

The Role of Cystatin C in Cerebral Amyloid Angiopathy and Stroke: Cell Biology and Animal Models

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A variant of the cysteine protease inhibitor, cystatin C, forms amyloid deposited in the cerebral vasculature of patients with hereditary cerebral hemorrhage with amyloidosis, Icelandic type (HCHWA-I), leading to cerebral hemorrhages early in life. However, cystatin C is also implicated in neuronal degenerative diseases in which it does not form the amyloid protein, such as Alzheimer disease (AD). Accumulating data suggest involvement of cystatin C in the pathogenic processes leading to amyloid deposition in cerebral vasculature and most significantly to cerebral hemorrhage in patients with cerebral amyloid angiopathy (CAA). This review focuses on cell culture and animal models used to study the role of cystatin C in these processes.

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The cysteine protease inhibitor, cystatin C (23), also known as γ trace (67), is found in all mammalian body fluids and tissues (23). In vitro experiments have indicated that it can inhibit the cysteine proteases cathepsins B, H, K, L and S (for review, see 18). In addition to being a protease inhibitor, cystatin C itself is a target of proteolysis (138, 143) and is inactivated by proteolytic degradation by cathepsin D and elastase (1, 92). Cathepsins are lysosomal proteases required for housekeeping function during protein turnover and they differ in structure, substrate-specificities and biochemical characteristics (159). Uncontrolled proteolysis as a result of imbalance between active proteases and their endogenous inhibitors has been associated with different diseases such as Alzheimer disease (AD) (116), ischemia (70, 122, 129, 158, 180), rheumatoid arthritis (157), renal failure (80), multiple sclerosis (19), osteoporosis (41), muscular dystrophy (148), inflammatory periodontal disease (87), inflammatory lung disease (29), inflammation and trauma (12), and various types of cancer (30, 44, 154). For review of the involvement of proteases and their inhibitors in the processes of neuronal degeneration and repair of the nervous system, see (156). This review will describe the data implicating cystatin C in pathological

processes of the central nervous system, mainly in CAA and cerebral hemorrhage.

MOLECULAR MECHANISM OF VARIANT CYSTATIN C FIBRIL FORMATION AND AMYLOID DEPOSITION IN HEREDITARY CEREBRAL HEMORRHAGE WITH AMYLOIDOSIS, ICELANDIC TYPE

Hereditary cerebral hemorrhage with amyloidosis, Icelandic type (HCHWA-I) (9, 61), also called hereditary cystatin C amyloid angiopathy (HCCAA) (126), is an autosomal dominant form of CAA. Amyloid deposition in cerebral and spinal arteries and arterioles leads to recurrent hemorrhagic strokes causing serious brain damage and eventually fatal stroke (61). The amyloid deposited is composed mainly of a Leu68Gln variant of cystatin C (5, 36, 51, 95, 130). A heterozygous point mutation, identical to that found in the cystatin C gene of these patients, was also identified in a Croatian man with CAA and intracerebral hemorrhage (56). Thus, sporadic CAA in some patients may be associated with mutations in the cystatin C gene (56, 110).

The cystatin C isolated from the leptomeninges of HCHWA-I patients was found to be amino-terminally truncated, starting at position 11 of normal cystatin C (36, 51). A serine protease, elastase, cleaves

cystatin C in vitro, generating a very stable product that lacks the amino-terminal 10 residues of full-length cystatin C (4). There is evidence to indicate that this modification may occur extracellularly in vivo as well because a truncated form of cystatin C was isolated from human urine (64). However, cystatin C isolated from cerebrospinal fluid (CSF) of HCHWA-I patients and from unaffected individuals is amino-terminally intact (11, 99, 125, 172), although a minor component starting at residue Leu9 was also found (11). Moreover, cystatin C synthesized by monocytes of carriers of the mutation also has full-length sequence (11, 155), and cells transfected with either the wild-type or variant genes secrete only full-length cystatin C (172). The cleavage of the amino-terminal decapeptide of the amyloid may occur in vivo during long-term exposure of the protein to proteases before, during and/or after amyloid fibril formation. Structural studies revealed that the amino-terminal truncation likely is an event secondary to amyloid formation, and of no relevance for the development of the disease (48). Mass spectrometric characterization showed that after three months of storage of human CSF at -20°C , cystatin C was cleaved in the peptide bond between Arg8 and Leu9 and lost its eight amino-terminal residues, whereas this cleavage did not occur when the CSF was stored at -80°C , indicating a storage-related artifact rather than a physiological or pathological processing of the protein (31).

