

Papers

Danish type gelsolin related amyloidosis: 654G-T mutation is associated with a disease pathogenetically and clinically similar to that caused by the 654G-A mutation (familial amyloidosis of the Finnish type)

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Abstract

Background—Familial amyloidosis of the Finnish type (FAF, Finnish hereditary amyloidosis) is caused by a 654G-A mutation in the gelsolin gene on chromosome 9 resulting in the expression of mutant Asn-187 gelsolin which is abnormally proteolytically processed generating amyloidogenic fragments that polymerise into amyloid fibrils. We have recently shown that in a Danish and a Czech family with a clinical syndrome similar to FAF, including corneal lattice dystrophy, cranial neuropathy and skin changes, the disease is caused by another mutation at the same position, namely 654G-T predicting a Tyr-for-Asp substitution at 187 in secreted gelsolin.

Aim—To undertake a closer examination of the Danish subtype of FAF and report immunohistochemical and biochemical findings.

Results—Immunostaining of plasma gelsolin isolated from heterozygous FAF of the Danish subtype revealed a pattern similar to that found in FAF-Asn 187. The > 60 kDa gelsolin species contain an epitope characteristic of the amyloid forming region as revealed by an amyloid specific antibody, whereas the ~50 kDa fragments are devoid of it. Compared with the wild-type gelsolin peptide (Asp-187), the corresponding mutant peptide (Tyr-187) showed dramatically increased fibrillogenicity as revealed by quantitative thioflavine-T based fluorimetry; ultrastructurally, amyloid-like fibrils were formed by the mutant peptide. Immunohistochemistry showed that antibodies directed against residues 231–242 of secreted gelsolin, representing the carboxy terminus of the sequence forming the amyloid protein (residues 173–243) laid down in the tissues in a fibrillar form in FAF, specifically labelled the amyloid deposited in rectum and skin in the Danish (654G-T) subtype.

Conclusions—The 654G-T mutation in the gelsolin gene gives rise to an amyloid disease clinically and pathogenetically similar to that caused by the 654G-A mutation. (J Clin Pathol 2000;53:95–99)

Keywords: amyloidosis; Finnish familial amyloidosis; gelsolin mutation 654G-T; fibrillogenesis

Familial amyloidosis of the Finnish type (FAF, Finnish hereditary amyloidosis, amyloid polyneuropathy type IV) is a dominantly inherited systemic amyloid disease characterised by cranial neuropathy, corneal lattice dystrophy, skin changes, and renal and cardiac manifestations.^{1–3} The disease is caused by a point mutation, 654G-A, in the gelsolin gene^{4–6} resulting in the expression of amyloidogenic Asn-187 gelsolin and the accumulation of a 71 residue fragment of mutant gelsolin, Ala173-Met243, as amyloid fibrils in the tissues.^{7–8} The Asn-187 gelsolin mutation has been found in all FAF patients so far described from Finland, as well as in occasional American, Dutch, and Japanese families.^{9–14} In contrast, in a Danish family and a Czech family with clinical syndromes similar to FAF, we found another mutation, namely 654G-T, in the gelsolin gene predicting a Tyr for Asp mutation at position 187 in secreted gelsolin.¹⁰ We have now undertaken a closer examination of the Danish subtype of FAF and report the immunohistochemical findings and the detection of an abnormal fragmentation of mutant gelsolin in plasma. We furthermore show that the substitution of Asp with Tyr at position 187 in synthetic gelsolin peptides is associated with markedly accelerated fibrillogenesis.

