# Nonmetric cranial variation in human skeletal remains from the Armenian Highland: microevolutionary relations and an intergroup analysis

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### SUMMARY

Nonmetric traits are frequently analyzed in the field of anthropology to measure genetic relatedness, or biodistance within or between populations. These studies are performed under the assumption that nonmetric traits are inherited genetically. Historically, interpretations of both biological and cultural change within the Armenian Highlands, have cited large-scale population movements. Biological estimates of these changes have traditionally relied upon biodistance estimates, using odontologic, craniofacial measures of both deformed and nondeformed skulls. In order to evaluate whether large-scale prehistoric and historic migrations occurred in the Armenian Highlands, we examine the biodistance results from nonmetric cranial traits for 19 mortuary samples that represented all time periods on the Armenian Highlands. None of the distances between each pair of mortuary samples examined in this study was significant. These results suggest biological continuity in the populations of Armenia. The biodistance results also suggest endogamy within inland populations. The broader implications of these results are also discussed.

**Key words:** Armenian Highlands – Biological anthropology – Archaeology – Osteology – Nonmetric traits – Biodistance

### INTRODUCTION

In the last 80 years (from Berry and Berry, 1967; Movsesyan et al., 1975; Cesnys, 1986, 1988; Kozintsev, 1988, to the recent papers by Prowse and Lovell, 1996; Christensen, 1997; Ishida and Dodo, 1997; Sutter and Mertz, 2004), a large body of literature has been dedicated to the assessment of the biological significance and importance of nonmetric traits of the skull (also named "discontinuous", "epigenetic", "discrete"). The importance that both the environment (Piontek, 1979, 1988; Hauser and Bergman, 1984; Bergman and Hauser, 1985; Bergman, 1993; Rubini et al., 1997) and heredity have in their expression has been evaluated. Regarding the potential use of such traits in studies of human populations, it is assumed that the phenotype (observable characteristics) of an individual will provide direct information about his or her genotype (genetic constitution). With regard to the important contributions regarding heredity

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Dr. Anahit Khudaverdyan. Institute of Archaeology and Ethnography, National Academy of Science, Yerevan 0025, Charents st.15, Republic of Armenia. Phone: +(374) 077133-254 / +(374) 565-884. E-mail: ankhudaverdyan@gmail.com; akhudaverdyan@mail.ru (Torgersen, 1951a, 1951b; Berry, 1975; Reinhard and Rösing, 1985; Rubini, 1997), we would particularly underline the report by Sjövold (1984). Nonmetric traits of the skeleton are therefore often used to assess genetic relatedness within (Cheverud and Buikstra, 1981; Kohn, 1991) and between (Movsesvan, 2005; Cesnys and Tutkuvienė, 2007; Matsumura, 2007; Saunders and Rainey, 2008) past populations. Understanding these relationships in past populations (especially those without written histories) can provide information about migration patterns, residence patterns, population structures, and human origins and evolution (Hanihara et al., 2003; Hlusko, 2004; Mclellan and Finnegan, 1990; Lane and Sublett, 1972; Turan-Ozdemir and Sendemir, 2006).

The term "biodistance" is commonly used to describe genetic relatedness. Saunders and Rainey (2008) describe biodistance as a measure of the amount of divergence; less divergence is equal to a closer genetic relationship (Saunders and Rainey, 2008; Sherwood et al., 2008). Christensen (1998) used biodistance analyses to trace the spread of the Zapotecan language family throughout Oaxaca, Mexico. By analyzing both nonmetric traits and linguistic data, he determined that people migrating from a central area were able to establish themselves in other areas of Oaxaca. These groups became distinct from the parent population in both genetics and in language dialects. Alt et al. (1997) studied the nonmetric traits of individuals in a triple burial in Dolce Vestonice. The data collected by this research team led them to conclude that the three were part of the same family. There are also various researchers who discuss the numerous factors that confound the heritability of nonmetric traits (Williams et al., 2005). Some factors that have been found to have a noticeable effect on the expression of these traits are geography, habitat, sexual dimorphism (differences in physical appearance between individuals of different sexes in the same species), age, nutrition, disease, size, and intertrait correlations (Berry, 1975; Cheverud et al., 1979).

The main purpose of this research project is to gain some insight into the expression of nonmetric traits on human skulls from the Armenian Highlands (from the Bronze Age to the beginning of the 20<sup>th</sup> century). The study of the heritability of some discontinuous traits based on a skeletal collection of individuals from various areas of the Armenian Highlands with known family relationships, has provided a new stimulus in the scientific debate.

## PREHISTORY AND HISTORY OF THE ARMENIAN HIGHLANDS

The Armenian Highlands (also known as the Armenian Uplands, the Armenian Plateau, or simply Armenia) is the central and highest of three land-locked plateaus that together form the northern sector of the Middle East (Hewsen, 1997). The present Armenian Republic (Fig. 1) is located in the South Caucasus on the eastern end of the Armenian Plateau. In early history, the Armenian highlands were a crossroads linking East and West (Martirosyan, 1964). Recent genetic studies confirmed that this avenue served not only for commerce and cultural diffusion, but also for the exchange of genes (Balaresque et al., 2010). From 4th millennium BC to 1st millennium BC, tools and trinkets of copper, bronze and iron were commonly produced in this region and traded in neighboring lands where those metals were less abundant (Krupnov, 1966; Trifonov, 1991; Nechitailo, 1991; Pystovalov 2002, etc.).

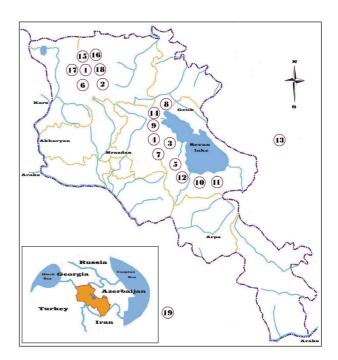


Fig. 1. Map showing the origins of the 19 Armenian Highland samples used in the present study: 1. Landjik, 2. Black Fortress, 3. Nerkin Getashen I, 4. Nerkin Getashen II, 5. Nerkin Getashen III, 6. Artik, 7. Karmir, 8. Sarukhan, 9. Arcvakar, 10. Karashamb, 11. Akunk, 12. Lchashen, 13. Shushi, 14. Karchakhpyur, 15. Shirakavan, 16. Beniamin, 17. Vardbakh, 18. Black Fortress I, 19. Crania Àrmenica

The invention in the Near East of wheeled vehicles and "kibetka-houses" on wheels allowed cattle drovers-farmers to move and survive with ease on the open steppes. Their movement across Eurasia in early times was not a military invasion, but a slow expansion caused by a decline in the child mortality rate and a resulting increase in population growth. The wide expanse of the Eurasian steppes, offering favorable conditions for human life and the spread of information and technology, facilitated a process of wide cultural integration in the Bronze Age throughout this area.

