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Protein S: Function, Regulation, and Clinical Perspectives

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Summary

This review is intended to advance understanding of the anticoagulant function of an important plasma protein, Protein S. Despite 40+ years of research, we have not had a complete description of PS biology as it pertains to control of blood coagulation. However, the picture of PS function has become sharper with the recent discovery of FIXa inhibition by PS. Hemostasis mediated by PS now includes regulation of FIXa activity alongside the cofactor activities of PS in the TFPI/APC pathways. In addition, the direct inhibition of FIXa by PS suggests that PS, particularly a small derivative of PS, could be used to treat individuals with PS deficiencies or abnormalities that cause thrombotic complications.

Structured Abstract

Purpose of review—Protein S (PS) is an essential natural anticoagulant. PS deficiency is a major contributor to acquired hypercoagulability. Acquired hypercoagulability causes myocardial infarction, stroke, and deep vein thrombosis in millions of individuals. Yet, despite its importance in hemostasis, PS is the least understood anticoagulant. Even after 40 years since PS was first described, we are still uncovering information about how PS functions. The purpose of this review is to highlight recent findings that advance our understanding of the functions of PS and explain hypercoagulability caused by severe PS deficiency.

Recent findings—Protein S has long been described as a cofactor for Activated Protein C (APC) and Tissue Factor Pathway Inhibitor (TFPI). However, a recent report describes direct inhibition of Factor IXa (FIXa) by Protein S, an activity of PS that had been completely overlooked. Thrombophilia is becoming a more frequently reported disorder. Hereditary PS deficiency is an anticoagulant deficiency that results eventually in thrombophilia. In addition, PS deficiency is a predisposing factor for venous thromboembolism, but an effect of PS deficiency in arterial thrombosis, such as arterial ischemic stroke, is uncertain. Plasma PS concentration decreases in pregnant women. Inherited thrombophilias are important etiologies for recurrent pregnancy loss, and anticoagulation therapy is of benefit to women with recurrent pregnancy loss who had documented only PS deficiency.

Hypoxia is a risk factor for venous thromboembolism (VTE), and hypoxia downregulates plasma PS level. Importantly, COVID-19 can lead to hypoxemia because of lung damage from IL6-driven inflammatory responses to the viral infection. Because hypoxia decreases the abundance of the key

anticoagulant PS, we surmise that the IL6-induced cytokine explosion combined with hypoxemia causes a drop in PS level that exacerbates the thrombotic risk in COVID-19 patients.

Keywords

Protein S; Factor IXa; COVID-19; Hypercoagulability; Hypoxemia

Introduction

Protein S (PS) is a γ -carboxylglutamate-containing plasma glycoprotein synthesized and secreted primarily by hepatocytes (1), endothelial cells (2), and Leydig cells (3). About 2.5% of circulating PS exists within platelet α -granules (4, 5); platelet PS is derived exclusively from expression within megakaryocytes (6).

Protein S is encoded by a single *PROS1* gene in humans (7). Human *PROS1* is located on chromosome 3; the gene contains 11 introns (8). Protein S was the first blood coagulation protein mapped to a human chromosome (9). Protein S is synthesized as a 676 amino acid precursor protein that is processed to a mature multimodular protein of 635 amino acids. The mature protein is composed of a phospholipid binding γ -carboxylglutamic acid domain (GLA), a thrombin sensitive region, four Epidermal Growth Factor (EGF)-like domains, and a sex hormone binding globule (10-12) that contains two laminin G (LG)-type domains (13). Three different types of post-translationally modified amino acid residues are found in PS, 11 γ -carboxylglutamic acid residues in the GLA domain, a β -hydroxylated aspartic acid in the first EGF-like domain, and a β -hydroxylated asparagine in each of the other three EGF-like domains (Figure 1)(14). Gamma-carboxylation, the most-studied posttranslational modification, is essential for PS function.