The biochemical properties of the variant cystatin C have been extensively studied. It was demonstrated that the substitution does not affect the inhibitory activity, because wild-type and Leu68Gln variant cystatin C produced in *Escherichia coli* expression sys-

tems and by tissue culture cells effectively inhibited the cysteine protease cathepsin B (2, 172). However, the Leu68Gln is within the hydrophobic core of cystatin C (8), thus reducing the stability of the protein. It was hypothesized that this amino acid substitution may induce conformational changes, affecting the proteolytic susceptibility of the variant protein, as well as its predisposition to dimerization and amyloid fibril formation.

Effect of the Leu68Gln substitution on cystatin C dimerization. It was shown that recombinant wild-type cystatin C dimerizes when exposed to denaturing agents, low pH, or high temperature (45). These conditions lead to partial unfolding of cystatin C molecules with subsequent refolding into a dimeric state (45), in which the protein loses its capacity to inhibit papain-like cysteine proteases. Analysis of wild-type and variant cystatin C dimerization using recombinant proteins expressed in *Escherichia coli* (46, 49) has shown that the proteins differed in their tendency to dimerize and form aggregates. The wild-type protein was monomeric even after prolonged storage at elevated temperatures, while the variant proceeded to dimerize and lose biological activity immediately after refolding (2). The dimers and aggregates of the Leu68Gln variant recombinant protein were formed at normal body temperature, nearly 25°C lower than that needed for wild-type cystatin C dimerization (2). The most likely site for physiological dimerization would be in acidic cellular compartments. The pH in lysosomes is in the range of 4.6 to 5.0 which corresponds to the pH at which recombinant cystatin C dimerized to some extent in vitro. Consistent with this, a possible role of lysosomes in amyloid fibril formation has been proposed (52, 147). The lysosome could be the site of partial protein denaturation yielding an intermediate, which can associate into amyloid fibrils faster than it can be degraded. Similarly, it was shown that partial denaturation of transthyretin under conditions which mimic the acidic environment of a lysosome is sufficient to effect amyloid fibril formation by self-assembly of a conformational intermediate (37).

Cystatin C also forms dimers in mammalian cells overexpressing either the wild-type or variant forms of the gene (22, 112, 172).

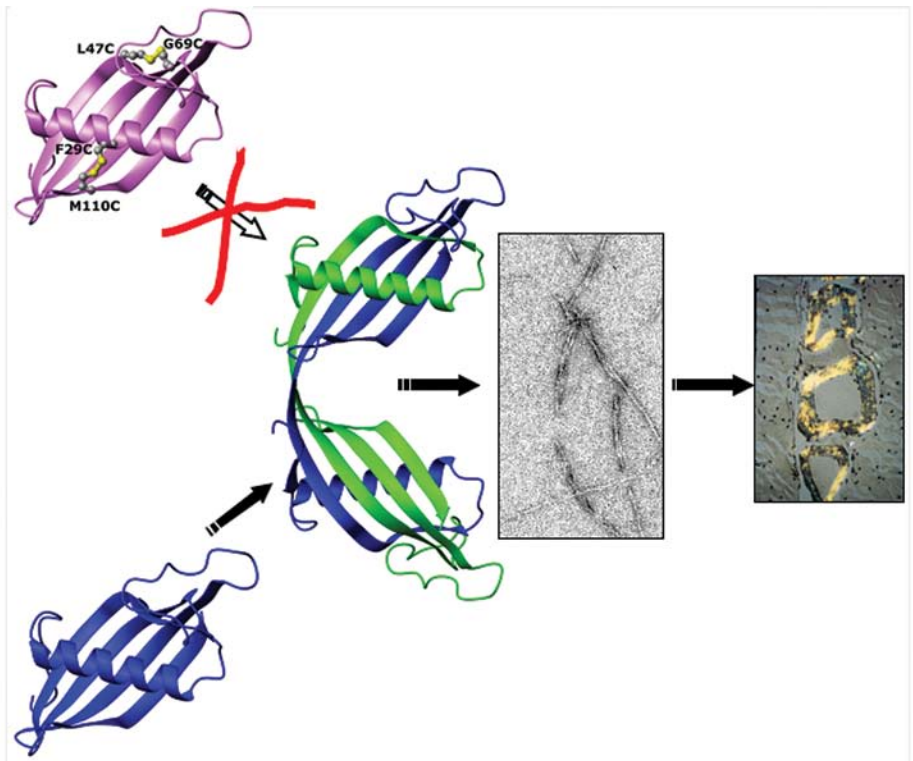


Figure 1. Domain swapping (blue and green) seems to be involved in the formation of amyloid fibrils from wild-type and Leu68Gln variant cystatin C. Proteins stabilized against domain swapping by disulfide bridges (purple) do not produce amyloid fibrils. Amyloid fibrils were formed following incubation of non-stabilized cystatin C (blue) at a concentration of 3 mg/ml in 10 mM glycine buffer, pH 2.0., containing 0.2% sodium azide, at 48°C.