Craniological data have allowed identification of alien Mediterranean characteristics influencing several ethnic Eurasian groups and revealed evidence of a migratory stream from the Armenian Highland and the Caucasus (Khudaverdyan, 2011a). The Armenian Highland samples (Kura-Araxes Culture) and the Catacomb culture samples from Kalmykia, Ukraine, Dnieper, exhibit very close affinities to one another. If we follow a hypothesis put forward and developed by Gamkrelidze and Ivanov (1984) considering the ancestral home of Indo-European areas of the Armenian Highlands and adjoining territories, whence other tribes entered the Northern Black coast both through the Caucasus and through Central Asia and the Volga region (carriers of a Catacomb culture ceremony), it is necessary to assign that movement to Aryan tribes, which were one of the first to reach the Black Sea coast steppes through the Caucasus (or possibly by sea?). Khlopin (1983) connects the Catacomb culture with the Indo-Aryans, because catacomb burial rituals had roots in Southwestern Turkmenistan since the early IV millennium BC (Sumbar cemetery). Fisenko (1966) suggest that the Catacomb people were Proto-Hittites. Kuzmina (1998) is also a supporter of the Fisenko's hypothesis. Anthony (2007) assumed the Catacomb people to be ancestors of the Greeks, while Berzin and Grantovsky (1962), and Klejn (1980, 1984) determined that the Indo-Aryans originated from the Catacomb culture.

The Armenian Highland samples (Kura-Araxes culture) serve as a phenetic link between the Dnestr region and the Ukraine samples (Tripolye Culture), which feature the closest affinities to each other. Hence, it is possible to outline the cultural and ethnic communications in antiquity and the known role of the Armenian Highlands (Kura-Araxes culture) as the intermediary between the ancient area of distribution of Tripolye cultures and the Eastern countries (Passek, 1949; Martiroyan and Mnacakanyan, 1973; Lang, 2005).

The Armenian Highland samples and the Albashevo, Fatianovo, Balanovo Cultures and Timber Grave samples from the Volga region exhibit very close affinities to one another (Khudaverdyan, 2011b). The presence of the Mediterranean components was also remarked by Trofimova (1949) in carriers of the Fatianovo culture, Shevchenko (1984, 1986) and Khokhlov (2000) in carriers of the Timber Grave cultures of the forest-steppe Volga region, and also by Yusupov (1989) in the Southern Ural Mountains.

craniological The and odontological research indicates some morphological association of the Siberian samples (Eluninskaya and Andronovo Cultures) with populations from Caucasia and Turkmenistan (Solodovnikov, 2006; Zubova, 2008; Tur, 2009; Khudaverdyan, 2011a). The different rates of genetic drift and external gene flow may have contributed to the morphological differentiation and diversification amongst the different Eurasian populations. The initial starting area (or one of the intermediate areas), as indicated by the anthropological data, would seem to be the Armenian highlands, and the Caucasus as a whole.

In the Ancient (Classical) time (1<sup>st</sup> century BC - 3rd century AD) in the Armenian Highland and Caucasus there was interaction of different ethno-cultural units - Iranianspeaking nomadic (Scythians, Sarmatians, Sauromatians, Saka) (Herodotus IV; Strabo XI) and local. The advancement of the Scythians, Sarmatians and Saka in the territory of Armenian Highland and Transcaucasia was accompanied by not only an interaction of various cultural elements, but also a mixture of them. Detailed analysis of the anthropological materials from the Armenian Highland allows us to explain not only the complicated anthropological compound of the population but also to discover the reason for the anthropological and ethnic non-homogeneity in the populations of the Classical Age. Intragroup analysis revealed two groups within the population (Khudaverdyan, 2000, 2012). The dolichocephaly type in both cases is presented. The male skulls of the first group have been

diagnosed as classical European group. The second is the same European type, but the horizontal profile of the face (group II) in them is a little weakened. The female skull group has the same analogical image as the males. It is necessary to state that carriers of this complex remind one of the Scythians from the territory of the Dnestr region, Steppes of Black Sea Coast, Ukraine, the Sarmatians from the Volga region and the Saka from the territory of Turkmenistan (Khudaverdyan, 2012). The invasions of the various tribes led, in stages, to a mixture of outsiders among the native Armenians, and the dilution of their ranks on the plateau. The artificial modification of skulls (such as bregmatic, ring deformations of a head, were known in the classical population of the Beniamin, Shirakavan and Karmrakar, Vardbakh) and teeth in ancient Armenia may be related to the emerging social complexity and the need to differentiate among people, creating a niche for such highly visual bodily markers (Khudaverdyan, 2011c).

### MATERIALS AND METHODS

Eleven samples from 19 Armenian Highland sites were examined in this study (Fig. 1, Table 1). In the Early Bronze period (4000-3000 BC) farmer and cattle-breeder Landjik represent the Kuro-Arexes population of the Shirak Plateau (Khudaverdyan, 2009).

The Late Bronze period samples are represented by remains of three Armenian highlands sites. The combination of the remains from these four sites is justified for three reasons. First, the small sample sizes for certain sites (Landjik, Black Fortress) were inadequate (from 10 to 15 individuals) for subsequent biodistance analysis. Second, the Landjik, Black Fortress, and Artik sites represent a cemetery from Shirak Plain. Indeed, the geographic distance among sites is short. Finally, analysis of all the nonmetric cranial traits examined in this study revealed that no significant differences were present among the remains from the four sites, and hence the data from these sites were combined for subsequent statistical analyses. An adequate number of remains were available

	Sample name	Date	Researchers
1	Landjik	c. 4000-3000 BC	Khudaverdyan, 2009
2	Black Fortress	c. XIV-XII BC	Khudaverdyan, 2009
3	Nerkin Getashen I	c. XV BC	Movsesyan,1990
4	Artik	c. XV/XIV-XI BC	Movsesyan,1990
5	Total group: Landjik, Black Fortress, Nerkin Getashen I, Artik	I period	
6	Sarukhan	c. XI-IX/ VIII BC	Movsesyan,1990
7	Nerkin Getashen II	c. XIII-XII BC	Movsesyan,1990
8	Nerkin Getashen III	c. IX-VIII BC	Movsesyan,1990
9	Arcvakar	c. XI-IX/ VIII BC	Movsesyan,1990
10	Akunk	c. XI-IX/ VIII BC	Movsesyan,1990
11	Karashamb	c. XI-IX/ VIII BC	Movsesyan,1990
12	Karmir	c. XI-IX/ VIII BC	Movsesyan,1990
13	Lchashen	c. 3000 - 2000 BC	Movsesyan,1990
14	Shushi	c. 3000- 2000BC	Movsesyan,1990
15	Total group: Sarukhan, Nerkin Getashen II and III, Arcvakar, Akunk, Karashamb, Karmir, Lchashen, Shushi	II period	
16	Shirak Plateau (total group): Landjik, Black Fortress, Artik	I period	
17	Sevan region (total group): Sarukhan, Nerkin Getashen II and III, Arcvakar, Akunk, Karashamb, Karmir, Lchashen	II period	
18	Karchakhpyur-Shirakavan	c. 1 BC – AD 3	Movsesyan,1990
19	Beniamin, Vardbakh, Black Fortress I	c. 1 BC – AD 3	Khudaverdyan, 2000, 2005
20	Total group: Karchakhpyur-Shirakavan, Beniamin, Vardbakh, Black Fortress I	c. 1 BC – AD 3	
21	Bingel Dag	20 century	Bunak, 1927; Movsesyan, Kochar, 2001