Protein S is a multifunctional protein at the intersection of blood coagulation, inflammation, and other cellular processes. Plasma PS concentration is 350 nM, 60% of which is bound to complement component 4 binding protein (C4BP) (15); the other 40% is termed as 'free' (16). Both forms of PS function as an anticoagulant; in addition, the C4BP-bound form of PS minimizes the effect of inflammation (13). Lastly, PS promotes efferocytosis (clearance of apoptotic cells by phagocytosis) through the TAM (Tyro-3, Axl and Mer) family of protein kinase receptors (17).

Protein S is a vital anticoagulant because homozygous PS deficiency causes severe thrombosis in newborns (18) and defects in vascular system development that may be fatal (19). However, despite 40 years of research, there was a lack of clarity about how PS exerts its anticoagulant activity. In hindsight, clarity was impaired because several proposed activities of PS as an anticoagulant were confounded by artifactual activity of PS multimers that form at low phospholipid concentration. More recent studies performed with physiological amounts of phospholipids have eliminated the effects of multimers and suggest three disparate functions of PS: 1) PS is a cofactor for Activated Protein C (APC) in inactivation of FVa and FVIIIa (20, 21), 2) PS is a cofactor for Tissue Factor Pathway Inhibitor (TFPI) in accelerating the inhibition of FXa (22, 23), and 3) PS binds and inhibits the function of FIXa in producing FXa from FX (24, 25).

Protein S Functions:

Protein S is a key negative regulator of blood coagulation, and it has a significant function in complement activation pathways (20). Protein S deficiency and dysfunction cause life-threatening thrombotic conditions, including neonatal purpura fulminans (18). Currently, the consensus is that PS inhibits coagulation either independently or by enhancing the anticoagulant function of Activated Protein C (APC).

Cofactor for APC

Protein S acts as a cofactor for APC by enhancing the inactivation of activated coagulation factors Va and VIIIa (20, 21, 26). Protein S amplifies APC's inactivating function by enhancing cleavage at Arg 306 of FVa (27, 28). The EGF1 domain residue Asp95 and the GLA domain residue E36 of PS are crucial for the APC cofactor function of PS (29).

There are two APC-independent functions of PS that stood out once studies were performed with appropriate phospholipid concentrations and the results were confirmed *in vivo*.

Cofactor of TFPI

Protein S acts as a cofactor of Tissue Factor Pathway Inhibitor by stimulating the inhibition of FXa by TFPI by 4~10-fold (23, 30, 31). The Kunitz domain 3 of TFPI binds the LG1 domain of PS (32-34), and a direct TFPI α -protein S interaction is required for the enhanced inhibition of FXa.

Inhibition of Factor IXa Activity

In 2012, another anticoagulant function of PS was established by the finding that PS directly binds and inhibits the procoagulant FIXa in the presence and absence of FVIIIa (24, 25). This direct inhibitory activity suggested a novel regulatory function of PS in blocking thrombin generation maximally in the presence of FIXa and FVIIIa (24, 25, 35, 36).

Plautz et al., showed that PS binds and inhibits FIXa *in vivo* by associating with the K132, K126, and R170 residues in the FIXa heparin-binding exosite; this interaction with PS regulates thrombin generation by inhibiting the intrinsic Xase complex (35). FIXa and PS also coimmunoprecipitated from plasma, corroborating their interaction in a physiological setting (25, 35). Importantly, Hemophilia B mice infused with a FIXa double mutant, K132A/R170A, that had full activity but could not bind to PS demonstrated an accelerated rate of fibrin clot formation compared with infusion of wild type FIXa (35). Crucially, disruption of the interaction between PS and FIXa caused an increased rate of thrombus formation in Hemophilia B mice (35). These data established that interaction of PS with the FIXa is an important APC-independent regulatory mechanism for the propagation phase of coagulation (35).