Methods

TISSUE AND PLASMA SAMPLES

We studied biopsy specimens from skin and rectum from three siblings of a Danish family with familial amyloidosis with slowly progressive cranial neuropathy and corneal lattice dystrophy.¹⁵ The 654G-T mutation in gelsolin in this family was identified by allele specific

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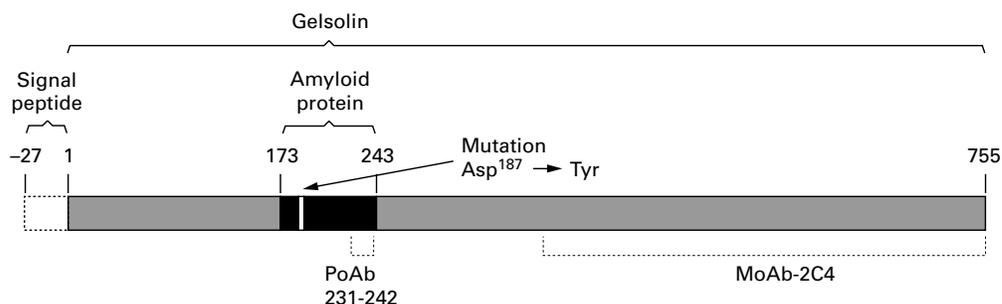


Figure 1 Schematic presentation of the amyloid forming region (residues 173–243) and mutation site in secreted gelsolin (residues 1–755) in Danish-type FAF, as well as the regions of the gelsolin molecule to which the amyloid specific polyclonal antibody (PoAb 231–242) and the monoclonal antibody MoAb-2C4 are directed.

oligonucleotide slot-dot hybridisation and by sequencing.¹⁰ Plasma gelsolin was isolated from a 57 year old male patient of the same Danish family. He had suffered from cranial amyloidotic polyneuropathy for about 10 years at the time of sampling. As controls we used plasma gelsolin isolated from Finnish subjects with FAF (two heterozygous (male) and one homozygous (female) patients with an identified 654G-A gelsolin mutation), plasma from healthy controls, and tissue samples from patients with FAF (Asn-187; three cases), AA amyloidosis (two cases), and AL amyloidosis (one κ and one λ case).

ANTIBODIES

We used two different antigelsolin antisera (fig 1): murine monoclonal antigelsolin antibody (MoAb-2C4) raised using a chymotryptic 47 kDa C-terminal actin binding fragment of gelsolin (Sigma), and polyclonal rabbit anti-amyloid antibodies (PoAb231–242) raised using synthetic VHVSEEGTEPEA amide corresponding to the carboxy terminus (residues 231–242) of the region forming the amyloid protein in FAF.^{7, 16} The polyclonal antibodies were purified by immunoaffinity chromatography on Sepharose coupled to a gelsolin peptide corresponding to residues 231–241 in secreted gelsolin.

IMMUNOBLOTTING

For western blots, proteins were separated on 10% SDS-PAGE, electroblotted on nitrocellulose membranes, and detected with murine monoclonal or rabbit polyclonal antibodies using as second antibodies peroxidase or alkaline phosphatase conjugated rabbit or swine immunoglobulins to murine or rabbit immunoglobulins (Daka), respectively. Development was performed with 3,3'-diaminobenzidine tetrahydrochloride and hydrogen peroxide or with nitroblue tetrazolium and 5-bromo-4-chloro-3-indolyl phosphate in dimethylformamide (AP Conjugate Substrate kit, Bio-Rad), respectively. As molecular standards, we used prestained protein marker, broad range (New England Biolabs) and prestained SDS page standards, low range (Bio-Rad).

AFFINITY CHROMATOGRAPHY

Isolation of gelsolin was carried out by immunoaffinity chromatography on a MoAb-2C4-Sepharose column using Prot On Kit 2 (Multi-

ple Peptide Systems) in the presence of 1 mM EGTA. Affinity chromatography on immobilised cibacron blue F3GA was performed in the presence of 1 mM EGTA as described previously.¹⁷ For routine testing, 100 μ l of plasma was chromatographed on 1.0 \times 1.5 cm cibacron blue column.