A study of stably transfected NIH/3T3 cells suggested that cells expressing the gene encoding Leu68Gln cystatin C had intracellular accumulation of insoluble variant cystatin C mainly in the endoplasmic reticulum (22). In Chinese hamster ovary transfected cells, variant cystatin C formed stable intracellular dimers that are partially retained in the endoplasmic reticulum and degraded. The bulk of mutant cystatin C that is secreted does not dissociate and is exported as an inactive dimer (16). Blocking exit from the endoplasmic reticulum with brefeldin A resulted in increased dimerization of cystatin C and removal of brefeldin A caused an increase in the intracellular monomeric form. It was suggested that cystatin C dimerization occurs in the endoplasmic reticulum and dimer dissociation occurs later in the secretory pathway (112). Some degree of intracellular dimerization was identified by chromatography on AcA-44 column of lysates of Chinese hamster ovary cells transfected with the wild-type cystatin C cDNA (79). Mass spectrometry and Western blot analyses of several human astrocytic glioma U251 and human kidney 293 stably transfected cell

lines, showed the existence of cystatin C dimers in cell homogenates with high levels of overexpression of cystatin C (172). Dimerization directly correlated with cystatin C concentration in the sample, although the variant cystatin C formed dimers at concentrations lower than those necessary for dimerization of the wild-type protein. The mature, active form of cystatin C was secreted as a monomer into the culture medium (112, 172). The temporal profile of cystatin C production and secretion by pulse-chase studies revealed that both wild-type and variant cystatin C were expressed and cleared from the cells at the same rate and to a similar extent (172). The variant was found along with wild-type cystatin C in plasma and CSF of patients. Approximately equal amounts of cystatin C dimers and monomers were demonstrated in plasma from HCHWA-I patients, whereas only monomers could be found in normal plasma. Leu68Gln-wild-type cystatin C heterodimers seem to be present in the dimeric cystatin C population. CSF from patients also contained cystatin C dimers and monomers, but the dimeric fraction was minute. CSF from control patients did not

contain dimeric cystatin C (21). These data suggest that dimerization of wild-type and variant cystatin C occurs in compartments with high concentrations of the protein and that the Leu68Gln variant dimerizes at lower concentrations than the wild-type protein.

The crystal structure of human cystatin C revealed that the protein refolds to produce very tight 2-fold symmetric dimers while retaining the secondary structure of the monomeric form (75). The dimerization occurs through 3-dimensional (3D) domain swapping, a process involving exchange of sub-domains of the monomeric proteins. The exchange of domains is symmetric, or closed-ended, leading to intertwined and closed dimeric species. It has been suggested that in an open-ended, run-away mode, the mechanism of domain swapping could lead to infinite linear polymerization (76), such as is characteristic of amyloid fibrils. Since domain swapping of cystatin C involves an extended β -sheet, the final aggregate could be consistent with cross- β structure, which is believed to be at the heart of the molecular architecture of amyloid fibrils (76). Association of 3D domain-swapped cystatin C dimers into larger structures with involvement of intermolecular β -sheet interactions has been confirmed in another crystallographic study (74). In addition, it has been demonstrated that cystatin C with amino-terminal truncation forms essentially identical 3D domain-swapped dimers (73). However, the absence of the N-terminal decapeptide may facilitate further association of the protein via β -sheet interactions. Prevention of domain swapping inhibits dimerization and amyloid fibril formation of cystatin C (119) (Figure 1). The process of 3D domain swapping destroys one of the key elements of the cysteine protease-binding epitope. This explains the complete loss of the protease inhibitory activity reported for dimeric cystatin C (45, 46). The dimerization process influences the equilibrium between the proteolytic and inhibitory activities, and may be, therefore, relevant for physiological regulation of cysteine protease activity, for amyloid formation, and for the occurrence of hemorrhages.

Effect of the Leu68Gln substitution on cystatin C structure and proteolytic susceptibility. Most native proteins are relatively resistant to cleavage, whereas unfolded

proteins are usually good substrates for proteolysis. Thus, the rate at which a protein is degraded by a protease is indicative of its stability, enabling comparison of a mutant protein to its wild-type counterpart. The fraction of molecules that are unfolded varies as a function of temperature for both proteins, but the transition for the mutant may occur over a lower temperature range than that for the wild-type protein. Some proteins aggregate because they are unfolded or incompletely folded, and thus escape proteolysis (128). Thus, a common mechanism of fibril formation emerges (101), and it was suggested that the effect of the Leu68Gln substitution on cystatin C is similar to that caused by amino acid replacements in other amyloid forming proteins: Glu22Gln variant of A β deposited in cerebral vasculature of patients with HCHWA, Dutch type (HCHWA-D) (94, 176), immunoglobulin light-chain associated with light-chain amyloidosis (69), transthyretin variants found in familial amyloid polyneuropathy (111), and amyloid formed in lysozyme amyloidosis (24). The first step of this mechanism is a change in conformation followed by the opening of new hydrophobic surfaces, finally leading to the fibrillogenic step.

