

Genetic polymorphism of Alpha-2-HS glycoprotein in Lombardy (Italy)

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Received February 27, 1990 / Received in revised form May 5, 1990

Summary. α 2HS-subtypes have been analysed in samples from 700 unrelated individuals from the Brescia area (Italy) by the isoelectric focusing technique and immunofixation. The observed allele frequencies were: α 2HS*1 = 0.7472; α 2HS*2 = 0.2507; α 2HS*V = 0.0021.

Key words: Blood groups – α 2HS – Glycoprotein polymorphism

Zusammenfassung: Bei 700 nicht verwandten Individuen aus Brescia (Italien) wurden die α 2HS Phänotypen mittels isoelektrischer Fokussierung und anschließender Immunofixation bestimmt. Die beobachteten Allelfrequenzen waren: α 2HS*1 = 0.7472; α 2HS*2 = 0.2507; α 2HS*V = 0.0021.

Schlüsselwörter: Blutgruppen – α 2HS – Glykoproteinpolymorphismus

Introduction

Alpha-2-HS-glycoprotein (α 2HS, A2HS) is a plasma glycoprotein (mol. wt. 49000), present in the serum of healthy individuals at levels between 45 and 80 mg/dl (Putnam 1984). It was first described by Heremans (1960) and later isolated by Schmidt and Burgi (1961). Alpha-2-HS glycoprotein is synthesized in the liver and deposited in mineralising human bone (Dikson et al. 1975; Triffitt et al. 1976). Its biological function is not yet entirely understood: in vitro, it enhances the phagocytic function of mouse macrophages (Lewis and André 1980) and human monocytes (Lewis and André 1981); it binds to lymphocytes which have been transformed by the Epstein Barr virus and this property can be of importance to remove these cells from circulation (Lewis and André 1982, 1983). A significant decrease is observed in patients suf-

fering from an acute inflammatory process of bacterial etiology (Lebreton et al. 1979).

The genetic polymorphism of this protein was first investigated by Anderson and Anderson in 1977 who used 2-dimensional electrophoresis. They described 2 different electrophoretic variants L and N. The presence of 2 common genetic types of α 2HS (A2HS*1 = L and A2HS*2 = N) has been confirmed by Umetsu et al. (1983) by means of the isoelectric focusing technique. Subsequent IEF methods have been developed with the detection of several variants which are polymorphic in some geographic areas. Cox and Andrews (1983) reported the existence of three alleles (α 2HS*1, 2 and 3) in a caucasian population, in 1984 Weidinger et al. described a new variant defined as α 2HS*4, while Umetsu et al. (1985) observed the A2HS*5 allele in a Japanese population. At present it is possible to distinguish a total of 15 alleles (Yuasa and Umetsu 1988). In 1985 Cox and Francke showed that the A2HSG locus is located on chromosome 3.

So far, genetic polymorphism of A2HS has not been extensively investigated and is not commonly used in paternity testing or bloodstain analysis. The aim of this study is to present a contribution to a better knowledge of the distribution of α 2HS polymorphism in Italy through a population sample from the Brescia area (Lombardy).

Materials and methods

Sera from 700 healthy and unrelated blood donors, made available by the Transfusion Centre of the "Spedali Civili" of Brescia (Lombardy – Italy) have been studied. Samples were stored at -20°C before use and tested within one month.

α 2HS typing was performed by thin layer isoelectric focusing on polyacrylamide gel ($250 \times 120 \times 0.5\text{ mm}$), according to Weidinger (1986), with minor modifications as suggested by Giari et al. (1988).

The gel was prepared by mixing the following solutions: 9 ml acrylamide stock solution (19% w/v acrylamide and 0.6% w/v N-N' Methylene bisacrylamide), 2.4 ml glycerol (87%), 5.5 ml distilled water, 1 ml Pharmalyte pH 4.2–4.9, 0.2 ml Pharmalyte pH 4.5–5.4 and 30 mg ACES (Serva).

Polymerization was performed after the addition of 6 μl TEMED and 180 μl ammonium persulfate (3%). After 1 h at room tempera-

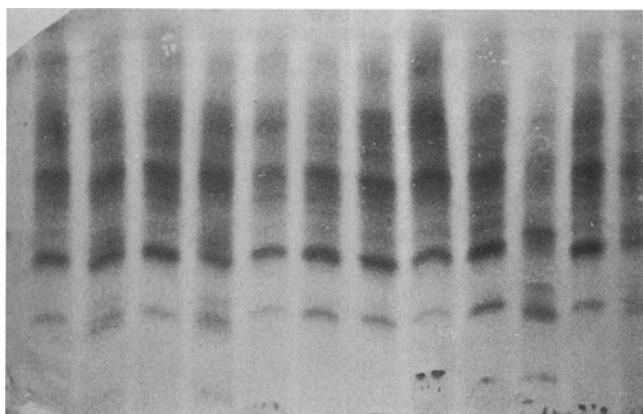


Fig. 1. Electrofocusing and immunofixation pattern of the common α 2HS phenotypes. From left to right: (1) 1; (2) 2-1; (3) 1; (4) 2-1; (5-9) 1; (10) 2; (11) 1; (12) 2-1

Table 1. α 2HS phenotype distribution and allele frequencies in Brescia (Lombardy - Italy)