HISTOCHEMISTRY

Tissue specimens were studied for the presence of amyloid by routine congo red staining and polarisation microscopy. Immunoperoxidase staining was carried out as previously described.¹⁶ For immunogold staining, tissue sections 1 μ m thick were deparaffinised and after three washings incubated with 5% normal goat serum diluted in phosphate buffered saline (PBS), pH 7.2, containing 0.1% bovine serum albumin and 0.1% sodium azide. The sections were then incubated with the purified rabbit PoAb 231–242 antibodies for 30 minutes, followed by three washings with PBS. Gold labelled goat antirabbit IgG (Auro Probe LM GAR, Amersham) at 1:40 dilution was added and incubated for one hour, followed by three washings with water. The sections were developed by a silver enhancement kit (Inten SEM, Amersham).¹⁸

IN VITRO FIBRILLOGENESIS

For fibril formation, synthetic peptides homologous to wild-type (SFNNGDCFILD-amide; residues 182–192) or mutant (SFNNGYCFILD-amide and SFNNGNCFILD-amide representing residues 182–192) gelsolin were used (Multiple Peptide Systems). The purity of the peptides was > 95% as judged by high pressure liquid chromatography on a Vydac C18 column (fig 2).

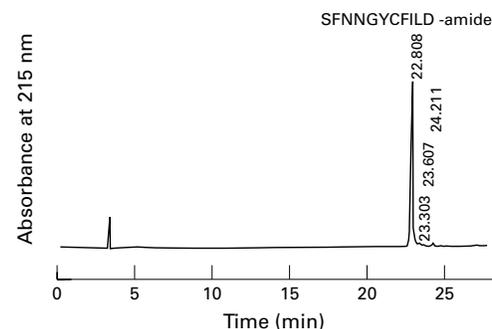


Figure 2 Reverse phase high pressure liquid chromatography on a Vydac C18 column of synthetic mutant 182–192 gelsolin peptide (Tyr-187).

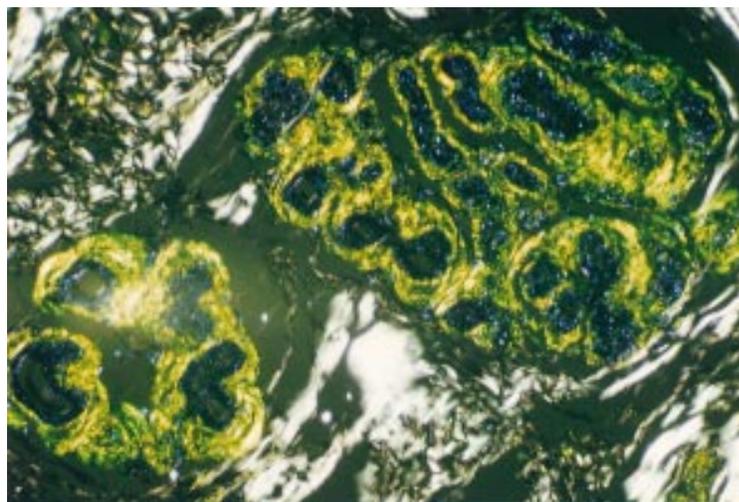


Figure 3 Dermal eccrine glands in Danish subtype (Tyr-187) familial amyloidosis showing prominent amyloid deposits in thickened basement membranes. Congo red staining, polarised light, magnification $\times 140$.

The peptides were dissolved in 10% acetic acid,¹⁹ pH 2.3, at a concentration of 5 mg/litre and analysed for Congo red positivity and green birefringence in polarised light after 15 minutes and overnight incubation at room temperature.

MONITORING OF FIBRIL FORMATION

Fibril formation was monitored by quantitative thioflavine-T fluorometry as described.^{20, 21} In this study we used the excitation and emission maxima of 450 nm and 482 nm, respectively, and a Perkin-Elmer Luminescence LS 30 spectrometer.