Circular dichroism, NMR spectroscopy, and 1-anilinonaphthalene-8-sulfonic acid binding analyses of the recombinant proteins isolated from *Escherichia coli* inclusion bodies showed that the Leu68Gln variant is structurally very similar to human wild-type cystatin C. However, the variant protein was stable only in a restricted range of conditions. Outside of this range, the variant differed from the wild-type protein by an increased amount of hydrophobic components on the protein surface (49). Experiments that used cystatin C proteins expressed in a mammalian tissue culture system indicate that the variant cystatin C has a more unfolded structure than the wild-type protein, with stretches of hydrophobic residues exposed to the solvent (172). The exposure of hydrophobic fragments to a polar environment is an unfavorable thermodynamic state that often results in protein oligomerization and aggregation. This could be the driving force of amyloid formation. Moreover, the unfolded structure of variant cystatin C may also expose critical residues of the protein to proteases, resulting in increased degradability. Thus,

the unfolding of the protein leads to either aggregation or degradation depending on the proteins' concentration. When variant cystatin C is present in a low concentration in body fluids, such as the CSF, it is highly susceptible to extracellular proteolysis by a serine protease. High concentration of the variant protein within neuronal cells, as was shown within the cytoplasm and cell processes of pyramidal neurons, mainly in layers three and four of the cortex of aged individuals (96), may result in its aggregation and fibrillogenesis.

The concentration of cystatin C in the CSF of HCHWA-I patients is lower compared to normal controls (60). Mass spectrometry of tryptic peptides of cystatin C isolated from the CSF of patients or from media of cultured monocytes isolated from patients revealed that only wild-type cystatin C was detectable, raising questions about the expression and fate of the mutant protein (11, 155). While accumulation of this protein in vessel walls may explain its depletion from the CSF, reduced secretion or enhanced proteolysis may be additional explanations. It was shown that the wild-type and variant cystatin C are similarly expressed and constitutively secreted by a variety of tissue culture cell lines transfected with either gene (28, 35, 39, 172). Both had a relatively short intracellular half-life (about 73 minutes) in human glial and kidney culture cells stably transfected with the cystatin C genes. The mature, active form of cystatin C was secreted as a monomer into the culture medium (112, 172). However, media of clones expressing the gene encoding Leu68Gln cystatin C had lower amounts of the protein than clones expressing wild-type human cystatin C (22, 172). It was shown that the secreted wild-type protein is resistant to extracellular proteolysis, whereas the secreted variant cystatin C is quickly degraded. Wild-type cystatin C secreted into the culture media was stable for at least 5 days. Conversely, the level of the Leu68Gln variant cystatin C was significantly reduced in the media in one day and only traces were observed after 2 days of culture (172). Protease inhibitory profile showed that the serine protease inhibitor, DFP, blocked the protease activity. Thus, a serine protease is involved in the observed reduction of the variant protein. This activity is cell line specific, since different levels of proteolytic activity are re-

leased by many cell types. Furthermore, a significant amount of the protease activity is also present in CSF of normal individuals (172). This suggests that the low level of cystatin C in CSF of affected patients is due to enhanced proteolysis of the variant protein.

INVOLVEMENT OF CYSTATIN C IN ALZHEIMER DISEASE

A β , a processing product of the larger β amyloid precursor protein (APP) (82), is the major constituent of the amyloid fibrils deposited in the brain of patients with AD, Down syndrome, sporadic CAA, and HCHWA-D (27, 53, 107, 178). Several APP missense mutations within or flanking the A β region were found in the APP gene in a few early onset familial AD pedigrees (151). APP amino acid substitutions located in the middle of the A β peptide, at positions 21, 22, and 23 (26, 65, 81, 94) are associated with various degrees of cerebrovascular pathology, some of which are extensive and result in cerebral hemorrhage (25, 26, 38, 55, 58, 63, 65, 81, 94, 102, 124, 162, 171). However, amyloid fibrils are usually formed from processing products of normal precursor proteins (152). Thus, mechanisms other than mutations expressed as protein variants might play a crucial role in the process of amyloid formation (88). Some proteins associated with AD lesions may have a role in the pathological processes leading to amyloidogenesis and neuronal degeneration and others may bind secondarily to amyloid deposits.