Pheno-type	Observed		Expected		Allele frequencies
	n	%	n	%	
1	395	56.43	390.82	55.83	α 2HS*1 = 0.7472
2-1	254	36.28	262.25	37.47	α 2HS*2 = 0.2507
2	48	6.86	44.00	6.29	α 2HS*V = 0.0021
1-V	2	0.29	2.20	0.31	
2-V	1	0.14	0.73	0.10	
Total	700	100.00	700.00	100.00	

$$\chi^2 = 0.7857, 2 \text{ d.f.}; 0.5 > P > 0.7$$

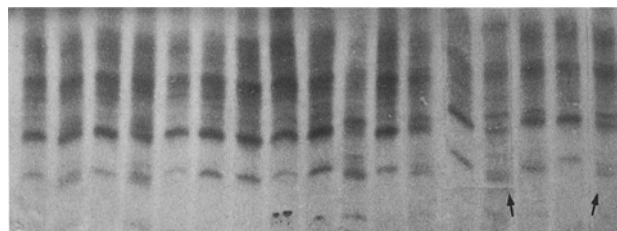


Fig. 2. α 2HS 2-V and 1-V. From left to right: (1) 1; (2) 2-1; (3) 1; (4) 2-1; (5-9) 1; (10) 2-2; (11) 1; (12) 2-1; (13) 1; (14) 2-V; (15) 1; (16) 1-V

Table 2. α 2HS allele frequencies in Italy

Region	n	α 2HS*1	α 2HS*2	α 2HS*V	Reference
Tuscany	350	0.7214	0.2771	0.0014	Giari et al. 1988
Latium	199	0.756	0.244	-	Scacchi et al. 1989
Abrutium	106	0.788	0.212	-	Scacchi et al. 1989
Sardinia	152	0.832	0.168	-	Scacchi et al. 1989
Lombardy	700	0.7472	0.2507	0.0021	This study

ture the gel was stored overnight at 4°C. Serum samples (8 μ l) were applied 1.5 cm from the cathode using applicator strips from Serva. As electrode solutions, a mixture of 0.025 M aspartic acid and 0.025 M glutamic acid was used for the anode and 0.1 M NaOH for the cathode.

Table 3. Geographical distribution of α 2HS allele frequencies

Country	n'	α 2HS*1	α 2HS*2	α 2HS*V	Reference
Norway	52	0.6000	0.3900	-	Olaison et al. 1981
England	382	0.6466	0.3469	0.0069	Westwood 1988
Germany	166	0.654	0.3220	0.0240	Weidinger et al. 1984
Germany	197	0.6550	0.3400	0.0050	Tarkkala Mender and Kühnl 1986
Germany	344	0.6642	0.3208	0.0150	Weidinger 1986
Germany	249	0.6310	0.3630	0.0060	Luckenbach et al. 1988
Germany	168	0.6250	0.3750	-	Yuasa and Umetsu 1988
France	240	0.7167	0.2750	0.0083	Robinet-Levy et al. 1988
Egypt	95	0.8579	0.1424	-	Abe et al. 1987
Libya	110	0.8364	0.1636	-	Sebetan and Heshmat 1988
Canada	215	0.6419	0.3555	0.0046	Cox et al. 1986
USA	150	0.6530	0.3470	-	Boutin et al. 1985
Japan	2050	0.7356	0.2639	0.0005	Umetsu et al. 1984
Japan	300	0.7233	0.2767	-	Yuasa et al. 1985
Japan	400	0.7325	0.2675	-	Yuasa et al. 1985
Japan	397	0.7670	0.2065	0.0264	Yuasa et al. 1985
Japan	283	0.7456	0.2544	-	Tamaki et al. 1985
Japan	256	0.7637	0.2363	-	Kishi et al. 1988
Taiwan	199	0.7286	0.2714	-	Yuasa and Umetsu 1988
Philippines	115	0.6870	0.3130	-	Umetsu et al. 1988
Nepal	140	0.7571	0.2429	-	Yuasa and Umetsu 1988

IEF was performed using an LKB Ultraphor chamber connected to the LKB 2197 Power Supply for a total of 3 h: 1 h at 2000 V, 20 mA, 8W and 2 h at 2000 V, 20 mA, 18W. The cooling unit was set at 8°C.

After separation, α 2HS phenotypes were identified by immunofixation. A cellulose acetate strip was soaked in anti-human α -2-HS-Glycoprotein (ATAB) diluted 1:3 in saline and placed on the gel surface at room temperature for 10 min. The strip was removed and washed twice in saline for 15 min, stained with nigrosine (1%) and destained in acetic acid (5%).

Results and discussion

Table 1 shows the phenotypic distribution and allele frequency values of α 2HS obtained in the population sample studied from Brescia.

Three common phenotypes α 2HS 1-1, α 2HS 2-1, α 2HS 2-2 and two rare variants were observed. These variants, defined as 1-V and 2-V, have a migration similar to the product of the allele AHSG*4, first described by Weidinger et al. in 1984.

Good correlation was found between the observed and expected phenotype distributions, assuming Hardy-Weinberg equilibrium conditions.

No significant differences have been found between the allele frequencies calculated in our population sample and other samples from Continental Italy (Table 2).

Table 3 gives the population data obtained by various authors showing the obvious variation in ethnic and geographic distribution. The estimated frequency of $\alpha 2\text{HS}^*1$ allele in our sample and in Continental Italy is higher than the frequency obtained in northern Europe and northern America but lower than that found in North Africa; it seems closer to the values recorded in Japanese.

On the basis of the allele frequencies estimated in the population sample from Brescia, the cumulative theoretical "a priori" probability of exclusion for $\alpha 2\text{HS}$ is 15.22% (the I class theoretical "a priori" probability of exclusion is 8.2%).

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