

Classification of α_2 -HS-Glycoprotein (α_2 HS) Types by Isoelectric Focusing

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Summary. A sample of 300 sera from unrelated individuals from Northern Japan was examined by isoelectric focusing on polyacrylamide gels. Three common types, α_2 HS 1-1, 2-1, and 2-2 were differentiated. The frequencies of the α_2 HS alleles in our sample were found to be: α_2 HS¹=0.7250 and α_2 HS²=0.2750. Analysis of 16 parents with 21 children did not show deviations from the expected mode of inheritance.

Key words: Serum groups, α_2 -HS-glycoprotein (α_2 HS) polymorphism – Paternity examinations, α_3 HS

Zusammenfassung. Mit Hilfe der isoelektrischen Fokussierung in Polyacrylamidgelen wurden 300 Proben von nicht verwandten Personen aus Nordjapan untersucht. Es wurden drei häufige Gruppen, α_2 HS 1-1, 2-1 und 2-2 differenziert. Die Allelfrequenzen in dieser Stichprobe betrugen: α_2 HS¹= 0,7250 und α_2 HS²=0,2750. Die Untersuchung von 16 Elternpaaren mit ihren 21 Kindern ergab keine Abweichung vom angenommenen autosomal kodominanten Erbgang.

Schlüsselwörter: Serumgruppen, α_2 -HS-Glykoprotein-Polymorphismus - Vaterschaftsuntersuchung, α_2 HS

Introduction

Schultze et al. (1962) isolated a new kind of glycoprotein from serum and named α_2 -HS-glycoprotein (α_2 HS). α_2 HS seems to have an essential function in the opsonic effect of human leucocytes (van Oss et al. 1974) but information of its role in vivo is lacking. Naturally, we have never seen a report related to α_2 HS polymorphism.

Using the technique of isoelectric focusing in polyacrylamide gel, we have examined the α_2 HS in a large number of sera from unrelated individuals and from members of normal families. The results of this survey are the subject of this report.

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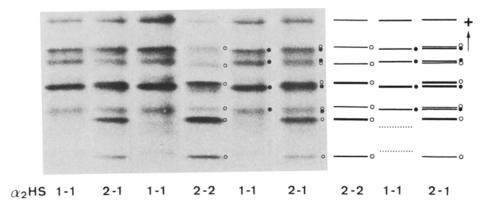


Fig. 1. Electrofocusing band patterns of the common α_2 HS phenotypes. Filled circles indicate α_2 HS¹-bands and empty circles α_2 HS²-bands

Materials and Methods

The sera were obtained from unrelated healthy individuals (n=300) and 16 families with 21 children residing in Yamagata prefecture, a northern area of Japan.

The Servalyt-containing polyacrylamide gels $(230 \times 110 \times 0.5 \text{ mm})$ were prepared by mixing 2.8 ml acrylamide solution (29.1 g/100 ml), 2.8 ml N,N'-methylene bisacrylamide (0.9 g/100 ml), 1.0 ml Servalyt 4.5-5.0 (Serva), 9.8 ml distilled water, and 0.4 ml ammonium persulfate (10 mg/ml). Six-millimeter electrode strips of a thick cellulose paper (Serva) were placed at both ends of the gel. The strips were soaked in the following electrode solutions: 0.025 M aspartic acid and 0.025 M glutamic acid at the anode, and 2 M ethylenediamine, containing 0.025 M arginine and 0.025 M lysine, at the cathode. Isoelectric focusing was carried out at 6° C with 7 W. We allowed it to prefocus (without samples) during first 30 min. Then 3 µl of the samples were applied with filter paper (Whatman No. 3) at a distance of 1.5 cm from the cathode. After 30 min, the paper pieces were removed, and focusing was terminated after a total time of 4 h.

The immunofixation was carried out using the monospecific anti- α_2 -HS-glycoprotein (Behring). A strip of cellulose acetate membrane was soaked with the antiserum and placed on the α_2 HS region of the gel for 10 min. Subsequently, the strip was removed, washed 3 h with saline, and stained with Acid Violet 49 (Serva).

Results and Discussion

Figure 1 shows the band patterns of three common human α_2 HS-polymorphism phenotypes electrofocused in pH 4.5–5.0 gradient. The homozygous α_2 HS 1-1 and α_2 HS 2-2 consist of four or six α_2 HS¹- and six α_2 HS²-bands, respectively. While the heterozygous phenotype α_2 HS 2-1 exhibits composite patterns of α_2 HS¹- and α_2 HS²-bands.

Distribution of the observed phenotypes is given in Table 1. The estimated allele frequencies of our population sample were: $\alpha_2 \text{HS}^1 = 0.7250$ and $\alpha_2 \text{HS}^2 = 0.2750$. A good agreement was found between the observed and expected phenotype distributions assuming Hardy-Weinberg equilibrium condition ($\chi^2 = 0.24$, df = 1, 0.5 < P < 0.7).

Table 2 shows the results of family studies. The results are in agreement with an autosomal codominant mode of inheritance.

Table 1. α ₂ HS	phenotypes	and allele	frequencies	in the	Japanese population
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Phenotypes	Phenotypes		Allele frequencies		
	observed	expected			
α ₂ HS 1-1	156	157.7	$\alpha_2 HS^1 = 0.7250$		
2-1	123	119.6	$\alpha_2 HS^2 = 0.2750$		
2-2	21	22.7			
Total	300	300.0			

Table 2. Distribution of α_2 HS phenotypes in 16 families with 21 children

Phenotypes of parents	Number of mating	Number of children	Children			
			1-1	2-1	2-2	
1-1×1-1	2	3	3 (3)	_	_	
$1-1 \times 2-1$	10	14	8 (7)	6 (7)	_	
$1\text{-}1\times2\text{-}2$	1	1	_	1 (1)	_	
$2-1 \times 2-1$	3	3	2 (0.75)	1 (1.5)	0 (0.75)	
Total	16	21	13	8	0	

Expected values are given in parentheses

From these observations the α_2 HS types are determined by two alleles, α_2 HS¹, and α_2 HS². As a result of the favorable distribution of α_2 HS genes in the Japanese population, this system is of high value for paternity testing. We calculated the isolated theoretic exclusion rate for this system to be 16.0%.

References

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