

Y-chromosomal STR haplotypes in an Arab population from Libya

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Abstract. We analyzed eight Y-chromosomal STRs (DYS385, DYS19, DYS 389I and II, DYS390, DYS391, DYS392, DYS393) in a population sample of 63 males from Tripolis (Libya) and found 63 different haplotypes. © 2005 Elsevier B.V. All rights reserved.

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1. Introduction

Y-chromosomal STRs are increasingly used in population genetic, linkage evolution and for forensic applications [1]. In the last years a large set of population data have been collected and multiplex protocols have been reported. However, an in-depth evaluation of their population genetic properties requires a large number of haplotype frequencies from as many populations as possible.

2. Material and methods

To establish a DNA database on Libyans, blood samples were taken from 63 unrelated Arab individuals, the area of Tripolis. DNA was extracted from these unrelated male blood samples according to standard Qiagen procedures QiAmp Blood Kit. Amplifications were

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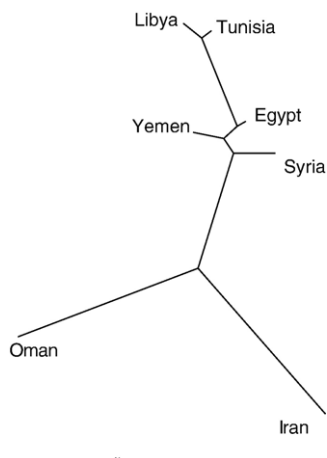


Fig. 1. Cluster analysis based upon R_{ST} with data from Libya and six other Arab populations.

performed using fluorescent dye labelled primers according to Elmoznino and Prinz (<http://www.ystr.org/europe/>). A set of eight Y-chromosomal STR loci (DYS19, $DYS389-I$, $DYS389-II$, $DYS390$, $DYS391$, $DYS392$, $DYS393$, $DYS385$) were analyzed.

Table 1
Allelic frequencies and forensic efficiency parameters of Y-STR loci in 63 unrelated Libyans

	DYS 19	DYS 389 I	DYS 389 II	DYS 390	DYS 391	DYS 392	DYS 393	DYS 385-a	DYS385-b
Libya_1	15	12	30	24	10	10	13	14	15
Libya_2	14	13	30	22	11	11	12	13	18
Libya_3	14	13	30	22	11	11	12	13	18
Libya_4	14	13	30	23	11	11	12	14	18
Libya_5	13	12	29	25	11	11	13	16	18
Libya_6	13	13	31	20	11	11	13	11	14
Libya_7	13	14	30	24	11	11	13	13	14
Libya_8	14	13	29	24	11	11	13	13	19
Libya_9	15	14	30	24	10	13	13	13	15
Libya_10	14	12	29	23	10	11	13	16	17
Libya_11	15	13	30	22	10	11	14	14	15
Libya_12	14	12	29	22	10	11	14	15	16
Libya_13	14	13	29	22	11	11	12	13	16
Libya_14	13	14	30	24	11	11	14	13	15
Libya_15	13	13	29	22	10	11	13	13	16
Libya_16	13	14	30	24	11	11	14	13	15
Libya_17	14	13	29	24	11	12	12	11	14
Libya_18	14	12	29	24	11	13	13	14	16
Libya_19	14	13	31	24	11	11	14	14	14
Libya_20	13	13	31	24	11	12	13	15	16
Libya_21	14	13	30	23	11	11	14	13	19
Libya_22	14	13	29	23	11	11	11	13	14
Libya_23	15	13	30	23	10	13	14	13	14
Libya_24	13	13	29	23	11	11	13	13	14
Libya_25	14	12	28	22	10	11	12	13	18
Libya_26	11	12	30	24	10	12	13	16	18
Libya_27	13	13	29	24	10	10	13	12	13
Libya_28	15	13	29	25	9	10	13	11	14
Libya_29	13	14	30	18	9	10	13	12	13
Libya_30	15	15	32	22	11	9	14	15	16
Libya_31	13	15	31	23	9	10	13	12	14
Libya_32	13	12	29	22	10	10	13	16	17
Libya_33	15	12	30	25	10	10	12	13	14

(continued on next page)

Table 1 (continued)

	DYS 19	DYS 389 I	DYS 389 II	DYS 390	DYS 391	DYS 392	DYS 393	DYS 385-a	DYS385-b
Libya_34	14	14	31	22	11	10	12	12	18
Libya_35	13	14	29	24	9	11	13	13	15
Libya_36	13	14	30	24	9	11	13	13	15
Libya_37	13	14	30	24	9	11	13	13	15
Libya_38	15	14	31	20	10	12	14	12	17
Libya_39	13	13	29	24	9	11	13	13	14
Libya_40	13	14	30	23	9	11	13	13	17
Libya_41	15	11	29	25	10	11	12	14	15
Libya_42	13	12	29	24	11	11	13	17	17
Libya_43	13	13	29	24	11	11	13	13	15
Libya_44	13	14	30	25	11	11	13	13	14
Libya_45	15	13	30	23	11	11	12	13	18
Libya_46	14	13	30	24	11	12	12	13	20
Libya_47	13	14	30	24	9	11	13	13	14
Libya_48	13	14	30	25	9	11	13	14	15
Libya_49	13	14	30	24	9	11	14	13	15
Libya_50	13	14	30	24	9	11	14	13	15
Libya_51	13	14	30	24	9	11	14	13	15
Libya_52	16	12	29	22	10	10	14	16	16
Libya_53	13	14	30	24	9	11	13	13	14
Libya_54	13	14	29	24	9	10	13	13	15
Libya_55	13	13	29	24	9	11	13	13	15
Libya_56	14	14	30	23	11	11	12	13	13
Libya_57	15	12	29	25	10	11	12	13	17
Libya_58	13	14	30	24	9	11	13	13	15
Libya_59	15	13	31	24	10	12	12	16	17
Libya_60	13	14	30	24	9	11	13	13	14
Libya_61	13	12	29	22	10	11	13	16	17
Libya_62	15	12	31	25	10	11	12	14	15
Libya_63	13	13	29	24	9	11	13	13	15

Amplification products were analyzed by capillary electrophoresis using the ABI 310® Genetic Analyzer according to the manufacturer's recommendations (Applied Biosystems) and the genotyper software.

Forensically relevant statistical analyses were calculated using the Arlequin software.

3. Results and discussion

Among the 63 Libyan males studied 63 different Y-STR haplotypes were observed (Table 1).

Cluster analysis based upon R_{ST} revealed that the Y-STR core haplotype distribution of the Libyan group is close to the neighbouring Tunisian population whereas another North-African population (Egypt) and populations from the Near East (Syria, Yemen, Oman and Iran) are more distant, reflecting the growing geographical and cultural distance from Libya (Fig. 1).

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Reference

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