Immunohistochemical studies revealed that cystatin C co-localizes with A β predominantly in amyloid-laden vascular walls, and in senile plaque cores of amyloid in brains of patients with AD, Down syndrome, HCHWA-D, and cerebral infarction (62, 71, 96, 104, 167) and of non-demented aged individuals (96). Cystatin C also co-localizes with A β amyloid deposits in the brain of aged rhesus and squirrel monkeys (173), dogs (160, 161), and transgenic mice overexpressing human APP (96, 149) (Figure 2).

The neuronal staining of cystatin C in AD brains was primarily limited to pyramidal neurons in cortical layers duplicating the pattern of neuronal susceptibility in AD brains: the strongest staining was found in the entorhinal cortex, in the hippocampus, and in the temporal cortex; fewer pyrami-

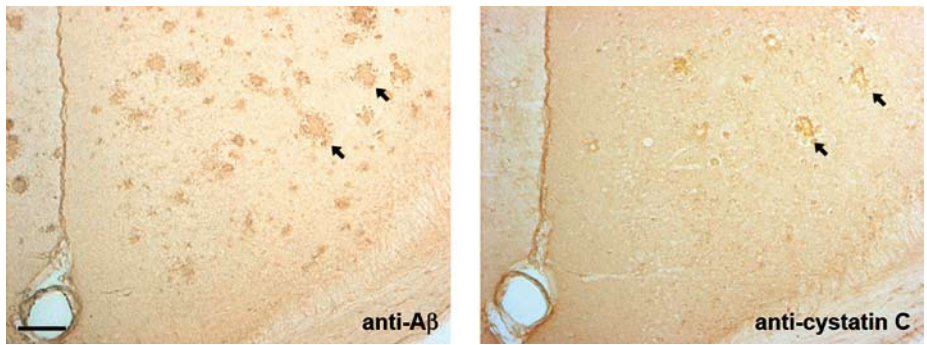


Figure 2. Immunohistochemical staining of amyloid deposits in the brain of an APP Tg2576 transgenic mouse at 18 months of age with anti-A β antibody (6E10) and with anti-cystatin C antibody. Arrows indicate same plaques stained with both antibodies and scale bar represents 50 μ m.

dal neurons were stained in the frontal, parietal, and occipital lobes (42). Immunostaining of cystatin C within neurons showed similar distribution to that of the endosomal/lysosomal proteases cathepsin B (42) and cathepsin D (33). Upregulation of cathepsin synthesis in AD neurons and accumulation of hydrolase-laden lysosomes indicate an early activation of the endosomal/lysosomal system in vulnerable neuronal populations, possibly reflecting early regenerative or repair processes (32, 33). These neuropathological observations support an association between cystatin C and cathepsin B and cathepsin D in AD, and suggest a model of cystatin C involvement in the process of neuronal death in AD. Using an antibody specific to the carboxyl-terminus of A β 42, intracellular immunoreactivity was observed in the same neuronal subpopulation strongly stained for cystatin C (96). This suggests that A β accumulates in a specific population of pyramidal neurons in the brain, the same cell type in which cystatin C is highly expressed. Co-localization of cystatin C with APP has been demonstrated in transfected human embryonic kidney HEK293 cells, mouse neuroblastoma N2a cells (145), in APP-overexpressing cultured human muscle fibers, and in muscle cells of patients with sporadic inclusion-body myositis (s-IBM) (163).

Co-localization of cystatin C with A β results from binding of cystatin C to A β , as well as to its precursor protein. Western blot analysis of immunoprecipitated cell lysate or medium proteins revealed binding of cystatin C to full-length APP and to secreted APP α (145). Deletion mutants of APP localized the cystatin C binding site within APP to the extracellular region of A β . This binding location seems to protect APP from β -secretase processing, resulting

in an increase in the non-amyloidogenic α -secretase cleavage, with no effect on the γ -secretase cleavage site. Accordingly, co-expression of cystatin C and APP in neuroblastoma cells resulted in increased secretion of APP α , while production of both A β 40 and A β 42 remained unchanged (145). The same observation was made in vivo, in transgenic mice expressing the human cystatin C gene (134). Enzyme-linked immunosorbent assay (ELISA) analysis of A β 40 and A β 42 concentrations in the brain of these mice showed no difference between cystatin C transgenic mice and their non-transgenic littermates (134). Thus, in vitro and in vivo overexpression of human cystatin C does not affect the levels of the potentially toxic A β .

Cystatin C binds not only to A β sequences within APP, but also to the peptide itself. Analysis of the association of cystatin C and A β by ELISA demonstrated that cystatin C interacts with both A β 40 and A β 42 in a concentration dependent manner at physiological pH and temperature. A specific, saturable and high affinity binding between cystatin C and A β was observed (145). Electron microscopic analysis of fibril formation revealed that incubation of cystatin C with either A β 40 or A β 42 inhibits A β fibril formation in a concentration-dependent manner (145). Thus, binding of cystatin C to soluble A β inhibits amyloid fibril formation, and the occurrence of the inhibitor in amyloid deposits may represent a residual effect of this association.