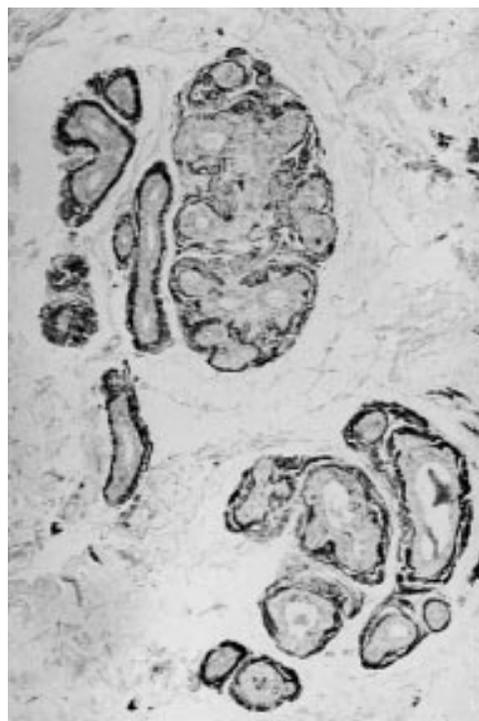


Figure 4 Immunogold-silver staining with amyloid specific antibody PoAb 231–242 of a skin biopsy in Danish subtype (Tyr-187) amyloidosis. Amyloid located in basement membranes is specifically stained. Magnification $\times 134$.

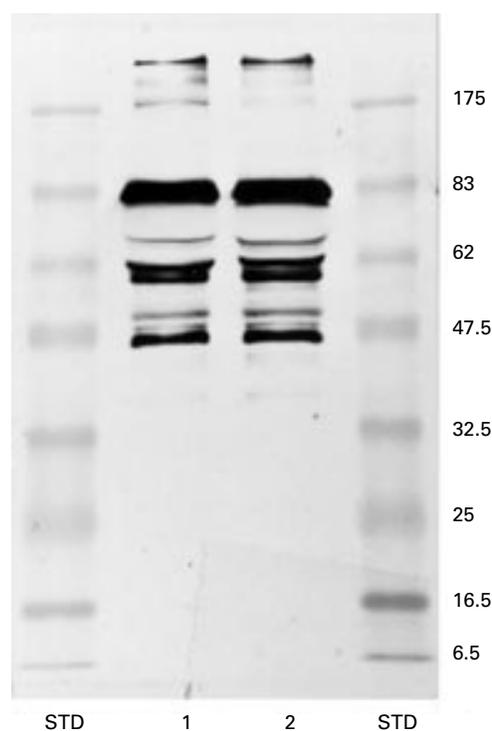


Figure 5 Comparison of heterozygous Danish subtype FAF (Tyr-187) and heterozygous Asn-187 FAF by immunoblotting by the monoclonal MoAb-2C4 antibody. Lane 1, FAF Asn-187; lane 2, Danish FAF (Tyr-187); STD, prestained molecular weight markers (kDa).

ELECTRON MICROSCOPY

Negative staining electron microscopy was carried out on peptide samples that were dissolved in water or 10% acetic acid at a concentration of 5 mg/litre, incubated at room temperature for 15 minutes and then kept frozen at -20°C . After thawing, the samples were diluted 1:1 in water, applied to a copper grid coated with pioloform and carbon, and stained with phosphotungstic acid adjusted to pH 6.5 with potassium hydroxide. The grids were examined with a JEOL-JEM-100CX transmission electron microscope at an operating voltage of 60 kV.

Results

HISTOCHEMISTRY

Prominent amyloid deposits were found along the collagen fibres in rectal mucosa and the thickened basement membranes of the dermal eccrine glands (fig 3). The amyloid deposits were specifically stained by the antigelsolin antibody PoAb231–242 directed against the carboxy terminus of the region forming the amyloid subunit protein (figs 1 and 4). PoAb231–242 did not stain tissue amyloid in AA or AL amyloidosis.