Table 1. Armenian Highland craniological samples.

from the Artik site (Movsesyan, 1990), and were therefore analyzed as a single sample. Nine Late-Period (XI-IX/VIII BC) samples were analyzed in this investigation (Movsesyan, 1990). The different site designations for Nerkin Getashen I, Nerkin Getashen II, and Nerkin Getashen III represent different time periods, rather than spatially discrete cemeteries. The sites (Sarukhan, Nerkin Getashen II, Nerkin Getashen III, Arcvakar, Akunk, Karashamb, Karmir, Lchashen) (Movsesyan, 1990) included in Period II are located in the Sevan region. The remains from the Akunk and Lchashen sites (Movsesyan, 1990) were treated as independent samples, because a sufficient number of crania from these spatially discrete cemeteries was available for study. The Classical period (1st century BC – 3rd century AD) samples examined in this study include the remains from Karchakhpyur, Shirakavan (Movsesyan, 1990), Beniamin, Vardbakh, and Black Fortress I (Khudaverdyan, 2009). After the Armenian genocide in 1915, Bunak has

gathered a large collection (Museum of Anthropology, Moscow) of human skulls (i.e. the victims a genocide). The modern population include remains from these people (Bingel Dag: Armenians from Musha) (Bunak, 1927).

For this study, 24 non-metric (i.e., epigenetic) cranial and mandibular traits were used to assess the biological affinities (Table 2) among the 19 prehistoric and historic Armenian Highland mortuary samples examined here (Table 1). All traits examined in this study were successfully used in other biodistance studies, and their scoring procedures and descriptions are well-known in the literature (Berry and Berry, 1967; Movsesyan, 1975, 1990; Kozintsev, 1980, 1988; Česnys and Tutkuvienė, 2007). Non-metric cranial traits have successfully been used to evaluate the evolutionary relations and biological affinities among numerous archaeological samples (e.g., Blom, 1998; Ishida and Dodo, 1997). Non-metric cranial traits have the advantage of being scoreable from highly frag-

Table 2. A complete list of nonmetric traits analyzed and the methods used to score them.

Trait	Scoring Method
Sutura metopica	absent, complete
Foramen supraorbitale	presence/absence
Foramen infraorbitale accessorium	two distinct foramina, more than two distinct foramina
Foramen parietale	present (on parietal), present (sutural). Absent
Os bregmaticum	presence/absence
Os epiptericum	presence/absence
Os apicis lambdae	presence/absence
Os asterii	presence/absence
Ossa suturae coronalis	presence/absence
Ossicula suturae squamosae	presence/absence
Ossa suturae sagittalis	presence/absence
Ossa suturae lambdoïdeae	presence/absence
Canalis condylaris intermedius	patent, not patent
Canalis hypoglossi bipartite	complete (within canal)
Foramen mastoideum absent	absent, 1, 2, more than 2
Foramen mastoideum exsutural	absent, 1, 2, more than 2
Foramen spinosum bipertitum	partial formation
Foramina alatine minoranus	absent, 1, 2, more than 2 (the lesser palatine foramina lie on both sides of the posterior border of the hard palate, immediately posterior to the greater palatine foramen)
Foramina mentale accessorium	absent, 1, 2, more than 2
Cribra orbitale	presence/absence
Os zygomaticum bipartitum	presence/absence
Tuberculum praecondylare	presence/absence (immediately anterior and medial to the occipital condyle)
Torus palatinus	presence/absence

mented skeletal materials. Although some studies have reported that some non-metric cranial traits are influenced by cranial deformation (Konigsberg et al., 1993; Ossenberg, 1970), other studies have indicated that most non-metric cranial trait expressions are largely free of influence from artificial cranial deformation (Griffin, 1995; Khudaverdyan, 2000).

Data are subjected to the component and cluster analysis. The clustering procedures

produce branching-tree diagrams to illustrate similarities/differences among cases in complex data matrices, by forming clusters that minimize intracluster variation while maximizing intercluster variation. Inspection of the 24 non-metric cranial traits and frequencies retained for biodistance analysis indicate that some of the trait expressions demonstrate a geo-temporal trend.

 Table 3. Number of affected and observed crania, and their dichotomized trait frequencies by mortuary sample for 24 nonmetric cranial traits used in biodistance analysis\*.

