

Potential Role of Protease Nexin-2/Amyloid β -Protein Precursor as a Cerebral Anticoagulant^a

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One of the hallmark features of Alzheimer's disease (AD) is deposition of the amyloid β -protein in extracellular neuritic plaques and within cerebral blood vessel walls.¹⁻⁵ Similar neuropathology is observed in brains of individuals with hereditary cerebral hemorrhage with amyloidosis-Dutch type.⁶ The amyloid β -protein is an insoluble, self-aggregating peptide with a molecular mass of 4.2 kDa.^{1,4} CDNA cloning studies revealed that the amyloid β -protein is derived from a larger precursor protein, termed the amyloid β -protein precursor (APP), which is encoded by a gene located on chromosome 21.⁷⁻¹⁰ Subsequent studies showed that APP is translated from at least three alternatively spliced mRNAs resulting in polypeptides of 695, 751 and 770 amino acids.¹¹⁻¹³ The latter two species contain an additional insert which is structurally similar to the inhibitory domain of Kunitz-type serine protease inhibitors (KPI). The secreted form of APP containing the KPI domain has been shown by us and others to be identical to the serine protease inhibitor, protease nexin-2 (PN-2).^{14,15} The identity of APP and PN-2 was demonstrated by amino terminal and additional peptide amino acid sequences of PN-2 aligning with amino acid sequences deduced from cDNAs for APP. Full-length APP is normally cleaved at the carboxy terminal side of Lys¹⁶ in the amyloid β -protein domain, resulting in a secreted PN-2 that contains the first fifteen amino acids of the amyloid β -protein at its carboxy terminus.¹⁶⁻¹⁸ Currently, little is known about the physiological function(s) of PN-2 and the other APP isoforms. Recent findings suggest a potential physiological role for PN-2/APP as a cerebral anticoagulant.

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PN-2 is the Predominant Isoform of APP in Human Brain

Studies have demonstrated that mRNA encoding the forms of APP that contain the KPI domain are present in most tissues.^{11-13,19} On the other hand, mRNA encoding the form of APP that lacks the KPI domain is found primarily in brain and has been suggested to be the major isoform of APP mRNA in this tissue. In human brain two major APP proteins have been identified with approximate molecular masses of 120 kDa and 105 kDa.²⁰⁻²² The 120-kDa isoform of APP protein in human brain has been shown to contain the KPI domain based on immunoblotting studies with a polyclonal antiserum raised against a synthetic peptide corresponding to a region of the KPI domain.²¹ The 105-kDa isoform of APP protein did not react with this same antiserum, suggesting that this APP isoform lacks the KPI domain.²¹ The seemingly abundant amount of APP protein that lacks the KPI domain in brain appeared to be consistent with the high levels of its corresponding mRNA in this tissue.

Recently we described a qualitative and quantitative study that analyzed the different APP protein isoforms in human brain.²³ In addition to immunoblotting analyses, our studies employed functional assays that were specific for KPI-containing isoforms of APP proteins. These assays included formation of sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) stable complexes with an ¹²⁵I-labeled protease²⁴⁻²⁶ and precipitation by trypsin-agarose. The immunoblotting and functional assays showed that human fibroblasts and human platelets contained a 120-kDa, KPI-containing form of APP protein. On the other hand, immunoblotting showed that normal and AD brain contained 120-kDa and 105-kDa forms of APP protein. However, the functional assays indicated that both forms of APP proteins in the brain contained the KPI domain.²³ Quantitative functional assays indicated that >80% of the total APP protein in human brain contains the KPI domain. The findings in this study were extended by functional analyses comparing APP proteins that were purified from brain and cerebrospinal fluid of normals and AD patients. A previous study indicated that <10% of the APP proteins in cerebrospinal fluid contain the KPI domain.²⁷ FIGURE 1 shows that APP proteins purified from cerebrospinal fluid formed <10% of the amount of SDS-PAGE stable complexes with ¹²⁵I-labeled protease than did APP proteins purified from brain. Similar findings were observed with purified APP proteins from normals and AD patients. Together, these KPI-specific functional assays indicate that the majority of APP in human brain contains the KPI domain, regardless of its molecular mass on SDS-polyacrylamide gels.