Regulation of Protein S:

Protein S deficiency is a genetic disorder caused by mutation in the *PROS1* gene. Protein S deficiency has been difficult to study because of the peculiar PS biology. PS has an anticoagulant function but no enzymatic activity, and PS interacts with plasma components

that operate in both hemostasis and inflammation. Heterozygous PS deficiency is an autosomal dominant condition because half the normal expression of PS increases the risk of thrombosis, as judged from familial and population studies. Heterozygous individuals have an increased risk of developing blood clots in the legs (deep venous thrombosis); the clots can break loose and travel to the lungs, an event that is termed pulmonary embolism. Rarely, two abnormal PS genes are inherited (congenital) that can cause widespread small clots and a life-threatening disease, purpura fulminans (18). Furthermore, murine PS knockouts are embryonic lethal (22) (23). However, a heterozygous genotype is not directly associated with lethality, unlike a homozygous genotype. Heterozygous PS deficiency is associated with increased risk of venous thrombosis (37-39) and pulmonary embolism (40).

Occasionally, PS deficiency may be acquired because of conditions such as kidney disease, malignancy, pregnancy, or the use of oral (estrogen-based) contraceptives (41, 42). Patients with chronic kidney disease who undergo hemodialysis experience hypercoagulability. There are reports that PS deficiency is a cause of hypercoagulability in these patients (43). Protein S is also an important signaling molecule that binds and activates a family of tyrosine kinase receptors known as TAM (Tyro3, Axl and Mer) receptors. TAM receptors are often upregulated in cancers. We recently reported that PS suppresses the growth of pancreatic cancer cells in vitro and that *PROS1* (PS) gene expression is associated with better survival in pancreatic cancer (44). Acquired PS deficiency occurs in women with high levels of estrogen, either because of pregnancy or from the use of estrogen containing oral contraceptive pills (45-47). There was a report that estrogen represses PS production because the estrogen receptor (ER α) downregulates the activity of the PS gene promoter (46). However, the details of this mechanism are not known. Sickle cell anemia is another seldom recognized cause of acquired PS deficiency (48). Patients with sickle cell disease bear a high risk of developing thromboembolic disorders, and PS deficiency was one of the major reasons for thromboembolism (49). It has been hypothesized that acquired PS deficiency in sickle cell disease is due to direct binding of PS to sickling red cells that express increased surface phosphatidylserine (50). In addition, PS expression is depressed in hypoxic conditions (51). Stabilization of HIF1 α in the liver, a normal response to hypoxia, is associated with reduced PS expression and plasma level, which, in turn, increases the likelihood of thrombosis. HIF1 α is a well-documented, general transcriptional activator, and Pilli et al. (51) showed that HIF1 α directly downregulates PS expression (51). There is a report of a rare case of PS deficiency associated with assisted reproductive treatment with ovarian hyperstimulation syndrome (OHSS), following right neck venous thromboembolism (1). Thromboembolism is a life-threatening complication among women with OHSS; therefore, thrombophilia workup should be considered for women with thrombotic events.

Mutations of Protein S

Almost 300 different mutations have been identified in *PROS1*, spread over all exons, the promoter region, and within several introns (54). Most alterations are missense mutations, but several other types of mutations have been reported.

Here, we describe further three *PROS1* mutations. We selected these mutations because they are unique and have been observed only recently. The molecular genetic analysis of

a 23-year-old patient who experienced venous thrombosis showed a homozygous missense mutation, G664>A, in exon7 of *PROS1* (52). This mutation results in replacement of glycine by arginine at codon 222, and the mutation causes a reduction in PS level (52). This alteration was the first reported mutation of PS codon 222 (52). Hereditary PS deficiency caused by a mutation/deletion in *PROS1* is autosomal dominant, which means that an individual who inherits only one mutant *PROS1* gene nevertheless has an increased chance of developing symptoms of PS deficiency. A recent report describes a heterozygous deletion of eight base pairs (Leu313Ser) in *PROS1* exon 9 and a missense variant (Ser538Phe) in two different patients. These mutations resulted in PS deficiency and severe thrombophilic diathesis (53). Further, a rare C39>T mutation in the *PROS1* 5'UTR was reported to create a new upstream translation initiation codon that caused PS deficiency and venous thromboembolism (54).