Traits	Bronze	Age	Bronze	Iron Age	Artik		Akunk		Lchashe	n	Classica	Age	Classical	Age	Classical	Age	Modern	Arme- nians
Traits	I	period	II	period	Ι	period	П	period	II	period	Beniamin	et all	Shirak	avan	Total	group	Bingel	Dag
	A/	0	A/	0	A/	0	A/	0	A/	0	A/	0	A/	0	A/	0	R	
1 Sutura metopica	21.2/	54	12.3/	259	19.47/	36	4.35/	28	13.49/	126	17.8/	58	6.67/	30	12.3/	88	225	
2 Foramen supraorbital	54.5/	54	45.1/	222	36.11/	36	27.27/	22	44.00/	125	44.95/	82	40.7/	27	42.9/	108	38	
3 Cribra orbitale	48.4/	59	16.4/	235	11.11/	36	9.09/	22	8.00/	125	27.3/	76	8.33/	24	17.9/	100	105	
4 Foramen infraorbitale acces.	19.8/	65	9.71/	245	11.11/	36	4.76/	21	8.80/	125	38.2/	50	5.0/	20	21.6/	70	97	
5 Os zygomaticum bip.	13.92/	67	11.3/	251	2.78/	36	1.19/	21	6.50/	123	39.6/	52	5.56/	18	22.6/	70	46	
6 Os bregmaticum	2.8/	60	1.9/	259	2.78/	36	1.08/	23	0.22/	125	1.93/	52	0.83/	30	1.38/	82	10	
7 Ossa suturae coronalis	15.99/	67	10.4/	229	2.78/	36	4.35/	23	3.22/	125	10.1/	113	3.45/	29	6.8/	142	25	
8 Os epiptericum	35.9/	67	23.4/	220	17.14/	35	4.76/	21	20.00/	120	20.2/	57	21.74/	23	21.0/	80	163	
9 Ossicula suturae squamosae	18.7/	66	6.2/	228	2.78/	36	4.35/	23	3.34/	120	18.5/	89	0.92/	27	9.8/	116	15	
10 Os asterii	18.9/	65	7.8/	239	8.33/	36	1.08/	23	6.50/	123	10.9/	46	3.44/	29	7.2/	75	132	
11 Foramen parietale	56.4/	58	53.97/	236	47.22/	36	34.78/	23	43.65/	126	54.5/	80	46.42/	29	50.5/	109	387	
12 Os apicis lambdae	15.6/	66	12.7/	234	2.78/	36	8.69/	23	4.84/	124	10.0/	50	10.71/	28	10.4/	78	62	
13 Ossa suturae sagittalis	19.1/	65	3.97/	250	0.69/	36	1.08/	23	0.81/	124	5.4/	92	0.83/	29	3.2/	101	-	
14 Ossa suturae lambdoïdeae	48.8/	68	40.6/	253	22.22/	36	47.82/	23	25.60/	125	58.7/	101	28.58/	28	43.7/	129	341	
15 Foramen mastoideum absent	57.6/	65	42.98/	255	37.14/	36	21.40/	23	34.92/	126	65.8/	73	28.57/	28	47.2/	92	294	
16 Foramen mastoideum exsutural	29.6/	64	23.7/	252	17.14/	35	30.43/	23	34.92/	126	33.6/	87	28.57/	28	31.1/	115	248	
17 Canalis condylaris intermed.	62.3/	60	61.4/	228	53.33/	30	44.44/	18	63.93/	122	61.5/	45	46.66/	15	54.1/	60	697	
18 Canalis hypoglossi bipartite	25.5/	60	27.6/	227	23.33/	30	33.33/	18	22.13/	122	37.5/	24	46.66/	15	42.1/	39	333	
19 Tuberculum praecondylare	21.1/	60	8.9/	250	3.33/	30	5.55/	18	5.73/	122	27.3/	22	6.67/	15	16.99/	37	85	
20 Foramen spinosum bipertitum	30.5/	64	9.3/	227	11.42/	35	4.54/	22	8.06/	122	40.0/	25	25.92/	17	32.96/	42	62	
21 Torus palatinus	39.6/	62	22.7/	233	15.15/	33	16.67/	18	14.63/	124	40.9/	89	1.47/	17	21.2/	106	198	
22 Foramina palatina minoranus	52.9/	63	43.8/	230	40.24/	33	50.0/	14	38.84/	121	49.1/	60	52.94/	17	51.1/	77	-	
23 Sulcus mylobyoideus	18.4/	30	24.7/	116	-		-		19.35/	62	-		-		-		-	
24 Foramina mentale accessor	21.6/	29	9.2/	56	-		-		-		22.0/	55			22.0/	55	-	

\* O: number of crania actually observed; A: number of crania showing trait (affected); R - radians.

In total, the comparative analysis included 15 craniological series from the territory of Eurasia (2 samples /Fatianovo and Balanovo Cultures/ from the Volga region (Chesnis, 1986), 2 samples /Sapallitepe, Gonur Depe/ from Central Asia (Khodjaiov, 1977; Nevchaloda and Kufterin, 2008), 4 samples /Afanasevo, Andronovo, Karasukskaya and Tagarskaya Cultures/ from Siberia (Kozintsev, 1980), 4 samples /Chernyakhov Culture/ from Ukraine (Chesnis and Konduktorova, 1982), 2 samples /Budeshti, Malaeshti/ from the Dnestr region (Chesnis and Konduktorova,1982), 1 sample /Latgali/ from Latvia (Chesnis, 1986)). Kozintseva and Kozintseva's statistical package (Peter the Great Museum of Anthropology and Ethnography, St. Petersburg) was used.

### **RESULTS AND DISCUSSION**

The remaining 24 traits, their frequencies, and the number of individuals observed for each trait for the 9 Armenian Highland samples are provided in Table 3 (only Bingel Dag sample - in radians). More specifically, the presence of a sutura metopica, sagittal, squamosae ossicles, multiple infraorbital foramina, foramen spinosum, bridging of the mylohyoid groove, tuberculum praecondylare and palatine torus show a chronological trend between two samples (periods I and II: from the Bronze Age to the Iron Age). For the multiple infraorbital foramina, and os zygomaticum, the classical sample (Beniamin-Vardbakh-Black Fortress I) generally has a higher frequency of expression for these traits. For the palatine torus, the Artik, Akunk, Lchashen, and Karchakhpyur-Shirakavan samples have the lowest frequencies of expression, while the Beniamin-Vardbakh-Black Fortress I and all Bronze Age (period I) samples are characterized by relatively higher levels of expression. Coronal ossicles show a slighter temporal trend in Classical period samples than those of the Bronze period, characterized by higher frequencies of expression.

Analysis 1. Brothwell (1959) first applied an array of ten non-metrical characteristics to the study of multivariate distances among populations. The number of traits was further increased after Berry and Berry's paper (1967) was published. Those traits have frequently been employed not only to compare populations by the multivariate distances method, but also to study processes affecting the genetic variations in the population structure and to determine kinship among individuals, etc. The purpose of such an analysis is to gain some insight into the expression of non-metric traits on the 11 human samples from the Bronze Age to the beginning of the 20<sup>th</sup> century from the Armenian Highlands (N 4, 5, 10, 13, 15-21). The values of the first three factors are given in table 4. The differentiation that can be traced in the Armenian Highland populations is shown in Fig. 2.

Table 4. Elements of three initial components for 11 groups.

Trait	Ι	II	III
Sutura metopica (frontalis)	0.976	0.017	0.066
Foramen supraorbitale	-0.711	0.642	0.227
Foramen infraorbitale accessorium	0.486	0.693	-0.500
Os zygomaticum bipartitum	0.014	0.932	-0.253
Ossa suturae coronalis	-0.233	0.882	0.280
Os epiptericum	0.823	0.326	0.399
Os asterii	0.882	0.427	0.064
Foramen parietale	0.938	-0.090	0.007
Canalis condylaris intermedius	0.975	-0.003	0.114
Canalis hypoglossi bipartite	0.897	-0.290	-0.106
Values	58.403	29.200	6.361

As expected, the first axis accounts for the majority (58.5%) of the intergroup discrimination. Taking into account the character of connection of attributes in these coordinates. it is possible to tell that the large values till I coordinate axes correspond to groups with the sutura metopica (0.976), canalis condylaris intermedius (0.975), foramen parietale (0.938), canalis hypoglossi bipartite (0.897), os asterii (0.882) and os epiptericum (0.823). The negative weight gives a foramen supraorbitale (-0.711). The second factor (29.2%) is the maximum for os zygomaticum bipartitum (0.932), ossa suturae coronalis (0.882), foramens infraorbitale accessorium (0.693) and supraorbitale (0.642). The third factor accounts for 6.4% of the intergroup variance. The negative weight gives a foramen infraorbitale accessorium (-0.500).