Recent immunohistochemical and immunoblotting studies by Arai *et al.*²⁸ detected APP proteins in various tissues and cell types of the central nervous system. However, most nonneural tissues were notably devoid of APP immunoreactivity. Extending these findings we have recently conducted quantitative immunoblotting studies to determine the levels of PN-2/APP in a variety of nonhuman primate tissues. These studies showed that PN-2/APP was most abundant in brain with some presence in testis and kidney (unpublished data). It is noteworthy that similar studies showed that protease nexin-1 (PN-1), a potent antithrombin, is also found primarily in brain. These studies raise the intriguing question as to why there is a rich and relatively exclusive investment of PN-2/APP, as well as PN-1, in brain.

PN-2/APP Is an Abundant Platelet α Granule Protein and Potent Inhibitor of Coagulation Factor XIa

Several findings have suggested a vascular contribution of PN-2/APP and the amyloid β -protein. APP mRNA transcripts have been recognized in human umbilical

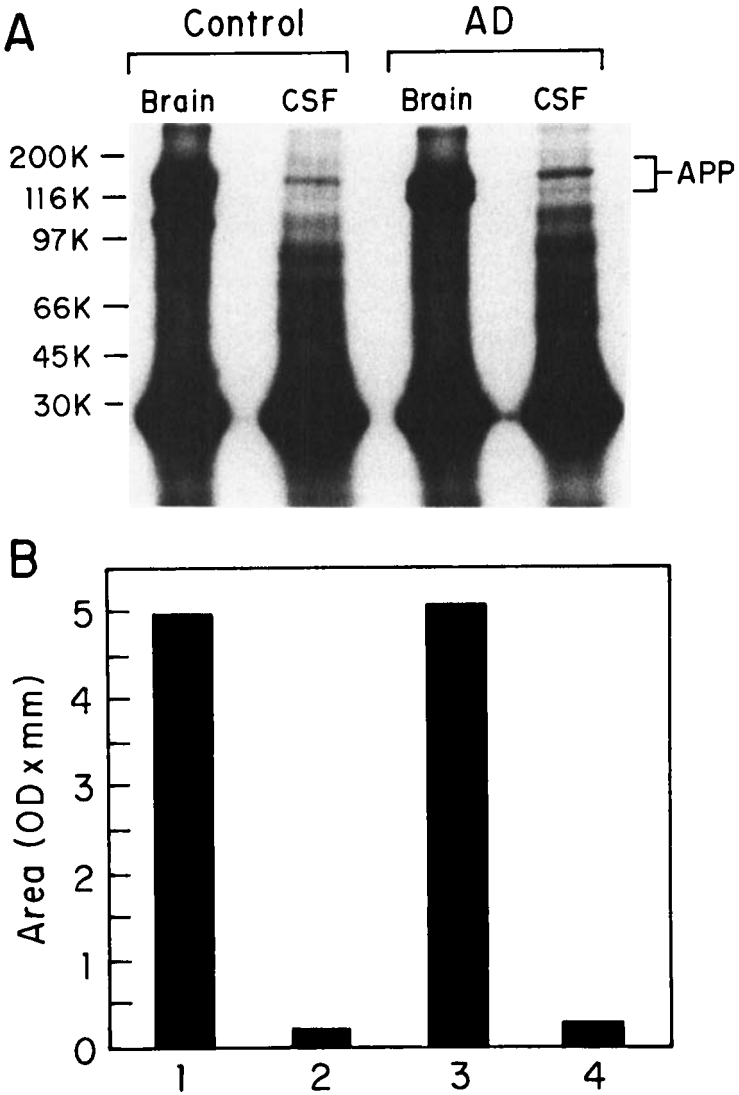


FIGURE 1. Quantitation of KPI-containing isoforms of APP in human brain and cerebrospinal fluid. (A) Equivalent amounts of APP proteins purified from normal brain tissue (*lane 1*), normal cerebrospinal fluid (*lane 2*), AD brain tissue (*lane 3*), and AD cerebrospinal fluid (*lane 4*) were incubated with a molar excess of ¹²⁵I-labeled epidermal growth factor binding protein and analyzed by SDS-PAGE with subsequent autoradiography. (B) The high molecular weight complexes between ¹²⁵I-epidermal growth factor binding protein and the KPI-containing isoforms of APP were quantitated by laser scanning densitometry. Purified cerebrospinal fluid APP contains < 10% the amount of KPI-containing isoforms than purified brain APP.

