141 148 /1387 /1 18 / **RAPD** 86/7/14: -1 2 3 E-mail: Gh.Elyasi@Gmail.com DNA (RAPD) DNA DNA RAPD 18 16 25 DNA 20 24 0/98 0/94 (0/06)RAPD-PCR

RAPD DNA

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Study of Turkey Genetic Diversity Using RAPD Markers

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Abstract

Application of DNA marker technology in poultry as a tool for species and strain identification has progressed rapidly during the last decade. Random amplified polymorphic DNA (RAPD) markers can be generated using short arbitrary primers to amplify genomic DNA, giving a genotype-specific pattern of bands. RAPD analysis should lead to the saturation of the genome without the requirement of previous genetic information. The purpose of this study was to evaluate genetic similarities and distances among four-colored phenotype of turkey population that kept in Turkey Research Station of Iran using RAPD technique. Blood samples were taken from 20 Black, 16 Golden, 24 Pied and 18 White local turkeys. Genomic DNA was extracted from 25µl blood samples. RAPD technique, using 15 primers, was applied to amplification of different fragments of genome. The genetic similarity between the groups varied from 0.94 to 0.98 between phenotypic groups. The highest genetic distance (0.06) was determined between the White and Golden phenotypic groups and the lowest genetic distance (0.02) was obtained between White and Black groups. The results of this research showed that RAPD-PCR is an appropriate tool for evaluation of genetic variation in poultry and color is not a usful character to differentiate turkey genetic groups.

Key Words: DNA marker, Genetic distance, RAPD, Turkey

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143 RAPD

.(2002)

(1996)

DNA .

12

(r=0/8)

.(1996

.(1996 DNA

DNA (1998)

(2003) (2001) ^{1}PCR .

²RAPD PCR . DNA

²Random Amplified Polymorphic DNA ¹Polymerase Chain Reaction

/1 18 /		144 1387
DNA	(1998)	1507
•		
DNA	RAPD 70 5	
DNA	•	
DNA	Δ	
	10	
.(1) 1	5	
1/5) 25	RAPD .	
20 dNTP 0/2 MgC12	4 4	-
Taq DNA 1	RAPD	
50 1 Polymeras	se	
PCR . (DNA		
3 94		
40		
37 45 9	04	
2 72 1	. RAPD	
10 72		
70 6 %2	. RAPD	
	10	
V. Y	16 18	
Popgene 3.2	20 24 EDTA %10	
1 018-110	EDIN	
.(1973) Nei	25 DNA	20
	(1995) (1990)	

145 RAPD

1

		5'→3'	
1	Moh 01	TGGACTCGAG	
2	Moh 02	GCACTGAGTA	
3	Moh 04	GCATGCGATC	
4	Moh 06	ACGTCGAGCA	
5	Moh 07	TACGCAGACT	
6	Moh 11	TGCATCGTAC	
7	Moh 12	ACGCCGTACG	
8	Moh 13	GCTGCTCGAGT	
9	Moh 26	CGAACCTGATC	
10	Moh 27	GCTTGCAGATC	
11	OPC 02	GTGAGGCGTC	
12	OPC 05	GATGACCGCC	
13	OPC 08	TGGACCGGTG	
14	OPC 16	CACACTCCAG	
15	OPD 05	TGAGCGGACA	

DNA DNA

RAPD

.(2 1)

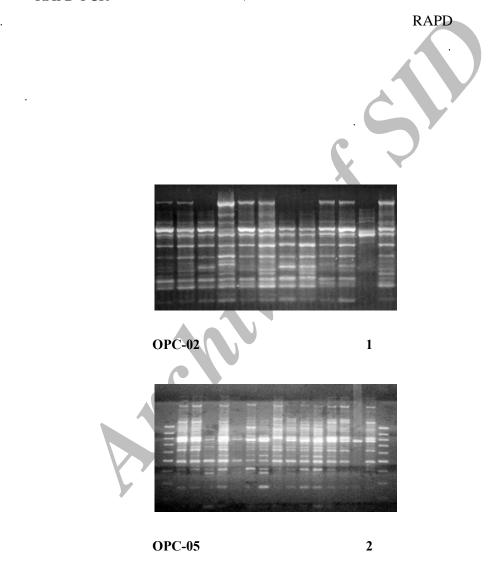
.5 (2)

RAPD .(2000

.(3 3 (2005) (2001) 2 5'→3'

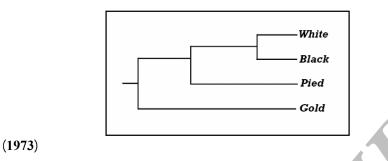
/1 18 /				146
_				1387
9	12	GTGAGGCGTC	OPC 02	_
14	18	GATGACCGCC	OPC 05	
9	12	TGGACCGGTG	OPC 08	
10	11	CACACTCCAG	OPC 16	
11	14	TGAGCGGACA	OPD 05	

RAPD-PCR



(1973) 3 0/96 0/97 0/98

47	RAPD		
0/95	0/94		
0/97			



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