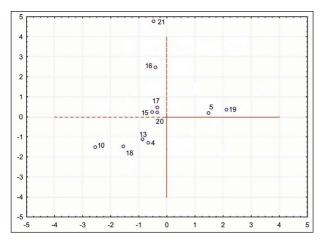


Fig. 2. Factor analysis: 4 – Artik, 5 – Armenian Highland /I period: Bronze Age/, 10 – Akunk, 13 – Lchashen, 15 – Armenian Highland /II period: Bronze and Early Iron Ages/, 16 – Shirak Plateau /I period/, 17 – Sevan region /II period/, 18 – Karchakhpyur-Shirakavan, 19 – Beniamin, Vardbakh, Black Fortress I, 20 – Armenian Highland /Classical Age/, 21 – Bingel Dag.

Figure 2, The graph obtained using the first two axes, shows how the groups from the Armenian Highland (II period and Classical Age) and the Sevan region (period II) are close to one another. Groups from the Karchakhpyur-Shirakavan, Shirak Plateau (Bronze Age: I period), Akunk and Bingel Dag are well distinguished from the groups from Armenian Highland. The graph shows how the groups from the Armenian Highland (Bronze Age (period 1) and Classical Age: Beniamin, Vardbakh, Black Fortress I) close to one another. The Artik and Lchashen samples also exhibit close affinities to one another.

Next, we applied cluster analysis (Fig. 3, table 5). In this diagram, the Lchashen, Akunk and Akunk samples are relatively close

to the Karchakhpyur-Shirakavan sample. Importantly, Ancient sample (Karchakhpyur-Shirakavan) is closely related to the previous samples (Bronze and Iron Ages). The most isolate Bingel Dag sample is shown in Figure 3. The prehistoric series, including the Shirak Plateau (I period), the Sevan region (II period) and the groups from periods I and II (Bronze and Early Iron Ages) are nearer the Ancient samples (total group and Beniamin-Vardbakh-Black Fortress I), as mentioned above.

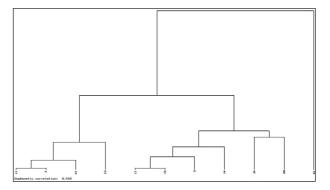


Fig. 3. Cluster tree: 4 – Artik, 5 – Armenian Highland /I period: Bronze Age/, 10 – Akunk, 13 – Lchashen, 15 – Armenian Highland /II period: Bronze and Early Iron Ages/, 16 – Shirak Plateau /I period/, 17 – Sevan region /II period/, 18 – Karchakhpyur-Shirakavan, 19 – Beniamin, Vardbakh, Black Fortress I, 20 – Armenian Highland /Classical Age/, 21 – Bingel Dag.

Analysis (2) of 11 series (N 4, 5, 10, 13, 15-21). The values for the first three factors are given in table 6. The characters are of different nature: sutural bones and varieties of openings, even cribra orbitalia, a pathological manifestation. It is known that pathological

Table 5. Matrix of distance, values for eleven Armen	an Highland mortuary samples examined in this study.
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Sample name	Lchashen	Akunk	Artik	Shirak Plain	Sevan region	Beniamin, Vardbakh, Black Fortress I	Karchakhpyur- Shirakavan	Armenian Highland /I period/	Armenian Highland /II period/	Armenian Highland /Classical Age/	Bingel Dag
Akunk	2.24										
Artik	0.54	2.08									
Shirak Plateau /I period/	3.68	5.62	4.03								
Sevan region /II period/	2.05	4.19	2.46	1.92							
Beniamin, Vardbakh, Black Fortress I	3.97	5.62	4.04	2.30	3.14						
Karchakhpyur-Shirakavan	0.89	1.49	1.02	4.37	2.76	4.72					
Armenian Highland /I period/	2.91	5.07	3.25	1.24	1.00	2.64	3.69				
Armenian Highland /II period/	1.71	3.85	2.07	2.04	0.54	2.85	2.48	1.23			
Armenian Highland /Classical Age/	1.94	3.60	2.03	2.51	1.95	2.07	2.65	2.21	1.43		
Bingel Dag	7.85	9.40	7.64	8.32	7.65	7.68	8.53	7.39	7.58	7.73	-

changes in bones manifested in the root of the orbit (i.e. cribra orbitalia) are frequently the result of acquired or congenital anaemia(thalassemia), caused by deficiency of Mg, Cl, Fe and folic acid in nutrition, or by helminthiases and malaria. With all the above in mind, one may assume that thalassemia is a reliable indicator of the socio-economic standard as regards the state of health of the population (including the level of hygiene, nutrition and others). Brothwell (1981) referred to them as "environmental indicators". Fornaciari et al. (1981) found them to be an appropriate test of the nutrition standard of a population and even of social stratification. Here this trait is called "the misery factor", because it informs about both the malady and the unfavorable environment bringing about this sickness (Bergman, 1987). In this work it was taken for granted that individuals in which no cribra orbitalia were found, lived in "on average better conditions" (as regards nutrition, hygiene and the state of health), in contrast to those with cribra orbitalia living in "on average worse conditions", following the assumed criteria. In contrast to grave equipment, cribra orbitalia provide a biological, organismdependent source of information about living conditions. In the Bronze Age (period I) and Modern era, higher frequencies of expression are found for cribra orbitale. Cribra orbitale impedes the connection of both parts of the facies condylaris into one, and produces such bony protuberances as the tuberculum praecondylare and the torus palatines. Clinical and osteological research suggests that malnutrition or non-specific systemic stress is strongly correlated with the incidence of vertebral anomalies (Bergman, 1993; See et al., 2008; Khudaverdyan, 2011d). See et al. (2008) documented numerous vertebral anomalies in the offspring of vitamin A-deficient rats, including cleft neural arches, occipital vertebrae, vertebral blocks etc. In addition, over 80% of the offspring exhibited basioccipital malformation of some variety (See et al., 2008). In their analyses of non-metric traits in human crania, Bergman (1993) and Khudaverdvan (2011d) have also found a statistically significant association between cribra orbitalia and the precondylar tubercle, which they attributed to a common morbid factor. However, many researchers in the analysis include criba orbitalia as an epigenetic variation (Chesnis and Konduktorova, 1982; Hauser and De

Stefano, 1989; Movsesyan, 1990; Zupanic, 2004; Česnys and Tutkuvienė, 2007; Khudaverdyan, 2009).

Factor I (46.4%) has tuberculum praecondylare (0.971) and cribra orbitale (0.911)as their strongest values. High values also correspond to the os apicis lambdae (0.725), ossicula suturae squamosae (0.702) and lambdoïdeae (0.686), with the foramen mastoideum absent (0.691) (Table 6). The positive weight (factor II, 32.2%) is given for maximum foramens mastoideum exsutural (0.947), mastoideum absent (0.636), ossa suturae lambdoïdeae (0.660) in contrast to the negative weight for os bregmaticum (-0.761), ossicula suturae squamosae (-0.618). The third component accounts for 9.9% of the intergroup variance. The negative weight gives a foramen spinosum bipertitum (-0.752).