vein endothelial cells.²⁹ Furthermore, interleukin I, which is elevated in brain tissue from patients with AD and Down's syndrome, upregulates APP mRNA in endothelial cells through protein kinase C.^{29,30} In addition, various sized PN-2/APP proteins between 105 and 135 kDa have been recognized in leptomeningeal and cortical blood vessel walls.³¹ Other vascular sources for PN-2/APP have been sought. The finding that heparinized plasma contained some PN-2/APP immunoreactivity³² suggested that circulating cells in blood could be releasing this protein. Our recent studies showed that PN-2/APP circulates in blood as a platelet α granule protein and is secreted upon activation of platelets by physiological agonists.³³ The presence of platelet PN-2/APP has been confirmed by three other laboratories.³⁴⁻³⁶ When compared with other prominent cells of the intravascular compartment, platelets appear to be a relatively specific, intravascular source for PN-2/APP, accounting for 99% of its concentration in the intravascular compartment.³⁷ Platelet PN-2/APP is also a major protein in platelets accounting for 0.5% of total platelet protein.³⁷ Furthermore, remnant APP mRNA is found in human platelets and some of the full length protein is present in platelet membranes.³⁶ Together, these studies clearly indicate that platelets are the major circulating repository for PN-2/APP and provide an effective mechanism for the delivery and expression of large quantities of this protein at specific sites throughout the vasculature.

Investigations by us and others have sought to identify physiologic target proteases that are inhibited by PN-2/APP. Kinetic inhibition studies showed that PN-2/APP is a potent inhibitor of several "trypsin-like" and "chymotrypsin-like" serine proteases including trypsin ($K_i = 4.2 \times 10^{-10}$ M), chymotrypsin ($K_i = 1.6 \times 10^{-9}$ M), epidermal growth factor binding protein ($K_i = 5.8 \times 10^{-9}$ M) and the γ subunit of nerve growth factor ($K_i = 9.1 \times 10^{-9}$ M).^{14,26,34,38,39} More specifically, studies defining the protease inhibitory properties of PN-2/APP revealed that it is a potent inhibitor of coagulation factor XIa ($K_i = 5.5 \times 10^{-11}$ M).^{34,38} Factor XIa is the first protease in the intrinsic coagulation pathway whose deficiency leads to bleeding.⁴⁰ The inhibition of factor XIa by PN-2/APP was augmented by heparin suggesting that glycosaminoglycans may play an important role in the regulation of certain physiologic target proteases by PN-2/APP.^{34,38} Together, the findings that platelet PN-2/APP is secreted upon platelet activation along with other α granule constituents and it is a very effective inhibitor of factor XIa, support the notion that PN-2/APP has a physiologic function in regulating hemostasis.^{33,34,37,38,41} Further, its other protease specificities suggest that it may play a role in regulating other proteolytic events associated with wound repair.

Regulation of Coagulation in the Systemic and Cerebral Vasculature

Recently, a revised hypothesis on the initiation of blood coagulation was proposed based on studies by Naito and Fujikawa⁴² and Gailani and Broze,⁴³ which showed that factor XI, the first committed hemostatic protein, is a better substrate of thrombin than factor XIIa in the presence of a negatively charged surface like dextran sulfate or glycosaminoglycans. In this revised model, coagulation is initiated by generating small amounts of thrombin at the site of injury via the extrinsic pathway (FIG. 2). Thrombin may then serve as the "trigger" to activate factor XI, which initiates the subsequent steps of the intrinsic pathway. Activation of the intrinsic pathway leads to the amplification of thrombin formation, which serves two important functions: (1) sustaining the activation of factor XI by a factor XIIa-independent mechanism, and (2) catalyzing the formation of fibrin leading to clot formation. It is noteworthy that this model implicates thrombin and factor XIa as two key enzymes involved with the amplification system of hemostasis (FIG. 2).

