Khoisan ethno-linguistic population groups (n=990).

Language Family	Branch	Sub-group	n
Niger-Congo	Narrow Bantu (Bantu)	Central K ¹	43
		Central R ²	58
		Central S ³	646
		Undeclared Bantu	5
Khoisan	Southern African Khoisan (SAK)	Khoe-Kwadi	133
		Kx'a	58
		Tuu	32
		Undeclared Khoisan	15

¹Central K- Subiya, Mbukushu

²Central R- Yeyi

³Central S- Kgalagadi, Kalanga, Tswana (and its dialects-Barolong, Bakwena, Bangwaketse, Balete, Bakgatla, Barolong, Bangwato, and Batawana)