

Originalarbeiten / Original Works

Pi Subtyping by Isoelectric Focusing: Further Genetic Studies and Application to Paternity Examinations

S. Weidinger, F. Schwarzfischer, and H. Cleve

Institute for Anthropology and Human Genetics, University of Munich, Richard-Wagner-Str. 10/I, D-8000 Munich 2, Federal Republic of Germany

Summary. Genetic variation of the protease inhibitor (Pi) a_i -antitrypsin was analyzed by isoelectric focusing on polyacrylamide gels in a sample of 347 unrelated individuals from Southern Germany. Six common subtypes of Pi M were observed as well as the relatively frequent variants Pi S and Pi Z and the rare variants Pi T, Pi < L, Pi L, Pi I and Pi F. Also, a variant called Pi Z1 was found. The frequency of alleles in this sample was Pi MI = 0.6917, Pi M2 = 0.1686, Pi M3 = 0.0865, Pi S = 0.0230, Pi Z = 0.0187, and Pi = 0.0115. In 82 families the distribution of Pi types was in agreement with an autosomal codominant mode of inheritance. The application of Pi classification in cases of disputed paternity is discussed.

Key words: Serum groups, a_1 -antitrypsin – Pi-subtypes, isoelectric focusing – Paternity examinations, Pi-subtypes

Zusammenfassung. Die genetischen Variationen des Protease-Inhibitors (Pi) a_1 -Antitrypsin wurden mit Hilfe der Isoelektrofokussierung in einer Stichprobe von 347 nicht verwandten Personen aus Süddeutschland untersucht. Es wurden sechs häufige PiM-Untergruppen und die relativ häufigen Varianten PiS und PiZ differenziert; zudem fanden sich die seltenen Varianten PiT, Pi<L, PiL, PiI, PiF sowie eine als PiZ1 bezeichnete Variante. In dieser Stichprobe wurden folgende Allelfrequenzen berechnet: Pi^{MI} = 0,6917, Pi^{M2} = 0,1686, Pi^{M3} = 0,0865, Pi^S = 0,0230, Pi^Z = 0,0187 und Pi* = 0,0115. In 82 Familien fand sich keine Abweichung vom angenommenen autosomal kodominanten Vererbungsmodus. Die Verwendbarkeit des Pi-Systems für die Paternitätsbegutachtung wird diskutiert.

Schlüsselwörter: Pi-Untergruppen, Isoelektrofokussierung – Vaterschaftsbegutachtung, Pi-Untergruppen – Blutgruppen, a_1 -Antitrypsin

Offprint requests to: S. Weidinger (address see above)

2 S. Weidinger et al.

Alpha₁-antitrypsin is the most prominent protease inhibitor (Pi) of human plasma. At least 25 different inherited Pi variants have been demonstrated by various electrophoretic procedures (Fagerhol and Laurell 1970; Cox and Celhoffer 1974; Johnson 1976). Application of the method of isoelectric focusing on polyacrylamide gels (PAGIF) revealed further genetic heterogeneity (Allen et al. 1974; Arnaud et al. 1975; Kueppers 1976; Genz et al. 1977; Klasen et al. 1977; Frants and Eriksson 1978; Kühnl and Spielmann 1978). By this method, six common Pi M subtypes can be distinguished which are called PiM1, PiM1M2, PiM2, PiM1M3, PiM2M3, and PiM3. Family studies indicate that these six subtypes are determined by three alleles Pi^{M1}, Pi^{M2}, and Pi^{M3} (Frants and Eriksson 1978; Kueppers and Christopherson 1978; Kühnl and Spielmann 1978; Cleve et al. 1979). Pi Z in the homozygous state is associated with severe chronic obstructive pulmonary disease (Eriksson 1965) or with childhood liver cirrhosis (Sharp 1973). Also, a rare allele Pi⁰ (null) has been described (Talamo et al. 1973).

In this study an improved separatory procedure is used. We provide data on the distribution of Pi subtypes and Pi variants in a sample of unrelated individuals from Southern Germany. Distribution of Pi types in 82 families is demonstrated. An unusual phenotypic variant of PiZ is presented. We discuss also the application of the Pi classification in cases of disputed paternity.