Genetic studies support a role for cystatin C in AD. The cystatin C gene (CST3), localized on chromosome 20 (3, 144) has three genetically linked base substitutions in the 3' region (14, 15). A G/A transition in exon 1 results in Ala/Thr variation in the coding region of CST3, within the signal

peptide. While the allele containing Ala at that position was called the A allele, the one containing Thr in the same position was called B allele. Several studies have linked CST3 gene polymorphisms with an increased risk of developing AD (20, 34, 40, 47, 54, 97, 127) and a possible interaction with apolipoprotein E (ApoE) genotype was noted. However, some studies have failed to show an association between the cystatin C gene and AD in a German cohort (43), a Dutch sample with early-onset AD (140), Japanese AD patients (103), and in early onset AD families (132).

Crawford et al (40) divided a multicentric AD population by age at onset and found an age related increased influence of the A allele of CST3. A significant interaction between the homozygous A genotype of CST3 and age of onset of AD was found, such that in the over 80 years age group this genotype was responsible for a 2-fold increased risk for the disease. This interaction was independent of the ApoE genotype (40). Another study of large European and American populations, with mean age at onset of 73.1 and 75.0 for AD and controls, respectively, showed linkage between the B allele and late onset AD with no synergistic association with ApoE allele (47). A synergistic association between the CST3 and ApoE ϵ 4 alleles was found in a Spanish sample. The CST3 B allele caused a 3-fold elevated risk of AD before age 70 and there was an 8-fold increase in risk for ApoE ϵ 4 carriers with this allele (20). In one study the combination of one or two CST3 B alleles and ApoE ϵ 4 carried a 14-fold increased risk for men and 16-fold for women. These risks apply to a shift in risk from ages 65 and older to younger ages (34). When it was attempted to determine the association between CST3 polymorphism and AD or vascular dementia, associations between CST3 B genotype and AD patients older than 75 years, or vascular dementia patients younger than 75 years were evident. A synergistic association of CST3 and ApoE ϵ 4 alleles was observed in predicting vascular dementia patients (97).

The amino acid exchange from Ala to Thr at the -2 position for signal peptide cleavage alters the hydrophobicity profile of the signal sequence (47), resulting in a less efficient cleavage of the signal peptide and thus a reduced secretion of cystatin C (17). The CST3 gene G73A polymorphism

functionally affects cystatin C plasma levels (123). Furthermore, a study of the targeting of the Thr haplotype in cultured retinal pigment epithelial and HeLa cells have shown that a proportion of the Thr protein undergoes incorrect trafficking. In contrast to the Ala haplotype that is targeted to the Golgi apparatus, the Thr variant was associated primarily with mitochondria, resulting in a substantial reduction in the efficiency of targeting cystatin C for secretion (131). A multicenter electroencephalographic (EEG) study analyzed the effects of CST3 haplotypes on resting cortical rhythmicity in subjects with AD and mild cognitive impairment. A relationship between the CST3 Thr haplotype and global neurophysiological phenotype (ie, cortical delta and alpha rhythmicity) was found. While ApoE ϵ 4 affects EEG rhythms in AD (77, 90, 91), the effects of CST3 polymorphism were independent of ApoE ϵ 4 co-presence (13).

Thus, multiple lines of research support a role for cystatin C in AD. It remains to be determined in what pathological, or alternatively, protective process(es) of the disease it is involved, and what the mechanism of its action is. Similar to a variety of activities that have been associated with other protease inhibitors in the brain (7, 68, 89, 139, 150), cystatin C has been implicated in the processes of neuronal degeneration and repair of the nervous system. Enhanced cystatin C expression was observed in response to injury, including facial nerve axotomy (113), perforant path transections (181), and hypophysectomy (83), in morphologically degenerative CA1 pyramidal neurons and reactive astrocytes of hippocampus following transient forebrain ischemia (70, 122, 129, 180) and after induction of epilepsy (10, 66, 100). It was proposed that cystatin C might facilitate the establishment of an axonal growth-promoting environment by protecting neurite growth-associated molecules from degradation by cysteine proteases in the deafferented hippocampus (6, 135). Oxidative stress also stimulates an increase in cystatin C expression in cultured neurons (120, 121) and in cerebral microvascular smooth muscle cells (170), suggesting a role in regulation of apoptosis. Moreover, a glycosylated form of rat cystatin C was found to be an autocrine/paracrine factor, required for the mitogenic activity of basic fibroblast growth factor on neural stem cells (153). These data suggest

that cystatin C may have a protective role in neurodegenerative diseases, including AD, in addition to its role in inhibition of A β fibrillogenesis.