Clinical Considerations:

Protein S – Clinical Perspectives

PS deficiency manifests clinically as thrombophilia, an imbalance of coagulation factors that leads to abnormal blood clotting. This pathophysiological disturbance puts patients at risk for disseminated intravascular coagulation, potentially causing disabilities such as cerebrovascular injury or chronic venous thromboembolism (VTE). We also consider unique case studies to discuss other recent developments of PS clinical research concerning pregnancy and management of PS deficiency. Lastly, Malas et al. reported that approximately 31% of COVID-19 patients admitted to the ICU experienced at least one thromboembolic event (55). Thus, it is important to discuss the possible correlation between COVID 19 morbidity and PS deficiency. Although there are certainly other factors that contribute to thrombophilia in severe COVID patients, when we consider the intersection of hypoxia, inflammation, and severity of symptoms associated with decreased PS, we suggest that Protein S likely contributes to thrombophilia in COVID-19 patients.

Estrogen Induced Acquired Protein S Deficiency

In general, estrogen-containing medications, such as oral contraceptives or hormone replacement therapy, induce a prothrombotic state that results from acquired PS deficiency. Pregnant women are at high risk for thromboembolic events because of estrogen-associated hypercoagulability [51]. Estrogen level steadily increases during pregnancy, which elevates the risk for clotting events [52]. Hereditary PS deficiency has long been a contraindication for contraceptive use because of the known effect of estrogen on downregulation of PS (46). The molecular mechanism of estrogen-induced PS deficiency has not yet been clearly defined.

Management of PS deficiency

Interestingly, a diminished PS level appears to correlate with lifestyle modifications to combat obesity and insulin resistance, despite overall reduction in thrombophilia with a decrease in prothrombotic factors. Recall that hypoxia can induce PS deficiency (51); thus, hypoxia secondary to intense exercise may be a factor in the negative correlation between lifestyle change and PS level. As for pharmaceutical management of PS deficiency, direct

oral anticoagulants (DOAC) have been found to be efficient in treatment and prevention of VTE in patients with mutations in the *PROS1* gene (56, 57). Traditionally, heparin or vitamin K antagonists, such as warfarin, have been used to prevent VTE; however, the DOACs described in these studies, rivaroxaban and apixaban, may offer more predictable outcomes in the case of PS deficiency. Another benefit of DOACs is that they offer fixed dosing, limited drug-drug interactions, and the potential to reverse anticoagulant activity with the administration of Andexanet alfa (58).

Protein S deficiency is associated with venous thromboembolism; however, it is unclear whether PS deficiency promotes arterial thrombosis, such as arterial ischemic stroke. Fearon et al. described a 79-year-old woman with PS deficiency due to a Trp149Cys mutation; this individual had an arterial thromboembolic event (59). Similarly, Naghavi et al. reported an effect of PS deficiency on arterial ischemic stroke (60). This group reported further that rivaroxaban was an effective option for prevention of recurrent arterial ischemic stroke caused by PS deficiency (60). Lastly, Ohashi et al. described a 59-year-old patient with ischemic stroke caused by a hereditary PS deficiency; the patient was treated with apixaban, another direct oral anticoagulant (57).

Further, Lou et al. described two patients with hereditary PS deficiency who had deep vein thrombosis even after warfarin therapy for two years (56). However, after the patients were treated with 20 mg/day rivaroxaban, they did not have any thromboembolic events even after one year follow up (56). A consensus should be reached for moving towards new oral anticoagulants such as rivaroxaban and enoxaparin-VKA, which can preserve PS level during clotting events.

COVID-19

Infection by SARS-CoV-2 not only results in hyperactivity of the immune system by a cytokine storm, but also the virus causes severe hypoxemia. We know hypoxia downregulates PS expression (51). Therefore, one may speculate that downregulation of PS during COVID-19 contributes to a hypercoagulable state (61). Several studies support the correlation between VTE in COVID-19 patients and decreased PS level (62) (63). Wood et al. showed that free, but not total, PS level was depressed in severe COVID-19 patients (64). Unpublished data from our group supported the findings of Wood et al. (64).