Material and Methods

Blood was collected from healthy individuals who were residents from Southern Germany. They were investigated in the course of paternity examinations. Sera were analyzed freshly and without prior treatment. Isoelectric focusing was carried out on thin-layer polyacrylamide gels (PAGIF). Gels $(245 \times 110 \times 1 \text{ mm})$ were prepared from 5 ml acrylamide solution (28%), 5 ml N,N'-methylenbisacrylamide solution (2%), 1 ml Servalyt pH 4-4.5, 0.5 ml Servalyt pH 4-5, 4 g saccharose, and 20 ml aqua dest. The mixture was freed from air bubbles by evacuation for 5 min. One milliliter ammonium persulphate solution (1%) was added. The mixture was filled into a prepared gel mould (LKB) with syringe. The gels polymerized within 30 min. Five microliter serum was applied on filter paper and placed on the gel 2 cm off the cathodal edge of the plate. The Multiphor chamber of LKB was used with the cooling system at a temperature of +10°C for the circulating fluid. At 40 mA and 24 W with a maximum of 1,600 V isoelectric focusing was carried out for 180 min. At the cathode 1 M glycine, at the anode 1 M H₃PO₄ was employed. Some gels were analyzed by immunofixation with monospecific α_1 -antitrypsinantisera (Atlantic Antibodies from Merz + Dade) soaked onto cellulose acetate strips and applied to the gel for 2 min immediately following the isoelectric separation. Fixation of proteins was accomplished by precipitation with sulfosalicylic acid for 30 min. Gels were stained with Coomassie Brillant Blue R 250.

Results and Discussion

In Table 1 the distribution of Pi phenotypes in 347 unrelated individuals from Southern Germany is shown. The observed distribution and the distribution expected at population equilibrium are in close agreement. The phenotype- and allele frequencies in this sample are different to the distributions reported earlier from our group only in two minor respects: In our first study (Genz et al. 1977) a relative deficit of the subtype M1M3 was observed which was not found in this

| Phenotypes | Observe | d | Expected | Allele | |
|------------|---------|-------|----------|--------|--------------------|
| | п | % | n | % | frequencies |
| Pi M1 | 171 | 49.28 | 166.02 | 47.84 | $Pi^{M1} = 0.6917$ |
| M1M2 | 79 | 22.77 | 80.93 | 23.32 | $Pi^{M2} = 0.1686$ |
| M1M3 | 35 | 10.09 | 41.52 | 11.97 | $Pi^{M3} = 0.0865$ |
| M2 | 9 | 2.59 | 9.86 | 2.84 | $Pi^{S} = 0.0230$ |
| M2M3 | 13 | 3.75 | 10.12 | 2.92 | $Pi^{Z} = 0.0187$ |
| M3 | 3 | 0.86 | 2.60 | 0.75 | Pi* = 0.0115 |
| M1S | 9 | 2.59 | 11.04 | 3.18 | |
| M2S | 4 | 1.15 | 2.69 | 0.78 | |
| M3S | 3 | 0.86 | 1.38 | 0.40 | |
| M1Z | 10 | 2.88 | 8.98 | 2.59 | |
| M2Z | 1 | 0.29 | 2.19 | 0.63 | |
| M3Z | 2 | 0.58 | 1.12 | 0.32 | |
| M1Z1 | 2 | | | | |
| MIT | 1 | | | | |
| M2 < L | 1 | | | | |
| M1L | 1. | 2.31 | 8.55 | 2.46 | |
| MII | 1 | | | | |
| M2I | 1 | | | | |

Table 1. Distribution of Pi phenotypes and Pi alleles in a sample from Southern Germany

 $\Sigma \chi^2 = 2.3037$, df = 5, P > 0.20; Pi* = Frequency of rare Pi alleles

100.00

1

347

M3F

Total

Note: Phenotypes of Pi^S. Pi^Z and Pi* were combined for χ^2 calculation. Pi Z1 has recently been found in a child and his presumptive father. Further genetic studies indicate the existence of further suballeles of Pi^M which are called Pi^{M4} and Pi^{M5}