INVOLVEMENT OF CYSTATIN C IN CEREBRAL AMYLOID ANGIOPATHY

While A β usually accumulates both in the cerebral blood vessels and in brain parenchyma as amyloid plaques, in some cases there is predominantly vascular A β deposition (165). The factors leading to vascular rather than parenchymal amyloid deposition are unknown and it is still unclear when CAA leads to hemorrhage. In vessels affected by CAA, local smooth muscle and elastic elements are lost and replaced by amyloid fibrils, thereby weakening the overall structure of the vessel (84, 104, 142, 164). Consequently, it was suggested that CAA predisposes towards cerebral infarction and cerebral hemorrhage. However, CAA is usually asymptomatic and only a subpopulation is at high risk of hemorrhage (57, 71, 78, 106, 109, 141, 164, 166, 168). Thus, CAA appears to be a prerequisite but not sufficient for vessel rupture, with additional factors playing important roles. Multiple genetic factors can be associated with the risk of CAA in the elderly. The role of ApoE in the genetics and pathogenesis of AD and of CAA both in AD and in non-demented elderly individuals has been well established. Whereas the ApoE ϵ 4 allele promotes deposition of A β in the cerebral vasculature (59, 108, 109, 117, 146, 177, 179), possession of ApoE ϵ 2 promotes the development of hemorrhage in vessels already laden with amyloid (57, 59, 109, 117, 118, 177).

A role for cystatin C in CAA-related hemorrhage is implicated from immunohistochemical studies of patients with AD, Down syndrome, HCHWA-D, intracranial hemorrhage, cerebral infarction, and of elderly patients without any neurological disorder, that revealed co-localization of cystatin C and A β in amyloid-laden vascular walls (62, 71, 104, 167). Maruyama et al (104) reported that only patients showing co-localization of cystatin C and A β immunoreactivity in their diseased cerebral vessels suffered fatal subcortical hemorrhages. The degree of cerebrovascular amyloid deposition in these patients was also greater than in patients without cerebral hemorrhages. Studies were conducted to find out

whether cystatin C exists as amyloid fibrils or as unpolymerized cystatin C absorbed onto or trapped within the bundles of A β amyloid fibrils. ELISA analysis of crude amyloid fibrils isolated from cerebral blood vessels of one patient revealed that cystatin C and A β have been included at the ratio of about 1:100 (115). In another case of sporadic CAA, isolation and chemical analysis of amyloid fibril proteins from leptomeningeal vessels revealed that while A β was fibrillar, cystatin C was soluble (105). It has been suggested that cystatin C deposition occurs secondarily to A β deposition and may increase the predisposition to cerebral hemorrhages (71).

Genetic studies show a role for cystatin C in CAA and stroke. In support of the interaction of CST3 with ApoE ϵ 4 in conferring vascular pathologies, a logistic regression analysis revealed that neither CST3 polymorphism nor its interaction with ApoE ϵ 4 were significant predictors of AD. However, a synergistic association of CST3 and ApoE ϵ 4 alleles was observed in predicting vascular dementia patients (97). Izumihara et al (72) suggested that while severe A β deposition and ApoE ϵ 2 and ϵ 4 alleles are risk factors for the occurrence of the hemorrhage, severe cystatin C accumulation is a risk factor both for the occurrence and enlargement of the hemorrhage, as well as for the tendency to induce recurrent hemorrhages. In addition, loss of vascular smooth muscle was observed in the intensely amyloid-laden vascular walls that showed cystatin C-immunoreactivity.

Cystatin C co-localization with amyloid, other than A β , was observed in a variety of disorders, such as hereditary gelsolin amyloidosis (familial amyloidosis, Finnish type) (85, 86) and familial cerebral amyloid angiopathy, British type, (50). Thus, cystatin C may have a role in CAA and hemorrhage in a variety of diseases that involve deposition of heterogeneous types of amyloid proteins.

CYSTATIN C IN ANIMAL MODELS OF CAA

A striking increase in CAA with aging was found in the APP transgenic mouse line APP23 (174). However, no significant correlation was found between CAA frequency or severity and amyloid load within age groups. 19-month-old APP23 mice had several focal cerebral hemorrhages that dramatically increased in frequency and size in

27-month-old mice. The anatomical distribution of the hemorrhages appeared very similar to the distribution of CAA. Antibodies to cystatin C showed appreciable staining of cerebrovascular amyloid in these mice, suggesting that mouse cystatin C is part of the amyloid. Similar to cystatin C staining of A β deposits in human brains, the cystatin C immunoreactivity was restricted to a subpopulation of amyloid-laden vessels and was clearly less intense than A β staining (96, 149, 174).