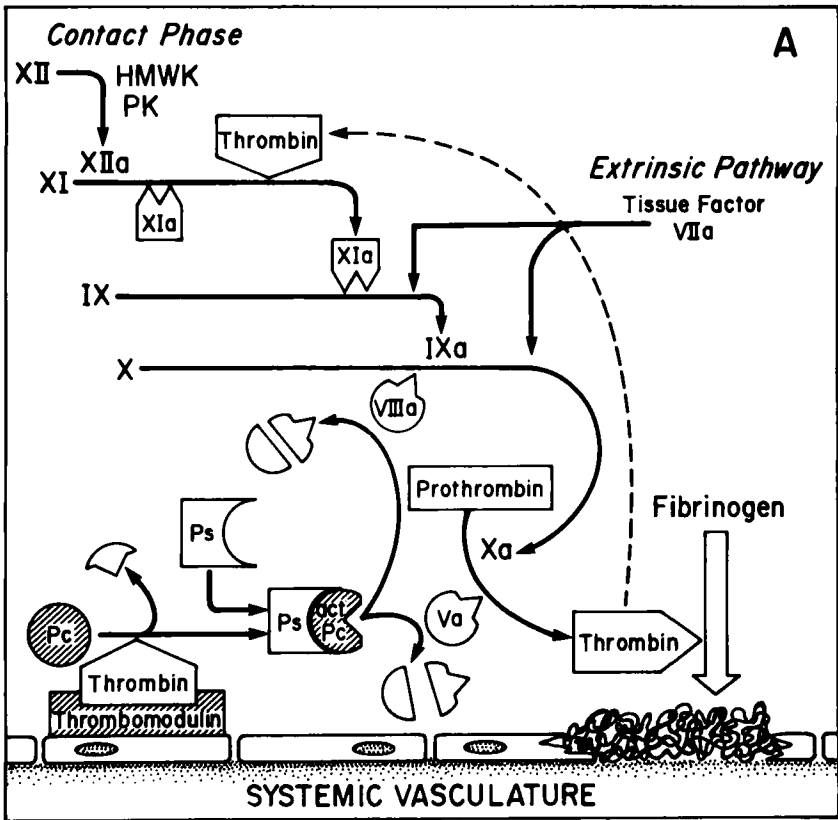


FIGURE 2A. Schematic representation of the regulation of the coagulation cascade in the systemic vasculature. Small amounts of thrombin generated by factor Xa via the extrinsic coagulation pathway can lead to activation of factor XI with subsequent amplification of the intrinsic coagulation pathway. Factor XIa can also autoactivate further contributing to the amplification of the intrinsic pathway. In the systemic vasculature, excess thrombin binds to thrombomodulin and converts protein C (Pc) into activated protein C (act. Pc). Optimal activity of Pc requires the presence of Protein S (Ps). Activated Pc-Ps complexes can inactivate coagulation factors Va and VIIIa via proteolysis.

Uncontrolled activation of the coagulation cascade can cause thrombosis leading to stroke and infarct. Systemically, regulation of coagulation is chiefly accomplished by two mechanisms, both of which involve the surface of the vascular endothelium.⁴⁴ The first anticoagulant mechanism is the heparin-antithrombin III system which involves glycosaminoglycans on the surface of the vascular endothelium. The other key anticoagulant molecule distributed on the plasma membrane of arterial, venous, capillary, and lymphatic endothelium is thrombomodulin (FIG. 2A). Thrombomodulin is an integral membrane protein which functions indirectly as an anticoagulant due to its ability to alter the substrate specificity of thrombin. When bound to thrombomodulin, thrombin loses its specificity for fibrinogen and becomes a potent activator