PLASMA GELSOLIN

Plasma gelsolin from heterozygous Danish subtype FAF was isolated by affinity chromatography and subjected to immunoblotting with MoAb-2C4 antibody. In addition to normal sized gelsolin, lower molecular weight fragments were detected; the pattern in Danish heterozygous FAF was similar to that found in

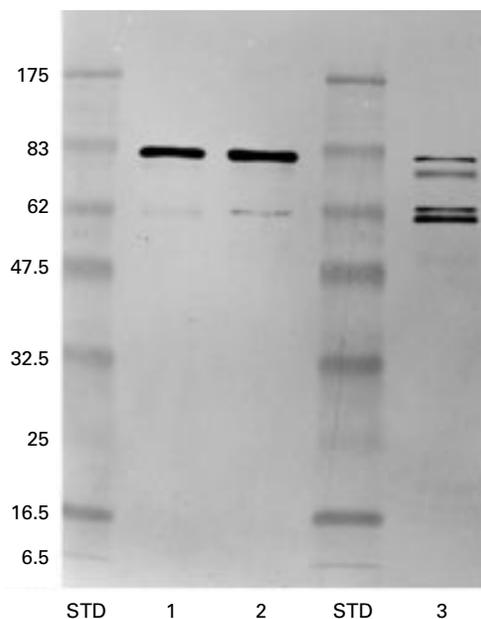


Figure 6 Immunostaining of the gelsolin fraction isolated from FAF plasma by affinity chromatography on an MoAb-2C4 Sepharose column by polyclonal anti-amyloid antibodies directed against the C terminus of the amyloid region (231–242). Lane 1, heterozygous FAF Tyr-187 (Danish); lane 2, heterozygous FAF Asn-187; lane 3, homozygous FAF Asn-187; STD, prestained molecular weight markers (kDa). Note that the ~50 kDa bands (fig 5) are not stained by the amyloid specific antibody as they are devoid of the amyloid forming sequence.

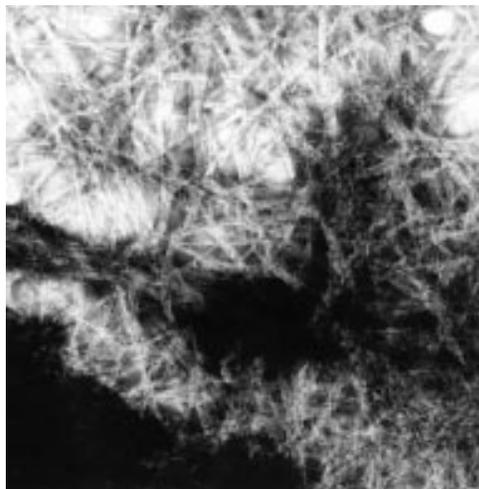


Figure 7 Electron micrograph of negatively stained protein assemblies formed by the mutant (Tyr-187) peptide, corresponding to residues 182–192 in gelsolin. Magnification $\times 59\,595$.

heterozygous Finnish patients with the Asn-187 mutation (fig 5). The antibody PoAb 231–242, directed against the amyloid forming region of gelsolin (fig 1) recognised the > 60 kDa gelsolin species, but not the ~50 kDa fragments (fig 6), which are devoid of the amyloid forming sequence.

IN VITRO AMYLOIDOGENESIS

Synthetic mutant Tyr-187 (Danish type) and Asn-187 (Finnish type) peptide, corresponding to residues 182–192 of gelsolin, formed amyloid-like fibrils in vitro (fig 7, Danish-type).

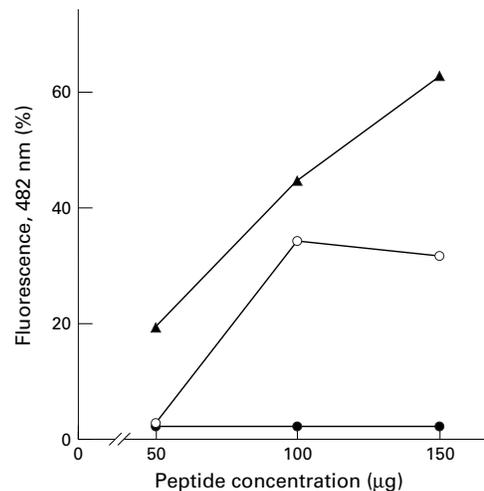


Figure 8 Comparison of amyloid fibril formation at different peptide concentrations by mutant (Tyr-187) gelsolin 182–192 peptide corresponding to Danish type amyloidosis (filled triangles), by mutant (Asn-187) gelsolin 182–192 corresponding to Finnish amyloidosis (empty circles), and wild-type (Asp-187) gelsolin 182–192 peptide (filled circles), as measured by quantitative thioflavine-T based fluorimetry.

