Genetic polymorphism of human α₂HS-Glycoprotein (AHSG) in the resident population of the Basque Country (Northern Spain)

Oscar García¹ and Antonio Alonso²

¹Sección de Biología, Laboratorio U.T. A.P., Departamento de Interior, Gobierno Vasco, c/Maria Diaz de Haro, 3-2, E-48013 Bilbao, Spain

²Sección de Biología, Instituto Nacional de Toxicología, Madrid, Spain

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Summary. The genetic polymorphism of human α_2 HS-glycoprotein (AHSG) was studied in a sample of 466 healthy unrelated individuals resident in the Basque Country (Northern Spain) by isoelectric focusing on micro-ultrathin polyacrylamide gels followed by immunoblotting. The allele frequencies obtained were AHSG*1= 0.7253, AHSG*2 = 0.2683 and AHSG*3 = 0.0064. These allele frequencies were compared with those reported in other European populations.

Key words: AHSG – Genetic polymorphism – Isoelectric focusing – Population study – European distribution

Zusammenfassung. Der genetische Polymorphismus des menschlichen α_2 HS-Glycoprotein (AHSG) wurde in einer Stichprobe von 466 gesunden unverwandten Personen untersucht, welche im Baskenland (Nordspanien) wohnen. Die Untersuchung erfolgte mit Hilfe der isoelektrischen Fokussierung auf mikro-ultradünnen Polyacrylamid-Gelen mit nachfolgendem Immunoblotting. Die Allelfrequenzen lauten: AHSF*1 = 0.7253, AHSG*2 = 0.2683 und AHSG*3 = 0.0064. Diese Allelfrequenzen wurden mit jenen aus anderen europäischen Populationen verglichen.

Schlüsselwörter: AHSG – Genetischer Polymorphismus – Isoelektrische Fokussierung – Populationsstichprobe – Europäische Verteilung

Introduction

Although the genetic polymorphism of AHSG has been investigated in many populations, the Basque population has not yet been considered. This study describes the polymorphism of AHSG in the resident population of the Basque Country (Northern Spain). This could be of importance for forensic purposes as well as for subpopulation studies.

Materials and methods

Serum samples were obtained from 466 unrelated donors resident in the Basque Country (Northern Spain). Isoelectric focusing was carried out on miniaturized polyacrylamide gels (inter-electrode distance 55 mm) as previously described (Alonso 1988). Native samples were analyzed according to Yuasa and Umetsu (1988). The AHSG band pattern was detected by a two-step enzyme immunoassay as previously described for the detection of GC protein (Alonso 1988) using the following antibodies: rabbit anti-human AHSG followed by swine anti-rabbit immunoglobulins/HRP.

The diversity between observed and expected results has been tested using the χ^2 test for independence and genetic equilibrium according to the Hardy-Weinberg law.

 Table 1. Allele frequencies of AHSG in Biscay, Guipúzcoa and Alava (Basque Country, Nothern Spain)

Population	n	AHSG allele frequencies			
		AHSG*1	AHSG*2	AHSG*3	
Biscay 256		0.7207	0.2715	0.0078	
Guipúzcoa	147	0.7279	0.2687	0.0034	
Alava	63	0.7381	0.2540	0.0079	

Table 2. AHSG phenotypes and gene distribution in the Basque Country (Northern Spain)

Phenotypes	n	Allele frequencies
1-1	249	
2-1	173	
2-2	38	AHSG*1 = 0.7253
3-1	5	$AHSG^{*2} = 0.2683$
3-2	1	$AHSG^{*3} = 0.0064$
3-3	0	
Total	466	

 $\chi^2 = 1.385388; df = 2; 0.60 > P > 0.50$

O. García and A. Alonso: AHSG in resident Basques (Northern Spain)

Table 3. Comparison between AHSG*1 andAHSG*2 allele frequencies found in thepresent study with those reported in otherEuropean populations

Population	Num- ber	AHSG			Reference
		1	2	VAR	
Norway	52	0.60	0.40	_	Olaisen et al. (1981)
Denmark	?	0.63	0.36	0.008	Eiberg et al. (1984)
England	382	0.65	0.34	0.006	Westwood et al. (1987)
Germany					
Münster	168	0.62	0.38	-	Yuasa and Umetsu (1988)
Bavaria	344	0.66	0.32	0.015	Weidinger (1986)
Region of Ulm	368	0.69	0.30	0.009	Lattke and Schönberger (1988)
France	240	0.72	0.27	0.008	Robinet-Lévy et al. (1988)
Italy	199	0.76	0.24		Scacchi et al. (1989)
Spain					
Madrid	489	0.71	0.28	0.008	Alonso et al. (1991)
Basque Country	466	0.72	0.27	0.006	This study

Results and discussion

The allele frequencies of AHSG in the Basque resident population are shown in Table 1. The allele and phenotype frequencies in the total sample from the Basque resident population are shown in Table 2. The observed numbers are in agreement with the expectation according to the Hardy-Weinberg law.

The AHSG*1 and AHSG*2 allele frequencies determined in this study have been compared with those reported in other European populations (Table 3). The values of AHSG*1 and AHSG*2 allele frequencies are within the range of variation reported in the resident population of Madrid (Central Spain), which is the only Spanish population previously studied (Alonso et al. 1991), France (Robinet-Lévy et al. 1988) and Italy (Scacchi et al. 1989). However, the comparison with Northern European populations such as Norway (Olaisen et al. 1981), England (Westwood et al. 1987) and Germany (Yuasa and Umetsu 1988; Weidinger 1986; Lattke and Schönberger 1988) showed significant differences that should be attributed to a progressive increase of the AHSG*2 allele frequencies and the consequent decrease of the AHSG*1 allele frequencies towards the North.

In conclusion, the AHSG population data from the resident population of the Basque Country is in the range of variations of other Southern European populations and is in agreement with the suggested north-to-south AHSG genocline (Yuasa and Umetsu 1988). The AHSG*3 variant has also been confirmed in this population study.

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