347.00

100.00

sample presumably due to more complete ascertainment of this class as a consequence of the improved separatory procedure. In our second study (Cleve et al. 1979) the frequency for Pi^Z was unexpectedly low, probably caused by an artefact, viz., the destruction of Pi Z in older serum specimen as a consequence of longer storage and repeated freezing and thawing. The distribution found in this study and given in Table 1 is in close agreement with the findings of Kühnl and Spielmann (1978) in a sample from Hessen/FRG. Apparently, there are no significant regional differences in the German population as far as the PiM subtypes are concerned. The subtype distribution is also similar in the Dutch population (Klasen et al. 1977; Frants and Eriksson 1978). Other European populations or populations of European origin may have slightly different distributions (Frants and Eriksson 1978; Kueppers and Christopherson 1978).

In Table 2 the results of a family study are summarized. In 82 families with a total of 82 children the distribution of Pi M and Pi phenotypes corresponded to an autosomal codominant mode of inheritance. Including this study, 273 families with a total of 397 children have been analyzed for the inheritance of Pi M subtypes and Pi types at the present time. The results confirm fully the genetic hypothesis.

S. Weidinger et al.

Table 2. Distribution of PiM subtypes in 82 parents with a total of 82 children

| Parents | n | Children | | | | | | | | | | |
|-------------------|------|----------|------|------|----|------|----|-----|-----|-----|-----|----|
| | | M1 | M1M2 | M1M3 | M2 | M2M3 | М3 | M1S | M2S | M1Z | M3I | SZ |
| $M1 \times M1$ | 17 | 17 | | | | | | | | | | |
| $M1 \times M2$ | 2 | | 2 | | | | | | | | | |
| $M1 \times M3$ | 1 | | | 1 | | | | | | | | |
| $M1 \times M1M$ | 2 11 | 5 | 6 | | | | | | | | | |
| $M1 \times M1M$ | 3 8 | 4 | | 4 | | | | | | | | |
| $M1 \times M2M$ | 3 5 | | 2 | 3 | | | | | | | | |
| $M1M2 \times M2$ | 3 | | 1 | | 2 | | | | | | | |
| $M1M2 \times M1M$ | 2 5 | 2 | 2 | | 1 | | | | | | | |
| $M1M2 \times M1M$ | 3 4 | 1 | 1 | 0 | | 2 | | | | | | |
| $M1M2 \times M2M$ | 3 4 | | 2 | 0 | 1 | 1 | | | | | | |
| $M1M3 \times M1M$ | 3 2 | 1 | | 1 | | | 0 | | | | | |
| $M1M3 \times M2M$ | 3 1 | | 1 | 0 | | 0 | -0 | | | • | | |
| M1 × M1S | 2 | 0 | | | | | | 2 | ٠ | | | |
| $M1 \times M2S$ | 1 | | 0 | | | | | 1 | | | | |
| $M2 \times M1S$ | 1 | | 1 | | | | | | 0 | | | |
| $M1M2 \times M1S$ | 1 | 0 | 1 | | | | | 0 | 0 | | | |
| $M1M2 \times M2S$ | 1 | | 0 | | 0 | | | 0 | 1 | | | |
| $M1M3 \times M2S$ | 1 | | 1 | | | 0 | | 0 | | | | |
| $M1M3 \times M3S$ | 2 | | | 0 | | | 1 | 1 | | | | |
| $M1 \times M1Z$ | 2 | 2 | | | | | | | | 0 | | |
| $M1 \times M2Z$ | 1 | | 0 | | | | | | | 1 | | |
| $M1M2 \times M1Z$ | 2 | 1 | 1 | | | | | | | 0 | | |
| $M1M3 \times M1Z$ | 1 | 1 | | 0 | | | | | | 0 | | |
| $M2M3 \times M1Z$ | 1 | | 0 | 1 | | | | | | | | |
| M1S ×M3Z | 1 | | | 0 | | | | | | 0 | | 1 |
| $M1 \times M1I$ | 1 | 1 | | | | | | | | | | |
| $M3S \times M2I$ | 1 | | | | | 0 | | | 0 | | 1 | |
| Total | 82 | 35 | 21 | 10 | 4 | 3 | 1 | 4 | 1 | 1 | 1 | 1 |