Table 6. Elements of three initial components for 11 groups.

Trait	Ι	II	III
Cribra orbitale	0.911	-0.179	0.169
Os bregmaticum	0.276	-0.761	0.460
Ossicula suturae squamosae	0.702	-0.618	-0.054
Os apicis lambdae	0.725	-0.227	0.163
Ossa suturae lambdoïdeae	0.686	0.660	0.144
Foramen mastoideum absent	0.691	0.636	0.102
Foramen mastoideum exsutural	0.306	0.947	-0.034
Tuberculum praecondylare	0.971	-0.040	-0.137
Foramen spinosum bipertitum	0.515	-0.334	-0.752
Values	46.398	32.173	9.841

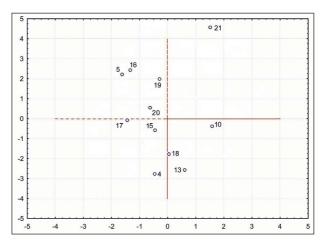


Fig. 4. Factor analysis: 4 – Artik, 5 – Armenian Highland /I period: Bronze Age/, 10 – Akunk, 13 – Lchashen, 15 – Armenian Highland /II period: Bronze and Early Iron Ages/, 16 – Shirak Plateau /I period/, 17 – Sevan region /II period/, 18 – Karchakh-pyur-Shirakavan, 19 – Beniamin, Vardbakh, Black Fortress I, 20 – Armenian Highland /Classical Age/, 21 – Bingel Dag.

Figure 4: The graph obtained using the first two axes shows how the groups from Armenian Highland (I period and Beniamin-Vardbakh-Black Fortress I) and Shirak Plateau /I period/ are close to one another. The most isolate Bingel Dag and Akunk samples. The Karchakhpyur-Shirakavan and Akunk samples exhibit close affinities to one another.

When considering the dendrogram (Fig. 5, table 7), we should note the significant similarity of the ancient group of the Karchakhpyur-Shirakavan to the samples from Bronze and Early Iron Ages (Lchashen, Akunk, Artik). The Shirak plateau sample (I period) and the samples from Classical period (Beniamin-Vardbakh-Black Fortress I) and Bronze Ages exhibit close affinities to one another. Here again, the Classical samples are closely related to the previous samples (Bronze Age). Again the most isolate Bingel Dag sample. According to the results presented here, it is possible to discern temporal trends among the Armenian Highland samples in this study. The results indicate a slight chronological trend, where populations from the Bronze to Ancient periods are more similar to one another, while the modern population appears to be more distant. Other biodistance studies using nonmetric data arrived at results similar to those reported here (Movsesyan, 1990; Movsesyan and Kochar, 2001; Khudaverdyan, 2009).

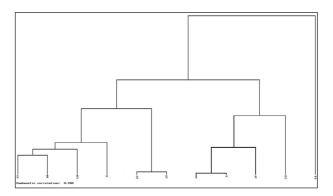


Fig. 5. Cluster tree: 4 – Artik, 5 – Armenian Highland /I period: Bronze Age/, 10 – Akunk, 13 – Lchashen, 15 – Armenian Highland /II period: Bronze and Early Iron Ages/, 16 – Shirak Plateau /I period/, 17 – Sevan region /II period/, 18 – Karchakhpyur-Shirakavan, 19 – Beniamin, Vardbakh, Black Fortress I, 20 – Armenian Highland /Classical Age/, 21 – Bingel Dag.

The next step in the analysis is to compare non-metric variation among the ancient inhabitants of the Armenian Highland with samples from the Eurasia, in order to help clarify the origins and interactions between the inhabitants of the Armenian Highland and neighboring Eurasia. An analysis of more than 14 groups of the Bronze and Iron Ages is undertaken here. The anthropological cover of Eurasia was generated during exclusively difficult historical events (Abdushelishvili, 1982, 2003; Khudaverdyan, 2011a, b). The advancement of the Mediterraneans in the territory of Eastern European was accompanied by not only an interaction of various cultural elements, but also by a mixture - a distribution sometimes at considerable distances from their centre of formation. On the basis of the

Table 7. Matrix of distance, values for eleven Armenian Highlands mortuary samples examined in this study.

Sample name	Lchashen	Akunk	Artik	Shirak Plain	Sevan region	Beniamin, Vardbakh, Black Fortress I	Karchakhpyur- Shirakavan	Armenian Highland /I period/	Armenian Highland /II period/	Armenian Highland /Classical Age/	Bingel Dag
Akunk	1.48										
Artik	1.85	1.41									
Shirak Plateau /I period/	5.75	4.80	5.52								
Sevan region /II period/	3.78	2.43	2.98	2.94							
Beniamin, Vardbakh, Black Fortress I	4.77	4.04	4.93	1.38	2.86						
Karchakhpyur-Shirakavan	1.12	1.17	1.57	4.76	2.89	3.87					
Armenian Highland /I period/	5.50	4.47	5.21	0.48	2.53	1.41	4.53				
Armenian Highland /II period/	3.29	1.98	2.68	3.07	0.53	2.74	2.42	2.68			
Armenian Highland /Classical Age/	3.96	3.78	4.33	2.97	3.49	1.99	3.06	3.06	3.22		
Bingel Dag	5.88	5.65	7.04	6.13	6.12	5.13	5.96	5.96	5.78	5.97	-

received information, cluster analysis has shown the epigenetic condensations of groups from Eurasia and factors of relatives or, conversely, distinctions between them.

Analysis 3. Placement of the 14 samples (Armenian Highland /I period: Bronze Age and II period: Bronze and Early Iron Ages/, Shirak Plateau /I period/, Sevan region /II period/, Artik, Akunk, Lchashen (Movsesyan, 1990; Khudaverdyan, 2009), Volga region /Fatianovo and Balanovo Cultures/ (Chesnis, 1986), Central Asia /Sapallitepe, Gonur Depe/ (Khodjaiov, 1977; Nevchaloda and Kufterin, 2008). Siberia /Afanasevo, Andronovo, Karasukskaya and Tagarskaya Cultures/ (Kozintsev, 1980)) determined by the values of factors I (35.2%) and II (28.3%) (Table 8). The positive weight (factor I) given for maximum foramen mastoideum (0.892) and parietale (0.799), ossa suturae lambdoïdeae (0.877), os apicis lambdae (0.788) and canalis condylaris intermedius (0. 794). Factor II has, as their strongest values, ossa suturae coronalis (0.896), sagittalis (847), sutura metopica (0.749), os asterii (0.708) and torus palatines (0.608). Factor III accounts for the 17.99% of the intergroup discrimination. Factor III has its strongest positive value a foramen supraorbitale (0.741) and negative weight a canalis hypoglossi bipartite (-0.850).

Table 8. Elements of three initial components for 14 groups.