Interestingly, Ruzicka et al. (65) hypothesized that the papain-like protease (PLpro) of SARS-CoV-2, the causative agent of COVID-19, may contribute to the coagulation and thrombotic dysregulation by dissolute cleavage of Protein S. A review of the three N-terminal cleavage sites recognized by the PL proteases of SARS viruses revealed a highly conserved LXGG motif. A BLAST search using the 10 amino acids that span the nsp3/4 cleavage site of CoV2 (IALKGG/KIVN) returned an eight amino acid “hit” within *PROS1*, including the core LXGG sequence (65). The “X” in the CoV2 polyprotein, a lysine, is replaced by a highly conserved arginine in Protein S. Therefore, it is rational to suggest that downregulation of PS during severe COVID 19 occurs by PLpro cleavage of PS.

Whether or not PS deficiency has a significant effect in the development of VTE, this research topic has potential for elucidating the mechanism of hypercoagulable states in

SARS-infected patients. Investigators have identified reduced vitamin K levels as a potential modifiable risk (66) factor in patients with COVID-19, which may be pertinent because PS is a vitamin K-dependent protein. Measurement of free PS level in the circulation is not currently part of the standard blood work-up for COVID-19 patients, but such measurements are warranted, and they will be informative for treatment.

Conclusion

The hypercoagulable state associated with congenital deficiency of PS demonstrates the importance of this protein in regulation of hemostasis. Protein S acts as an anticoagulant by performing three different functions: 1) Cofactor for APC, 2) Cofactor for TFPI, and 3) Inhibitor of FIXa. Downregulation of FIXa function by PS has not been previously considered. The allure of this inhibitory property of PS is related to the fact that mutant forms of FIXa that lack the binding site for PS would then exhibit enhanced procoagulant activity, which has translational potential regarding therapies for FIX upregulation or PS deficiency. Pregnancy, estrogen containing contraceptives, hypoxia, and most probably COVID-19 infection are factors that cause acquired PS deficiency, which, in turn, causes VTE. Measurement of PS in plasma must be considered as a routine for COVID-19 and during pregnancy, so that initial symptoms due to PS deficiency can be treated in a timely manner with DOACs such rivaroxaban.

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Key Points:

- Protein S enhances the activities of Activated Protein C and Tissue Factor Pathway Inhibitor and inhibits the activity of activated Factor IX. Thus, Protein S occupies a central position in regulation of hemostasis.
- Genetic deficiencies of Protein S, and acquired deficiencies of Protein S induced by, e.g., estrogen and hypoxia, are linked to a variety of venous and arterial thrombotic complications, including deep vein thrombosis and stroke.
- SARS COV2 virus-induced Protein S deficiency may be responsible for the often fatal hypercoagulated state of COVID-19 patients.

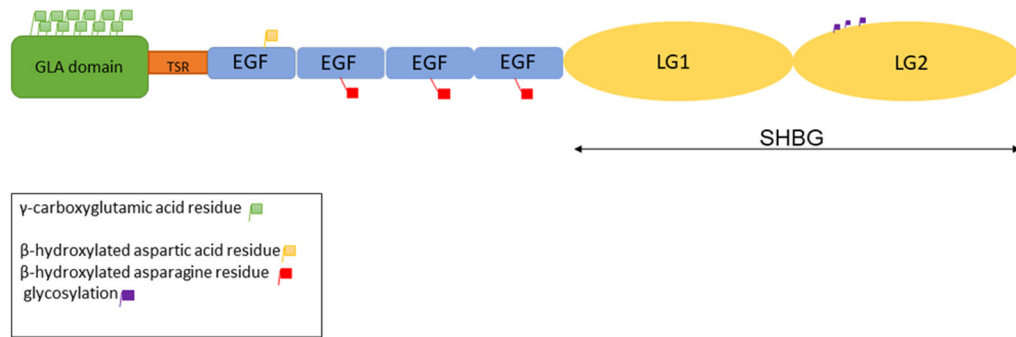


Figure 1-
The cartoon showing different domains of Protein S and sites for post translational modifications in different domains.