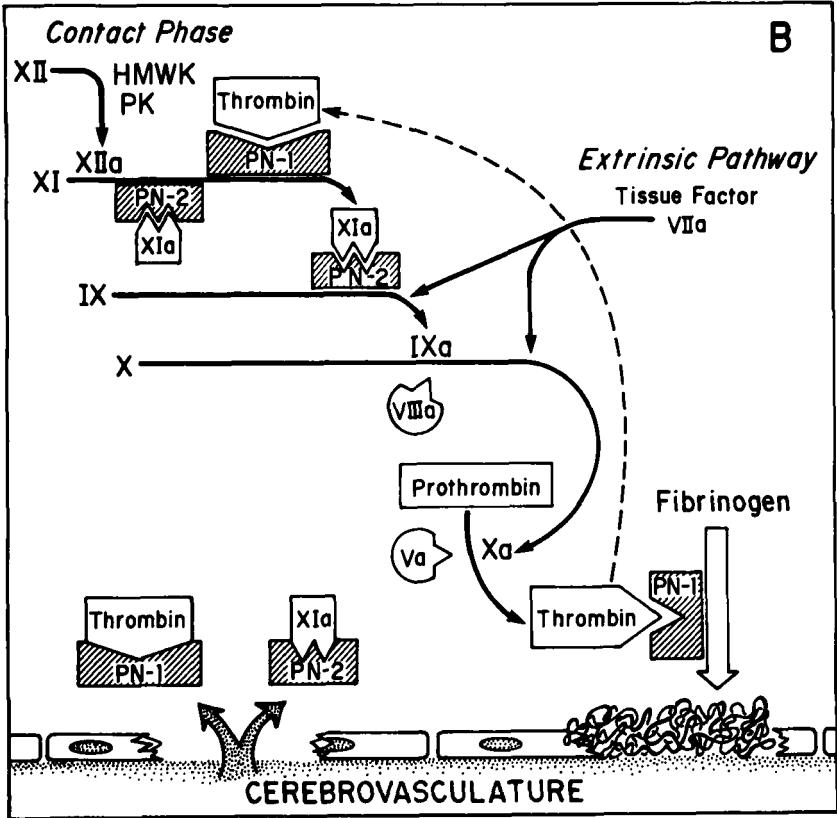


FIGURE 2B. Schematic representation of the proposed roles of PN-1 and PN-2 in the regulation of the coagulation cascade in the cerebrovasculature. Excess thrombin binds to PN-1 which blocks its ability to cleave fibrinogen or activate factor XI. Excess factor XIa binds to PN-2 which inhibits its ability to activate additional factor XI and the subsequent steps of the intrinsic coagulation pathway.

of protein C. Protein C is a serine protease which, in concert with protein S, exerts its anticoagulant effect by rapidly hydrolyzing coagulation factors Va and VIIIa. In order for thrombin to effectively activate protein *in vivo* it must be bound to thrombomodulin.

In the brain, however, thrombomodulin is virtually absent from much of the endothelium of cerebral blood vessels.⁴⁵ This finding suggests that the anticoagulant protein C system is not operative in the brain. Other regulatory molecules must be present in the brain and the cerebrovasculature to provide a means to control thrombin formation (Fig. 2B). As mentioned above, brain exhibits a very rich and exclusive investment of PN-2/APP, a potent inhibitor of factor XIa. In addition, normal brain, particularly around cerebral blood vessels, is richly invested in PN-1, a potent inhibitor of thrombin.^{46,47} We propose that abundant PN-2/APP and PN-1 working at the level of factor XIa and thrombin, respectively, could function in concert as major

intracerebral anticoagulants. Alterations in the levels or functional properties of PN-2/APP and/or PN-1 could lead to coagulation disorders in the brain.

SUMMARY

The amyloid β -protein precursor (APP) is the parent molecule to the amyloid β -protein which is a major constituent of neuritic plaques and cerebrovascular deposits in Alzheimer's disease (AD). The protease inhibitor, protease nexin-2 (PN-2), is the secreted form of APP that contains the Kunitz protease inhibitor (KPI) domain. We reported that the predominant isoform of APP in human brain contains the KPI domain and is thus PN-2. Quantitation of PN-2/APP in various tissues revealed that it is primarily found in brain. Circulating blood platelets are another rich source of PN-2/APP. Platelet PN-2/APP is contained in platelet α granules and is secreted upon activation of platelets by physiological agonists. Protease inhibition measurements demonstrated that PN-2/APP is a potent inhibitor of intrinsic blood coagulation factor XIa. These findings suggest that PN-2/APP may play a role in the regulation of blood coagulation and platelets may serve as a systemic vehicle to deliver large amounts of this protein to sites of vascular injury. In addition, we propose that the rich, and relatively exclusive, investment of PN-2/APP in brain suggests that it may function locally as an intracerebral anticoagulant.

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