Non-human primates are good models to study cerebral changes that occur through aging. Neuropathologies characteristic of AD and normal aging in humans were found in senescent non-human primates (137, 169, 175). In aged rhesus monkeys (*Macaca mulatta*), amyloid deposition predominates in senile plaques with relatively minor vascular involvement. However, cerebrovascular deposits in aged squirrel monkeys (*Saimiri sciureus*), usually are more conspicuous than senile plaques (169). Because CAA in squirrel monkey resembles that in HCHWA-D it was hypothesized that a species-specific amino acid difference in A β between rhesus and squirrel may contribute to CAA in aged squirrel monkeys. However, it was shown that the predicted amino acid sequence of A β in squirrel monkeys is identical to that of rhesus monkeys and normal humans (93). It was also demonstrated that species differences in loci of amyloid deposition are not related to ApoE allotype because both rhesus and squirrel monkeys are homozygous for the same ApoE allele (114, 136, 182). However, sequence analysis of rhesus and squirrel monkey cystatin C cDNA revealed that squirrel monkey has Met at position 68, which is Leu in the rhesus and wild-type human cystatin C and Gln in HCHWA-I patients (173). An additional difference between squirrel and rhesus monkeys in cystatin C sequence was found at position 10, a residue that was shown to affect the specificity of the inhibitor for different cysteine proteases (98). The species-specific cystatin C sequences in humans, rhesus, and squirrel monkeys may be responsible for the variability of the amyloid deposits observed.

In humans co-localization of cystatin C with A β elevates the risk of hemorrhagic stroke. While co-localization of A β and cystatin C was demonstrated in vascular

and parenchymal deposits in the brains of rhesus monkeys and in vascular amyloid in brains of aged squirrel monkeys, there was no evidence of intracerebral hemorrhage (173). Abundant amyloid deposition and mild hemorrhage was found in the brain of one female squirrel monkey, known as "Baker," who died at the estimated age of 27 years from renal failure (169, 173). However, no systematic, large-scale study of hemorrhage in aged monkeys has been conducted. Thus, the specific cystatin C sequences in squirrel monkeys may contribute to the predominant deposition of amyloid in cerebral vessel walls, but not to an increased risk of hemorrhage.

OVEREXPRESSION OF CYSTATIN C AS A CAUSE OF CEREBRAL HEMORRHAGE

Cystatin C transgenic mice were generated in order to elucidate the role of increased expression of this protease inhibitor in vivo (134). These mice express either human wild-type or the Leu68Gln variant cystatin C genes under the transcriptional control of its own promoter (95), overexpressing the transgene along with its endogenous counterpart in the appropriate tissues (134). Lines of mice expressing various levels of cystatin C in the brain were selected. All selected lines had very high concentrations of the transgene in the blood. None of the mice had amyloid deposits either in the vessel walls or in the neuropil. Neuropathological examination of dead or ailing aged transgenic mice revealed mice with cerebral or subarachnoid hemorrhages (133). Conversely, no hemorrhages were observed in their non-transgenic siblings. These data demonstrate that elevated brain and/or blood levels of cystatin C can cause hemorrhagic strokes in the absence of vascular amyloid deposits. In the brain of aged individuals and AD patients, cystatin C co-deposits with A β . It has been shown that the risk of cerebral hemorrhage increases when high levels of cystatin C are present in cerebrovascular A β deposits. We suggest that cystatin C binding to A β may cause local accumulation of the protease inhibitor, contributing to hemorrhages.

SUMMARY

Accumulating data implicate cystatin C in Alzheimer disease, in CAA, and in cerebral hemorrhage. A variant cystatin C with a single amino acid substitution can form

amyloid fibrils and directly cause CAA and the resulting hemorrhages. Wild-type cystatin C binds to proteins that form amyloid fibrils, such as A β , and the resulting local accumulation of the inhibitor can result in hemorrhages as well. Furthermore, animal models suggest that high systemic levels of cystatin C can cause cerebral hemorrhages. Future studies will determine the mechanism by which cystatin C causes damage to vessel walls and leads to hemorrhage.

Two proteins, the cysteine protease inhibitor cystatin C and ApoE, are linked to late onset AD and to CAA, and both proteins have been implicated in CAA-related hemorrhage. While the ApoE ϵ 4 and ApoE ϵ 2 isoforms are risk factors for the occurrence of hemorrhages, severe cerebrovascular cystatin C accumulation is a risk factor not only for the incidence but also for enlargement and recurrence of hemorrhages. While both ApoE and cystatin C are factors contributing to cerebral hemorrhage, neither factor is sufficient to always produce hemorrhages. The relative significance of each of the factors involved, and the possible synergy between them remain to be determined.

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