Trait	Ι	II	III
Sutura metopica	-0.465	0.749	-0.210
Foramen supraorbitale	-0.297	0.444	0.741
Ossa suturae coronalis	-0.316	0.896	0.044
Os asterii	0.100	0.708	0.245
Foramen parietale	0.799	0.039	0.376
Os apicis lambdae	0.788	0.202	0.069
Ossa suturae sagittalis	0.262	0.847	0.371
Ossa suturae lambdoïdeae	0.877	-0.081	0.148
Foramen mastoideum absent	0.892	-0.031	0.208
Canalis condylaris intermedius	0.794	0.023	-0.358
Canalis hypoglossi bipartite	0.097	0.429	-0.850
Torus palatinus	0.524	0.608	-0.552
Values	35.194	28.205	17.989

Inspection of the 12 nonmetric cranial traits, frequencies retained for biodistance analysis indicates that some of the trait expressions demonstrate a geographic or ethnic trend (Fig. 6). More specifically, the populations from the Volga regions and Siberia show

an ethnic trend between samples (a positive field). The Armenian Highlands samples and the Mediterranean samples from Central Asia exhibit close affinities to one another (a negative field).

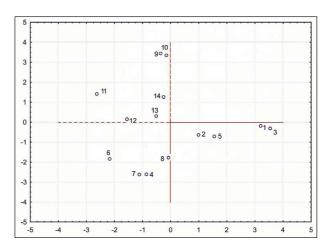


Fig. 6. Factor analysis: 1 – Armenian Highlands / period I: Bronze Age/, 2 – Armenian Highland / period II: Bronze and Early Iron Ages/, 3 – Shirak Plateau / period I /, 4 – Armenian Highland /Artik/, 5 – Sevan region / period II /, 6 – Armenian Highland /Akuk/, 7 – Armenian Highland /Lchashen/, 8 – Central Asia /Sapallitepe, Gonur Depe/, 9 – Volga region /Fatianovo Culture/, 10 – Volga region /Balanovo Culture/, 11 – Siberia /Andronovo Culture/, 12 – Siberia /Afanasevo Culture/, 13 – Siberia /Karakolskaya Culture/, 14 – Siberia /Tagarskaya Culture/.

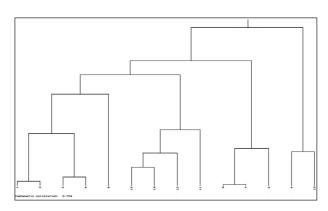


Fig. 7. Cluster tree: 1 – Armenian Highland / period I: Bronze Age/, 2 – Armenian Highland period /II: Bronze and Early Iron Ages/, 3 – Shirak Plateau / period I /, 4 – Armenian Highland /Artik/, 5 – Sevan region / period II /, 6 – Armenian Highland /Akunk/, 7 – Armenian Highland /Lchashen/, 8 – Central Asia /Sapallitepe, Gonur Depe/, 9 – Volga region /Fatianovo Culture/, 10 – Volga region /Balanovo Culture/, 11 – Siberia /Andronovo Culture/, 12 – Siberia /Afanasevo Culture/, 13 – Siberia /Karakolskaya Culture/, 14 – Siberia /Tagarskaya Culture/

The diagonal matrix of distance is provided in Table 9. The dendrogram gives a visual idea of the relationship between the various groups (Fig. 7). Cluster analysis provides a different representation of the distance matrix, because it is an unrooted tree whose branches have different lengths. Long branch lengths

Sample name	Armenia	Armenia	Shirak	Artik	Sevan	Akunk	Lchashen	Sapallitepe,	Fatianovo	Balanovo	Andronovo	Afanasevo	Karakolskaya	Tagarskaya
	/I period/	/II period/	Plateau		region			Gonur Depe	Culture	Culture	Culture	Culture	Culture	Culture
Armenian Highland														
/II period: Bronze and	2.38													
Early Iron Ages/														
Shirak Plateau /I period/	0.57	2.71												
Artik	5.27	3.06	5.49											
Sevan region /II period/	1.71	0.72	2.04	3.59										
Akunk	6.26	3.92	6.48	1.71	4.59									
Lchashen	5.18	2.90	5.44	0.46	3.48	1.68								
Central Asia /Sapallitepe, Gonur Depe/	4.06	2.74	4.52	3.39	2.90	4.63	3.11							
Volga region /Fatianovo Culture/	5.98	5.13	6.15	6.66	5.48	5.98	6.43	7.15						
Volga region /Balanovo Culture/	5.29	4.62	5.57	6.55	4.92	6.16	6.26	6.28	1.59					
Siberia /AndronovoCulture/	6.33	5.01	6.86	5.91	5.43	5.87	5.46	4.37	5.65	4.45				
Siberia /Afanasevo Culture/	4.93	3.32	5.45	4.16	3.77	4.47	3.71	2.59	5.53	4.52	1.92			
Siberia /Karasukskaya Culture	e/ 3.87	2.35	4.38	3.94	2.76	4.36	3.54	2.49	4.88	3.88	2.68	1.14		
Siberia /Tagarskaya Culture	e/ 3.69	2.58	4.18	4.65	2.91	4.90	4.29	3.41	4.08	2.94	2.84	1.95	1.05	-

Table 9. Matrix of distance, values for fourteen Eurasia mortuary samples examined in this study.

may be interpreted as an indicator of a large degree of morphological separation, while short branch lengths are indicative of a small degree of morphological separation between samples. The Artik sample features close affinity with those of the Lchashen and Akunk samples. The Armenian Highland samples from Periods I and II (Bronze and Iron Ages), the Shirak Plateau and the Sevan region serve as an epigenetic link between Central Asia (Sapallitepe, Gonur Depe) samples, which feature the closest affinities to one another. Within the dendrogram are samples from the Armenian Highlands featuring the closest affinities to one another. These four prehistoric skeletal series of different periods from Siberia are also similar to other series in the same region. Within the dendrogram are samples from the Volga region featuring the closest affinities to one another (Fatianovo and Balanovo Cultures). Thus, the cluster analysis of the 12 nonmetric cranial traits of the samples in the Bronze and Iron Ages from Eurasia indicates that in some of the traits, expressions demonstrate an ethnic trend.