In two different cases of disputed paternity an unusual Pi phenotype was observed. We have named this phenotypic variant PiZ1 because of its similarity to PiZ. As shown in Figs. 2 and 3 PiZ1 is characterized by a band slightly more cathodal than PiZ and, in addition, a broad cathodically located band. These bands were observed together with PiM1. The results were reproducible on repeated examinations and, in one case, on repeat samples from the same individual. The broad extra band reacts with a specific α_1 -antitrypsin-antiserum on immunofixation (Fig. 3). A family study will be carried out. Kühnl (pers. commun. 1980) has observed a similar phenotypic variant for which he considers a nongenetic origin.

In Figs. 1-3 the Pi phenotypes are illustrated. In Fig. 1 the six common PiM subtypes are shown and the variants PiS and PiT. The arrows point to the M3 bands in the so-called M8 region, the slight differences indicate that further heterogeneity of M3 may exist. In Fig. 2 several phenotypes of the variants PiS, PiZ and PiI are shown together with PiM1Z1. In Fig. 3 the patterns obtained after

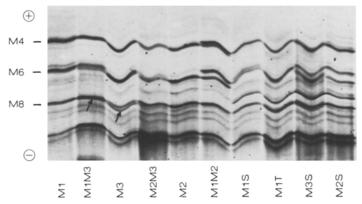


Fig. 1. Demonstration of Pi phenotypes by isoelectric focusing on polyacrylamide gels with a pH range of 4-5. Shown are the six common Pi M subtypes and the variants Pi S and Pi T. The arrows point to the Pi M3 bands (see text)

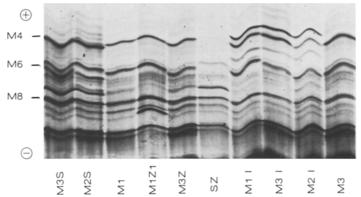


Fig. 2. Isoelectric locusing for r_1 classification; illustrated are several P_i phenotypes of the variants $S,\,Z,\,I,\,$ and $\,Z\,I$

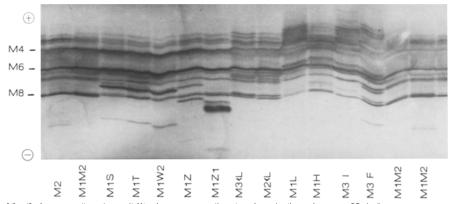


Fig. 3. Immunofixation of Pi phenotypes after isoelectric focusing at pH 4-5

6 S. Weidinger et al.

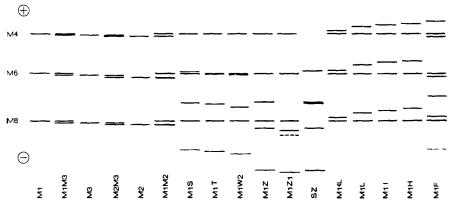


Fig. 4. Schematic presentation of Pi phenotypes as observed by polyacrylamide gel isoelectric focusing (PAGIF) at pH gradient 4-5

immunofixation are shown. The Pi banding pattern is very complex, reliable classification obviously requires a high degree of expertise.

In Fig. 4 we give a schematic presentation of the Pi phenotypes as observed by isoelectrofocusing at pH 4–5. M4, M6, and M8 refer to the different regions of the banding pattern in the pH gradient. The phenotypic notations are given at the bottom.