The mutant peptide aggregates were Congo red positive and showed green birefringence in polarized light. Fibril formation by the mutant peptides was dramatically enhanced as compared with the corresponding wild-type (Asp-187) peptide (fig 8).

Discussion

The Danish type of familial amyloidosis with cranial neuropathy and corneal lattice dystrophy was described in 1979 by Boysen *et al* in five siblings of a family who had lived in Denmark for several generations.¹⁵ No genealogical connections to Finland could be shown for the family. The clinical features of the disease are very similar to those described by Meretoja in the Finnish families with hereditary amyloidosis, FAF.¹ In contrast to FAF, where the gelsolin mutation has been identified as 654G-A,^{4–6} a 654G-T mutation causes the Danish subtype of FAF.¹⁰ This mutation predicts a tyrosine for aspartic acid substitution at position 187 in secreted gelsolin.

The results of this study show that the biological consequences of the 654G-T mutation in gelsolin are very similar to those of the 654G-A mutation. Immunoblotting of plasma gelsolin in heterozygous Danish subtype FAF reveals a fragmentation pattern similar to that found in FAF (Asn-187)^{22–23}; in addition to normal sized gelsolin, a series of lower Mr C-terminal fragments is found. Thus the substitution of aspartic acid at position 187 in gelsolin with either asparagine or tyrosine is associated with abnormal degradation. The most likely explanation for the abnormal fragmentation, and the generation of the disease specific 65 kDa fragment,^{22–23} is that residue 187 represents a critical site where a substitution of an amino acid with a charged side chain (Asp) with one with an uncharged (Asn) or hydrophobic (Tyr) side chain induces a novel proteolysis site at 172–173 by the mechanism of local unfolding.^{24–25} This

conclusion is also substantiated by cell culture studies that show that transfection of COS-1 cells²⁶ with cDNA of Asp 187Asn/Tyr gelsolin is followed by the secretion in the medium of an abnormal fragment of about 68 kDa size representing a C-terminal fragment of gelsolin, starting with Ala 173.

The experimental data also suggest that the gelsolin fragments containing the mutation site are highly fibrillogenic and rapidly polymerise into insoluble amyloid fibrils. Compared with the corresponding wild-type gelsolin peptide, mutant Tyr-187 peptide was very fibrillogenic as revealed by quantitative thioflavine-T based fluorimetry. Moreover, electron microscopy showed that the fibrils created by the mutant Tyr-187 peptide were amyloid-like. Thus the 654G-T mutation appears to have a dual effect on amyloidogenesis: it is associated with the generation of a novel proteolysis site in gelsolin (172–173) resulting in the release of abnormal fragments that polymerise to amyloid fibrils during further proteolysis owing to the mutation induced, highly fibrillogenic amino acid sequence.

As shown previously for FAF(Asn-187), the circulating ~65 kDa gelsolin fragment, containing the amyloid sequence and derived from mutant gelsolin only, is FAF specific and can be used for diagnostic purposes.^{22–23} The same test can be used for diagnosing the Danish subtype of FAF as the same sized fragment is generated in this disease. The disease specific fragment can be detected by the MoAb-2C4 antibody which recognises the non-amyloid region of the molecule (fig 1), or more specifically, by the amyloid directed PoAb 231–242 antibody (fig 1), as shown here.

In conclusion, the data presented here support the view that the 654G-T mutation in the gelsolin gene gives rise to an amyloid disease clinically and pathogenetically similar to that caused by the 654G-A mutation.

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