Analysis (4) of 10 series: Armenian Highlands /Beniamin-Vardbakh-Black Fortress I, Karchakhpyur-Shirakavan/ (Movsesyan, 1990; Khudaverdyan, 2000, 2005), Ukraine /Chernyakhov Culture: Dzuravka, Chernykhov-Romashki-Derevynnoe-Teleshovka, Gavrilovka-Voloshskoe, Koblevo-Ranjevoe-Viktorovka/ (Chesnis and Konduktorova, 1982), the Dnestr

region /Budeshti, Malaeshti/ (Chesnis and Konduktorova, 1982), Latvia /Latgali/ (Chesnis, 1986). Global processes led to cultural and genetic transformations within the Armenian Highlands and Transcaucasia. In the present study we investigated the potential effects of gene flow among the population samples of the Armenian Highland. The results of craniological analysis give a typical picture of infiltration, from the 8<sup>th</sup> century BC up to the 3<sup>rd</sup> century AD, alien to the ethnic groups of the Transcaucasia. This scenario is consistent with supporting archaeological and historical studies (Piotrovskii, 1959; Krupnov, 1960; Strabo, 1964; Ter-Martirosov, 1999). According to archeologists, the Scythian presence in the Caucasus had been permanent (Vinogradov and Dudarev, 1983: Petrenko, 1983: Il'inskava and Terenozhkin, 1983). Here we have undertaken a nonmetric traits analysis of more than 10 groups from the territory of the Armenian Highlands and Eastern Europe. The emplacement of the coordinate axis of the samples is determined by the values of factors I (38.8% of the total variability) and II (28.3% of the total variability) (table 10). Factor I has as their strongest values canalis condylaris intermedius (0.863), foramen parietale (0.826), os asterii (0.659) and ossa suturae lambdoïdeae (0.658). The highest negative value corresponds to the foramen supraorbitale (-0.802). The highest positive weights (factor II, 28.3%) are given for foramen infraorbitale accessorium (0.859) and ossa suturae lambdoïdeae

(0.630). The third component accounts for 14.7% of the intergroup variance/variability. The highest negative weight gives a canalis hypoglossi bipartite (-0.916).

Table 10. Elements of three principal components for 10 groups.

Trait	Ι	II	III
Sutura metopica	-0.047	0.871	-0.143
Foramen supraorbitale	-0.802	0.497	-0.024
Foramen infraorbitale accessorium	0.397	0.859	0.159
Os asterii	0.659	-0.033	-0.456
Foramen parietale	0.826	0.006	0.012
Ossa suturae lambdoïdeae	0.658	0.630	0.061
Canalis condylaris intermedius	0.863	-0.343	0.266
Canalis bypoglossi bipartite	-0.074	0.042	0.916
Values	38.788	28.276	14.611

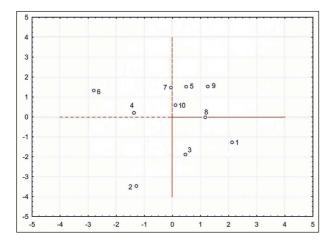


Fig. 8. Factor analysis: Armenian Highland /Beniamin-Vardbakh-Black Fortress I/, 2 – Armenian Highland /Karchakhpyur-Shirakavan/, 3 – Armenian Highland /Classical Age/, 4 - Ukraine /Dzuravka/, 5 – Ukraine /Chernykhov, Romashki, Derevynnoe, Teleshovka/, 6 – Ukraine /Gavrilovka, Voloshskoe/, 7 – Ukraine /Koblevo, Ranjevoe, Viktorovka/, 8 – the Dnestr region /Budeshti/, 9 – the Dnestr region /Malaeshti/, 10 – Latvia /Latgali/

The populations from Ukraine, the Dnestr region (Chernyakhov Culture) and Latvia (Latgali) show close affinities with one another. The Armenian Highland samples also exhibit close affinities with one another. Groups Karchakhpyur-Shirakavan are well distinguished from the groups from the Armenian Highland.

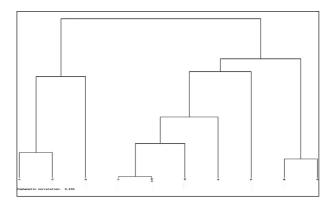


Fig. 9. Cluster tree: 1 – Armenian Highland /Beniamin-Vardbakh-Black Fortress I/, 2 – Armenian Highland /Karchakhpyur-Shirakavan/, 3 – Armenian Highland /Classical Age/, 4 – Ukraine /Dzuravka/, 5 – Ukraine /Chernykhov, Romashki, Derevynnoe, Teleshovka/, 6 – Ukraine /Gavrilovka, Voloshskoe/, 7 – Ukraine /Koblevo, Ranjevoe, Viktorovka/, 8 – Dnestr region /Budeshti/, 9 – Dnestr region /Malaeshti/, 10 – Latvia /Latgali/

The diagonal matrix is provided in Table 11. The dendrogram gives a visual idea of the relationship between the various groups (Fig. 5). Interestingly, the total sample from Beniamin-Vardbakh-Black Fortress I are the most similar to the Karchakhpyur-Shirakavan sample. When compared to other samples examined by this study (Dnestr region, Ukraine /Chernyakhov Culture/, Latvia) samples from Armenian Highland are the least similar. Given the small biological distances between the Ancient period samples from the Armenian Highland, the biological distances are most likely due to genetic drift and non-significant gene flow.

### CONCLUSION

The biodistance results reported in this study indicate that differences among prehistoric samples from the Armenian Highland are nonsignificant. Instead, when relying on nonsignificant biodistance results, it is suggested that an ancestral-descendant relationship existed among Armenian Highland populations from the Bronze Age through the Classical period. While it is recognized (craniometric studies and dental traits) that significantly different immigrant populations in the Classical period may have been present in the prehistoric Armenian Highland, they were not detected among the samples analyzed by this study. These conclusions are consistent with those reported by other biodistance studies that examined non-metric cranial analyses for Armenian Highland samples. Further,

1	niamin-Vardbakh Black Fortress I	Karchakhpyur- Shirakavan	Armenian Highland /Classical Age/	Dzuravka	Chernykhov, Romashki, Derevynnoe, Teleshovka	Gavrilovka, Voloshskoe	Koblevo, Ranjevoe, Viktorovka	Budeshti	Malaeshti	Latgali
Karchakhpyur- Shirakavan	4.21									
Armenian Highland /Classical Age/	1.82	2.45								
Dzuravka	3.78	3.75	3.03							
Chernykhov, Romash Derevynnoe, Teleshov	ki, 1.42 ka	5.96	4.15	2.94						
Gavrilovka, Voloshsk	oe 5.68	5.06	4.85	2.04	4.23					
Koblevo, Ranjevoe, Viktorovka	3.83	5.27	3.77	2.32	2.73	2.87				
Budeshti	3.10	4.62	2.91	3.58	4.24	4.57	2.37			
Malaeshti	3.49	5.75	3.85	3.50	3.21	4.28	1.45	1.69		
Latgali	2.83	4.57	2.88	1.68	1.88	3.14	1.34	2.64	2.09	-

Table 11. Matrix of distance, values for ten Eurasia mortuary samples examined in this study.

based on the biodistance results presented here, we suggest that at the beginning of the Bronze period there appears to have been a degree of genetic flow among inland populations. The biodistances reported here suggest that there was a decrease in isolation (i.e., increased gene flow) among the Classical populations during 1st century BC - 3rd century AD. This assertion requires further exploration. In spite of this possibility, it is clear that the techniques employed in this study would have made it more likely to find significant differences among the samples, if any existed. In conclusion, the biodistances from the non-metric cranial traits reported here indicate that no significant prehistoric gene flow occurred in the Armenian Highlands.

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