We applied Pi classification in 66 cases of disputed paternities. In six cases the constellation in the Pi system permitted an exclusion of paternity. This concerned two complainants, three defendants, and one witness. In each of this cases the exclusion was confirmed in other systems. From our results the practical exclusion rate appears to be 9.10%. The theoretical exclusion rate was calculated to be 23.37%. Pi classification which includes the determination of PiM subtypes has, therefore, great potential for paternity examinations. Adequate separation by isoelectric focusing is mandatory for reliable classification. Furthermore, fresh sera or sera which have been frozen and thawed only once or twice are required, since in our experience the bands of PiZ, PiS, and PiI are susceptible to repeated freezing and thawing during which treatment they gradually become fainter until they disappear.

Acknowledgement. We are grateful to Dr. P. Kühnl, Institut für Immunhämatologie, Universität Frankfurt, for control examinations and the provision of several standard sera. We thank Mrs. W. Patutschnick for her technical assistance.

References

Allen RC, Harley RA, Talamo RC (1974) New method for determination of alpha₁-antitrypsin phenotypes using isoelectric focusing on polyacrylamide gel slabs. Am J Clin Pathol 62: 732-739

Arnaud Ph, Chapuis-Cellier C, Creyssel R (1975) The Pi system: its study by means of thin-layer-gel electrofocusing in polyacrylamide gel. In: Peters H (ed) 22nd Colloquium, protides of the biological fluids. Pergamon, Oxford, pp 515-520

- Cleve H, Patutschnick W, Strecker K, Nevo S (1979) Inheritance of PiM subtypes. A study of 151 families with a total of 242 children and of 142 mother-child pairs. Hum Hered 29: 351-354
- Cox DW, Celhoffer L (1974) Inherited variants of α_1 -antitrypsin: a new allele Pi^N. Canad J Genet Cytol 16:297–303
- Eriksson S (1965) Studies in alpha₁-antitrypsin deficiency. Acta Med Scand (Suppl) 432:1-85
- Fagerhol MK, Laurell CB (1970) The Pi system—inherited variants of serum a_1 -antitrypsin. In: Steinberg AG, Bearn AG (eds) Progress in medical genetics, vol 7. Grune and Stratton, New York, pp 96-111
- Frants RR, Eriksson AW (1978) Reliable classification of six PiM subtypes by separator isoelectric focusing. Hum Hered 28:201-209
- Genz T, Martin JP, Cleve H (1977) Classification of a_1 -antitrypsin (Pi) phenotypes by isoelectrofocusing. Distribution of six subtypes of the PiM phenotype. Hum Genet 38:325-332
- Johnson AM (1976) Genetic typing of α_1 -antitrypsin by immunofixation electrophoresis. Identification of subtypes of PiM. J Lab Clin Med 87:152–163
- Klasen EC, Franken C, Volkers WS, Bernini LF (1977) Population genetics of a_1 -antitrypsin in the Netherlands. Description of a new electrophoretic variant. Hum Genet 37:303–313
- Kühnl P, Spielmann W (1978) Subtypisierung der genetisch determinierten Serumproteinpolymorphismen des Transferrins (Tf), der gruppenspezifischen Komponente (Gc) und des a₁-Antitrypsins (Pi). Archiv Genetik 51:15-16
- Kühnl P, Spielmann W (1979) Pi^T: a new allele in the alpha₁-antitrypsin system. Hum Genet 50: 221-223
- Kueppers F (1976) a₁-antitrypsin M₁: a new common genetically determined variant. Am J Hum Genet 28:370-377
- Kueppers F, Christopherson MJ (1978) Alpha₁-antitrypsin: further genetic heterogeneity revealed by isoelectric focusing. Am J Hum Genet 30:359-365
- Sharp HL (1973) Alpha₁-antitrypsin deficiency. In: Mc Kusick VA, Claiborne R (eds) Med Genet. Hospital Practice Publishing Company, New York, pp 131–140
- Talamo RC, Langley CE, Reed CE, Makino S (1973) α_1 -Antitrypsin deficiency: A variant with no detectable α_1 -antitrypsin. Science 181:70–71
- Weidinger S, Schwarzfischer F, Cleve H (1980) Classification of transferrin (Tf) subtypes by isoelectric focusing. Z Rechtsmed 85:255-261

Received May 7, 1980