



Coagulation and Skin Autoimmunity

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Several lines of evidence indicate that the immune system, inflammation, and coagulation are simultaneously activated in autoimmune and immune-mediated skin diseases. Pro-inflammatory cytokines such as interleukin-6 and tumor necrosis factor-alpha induce the expression of the main initiator of coagulation, i.e., tissue factor. The proteases of coagulation in turn act on protease-activated receptors inducing the expression of various pro-inflammatory cytokines triggering inflammation. The cross-talk among immune system, inflammation, and coagulation amplifies and maintains the activation of all three pathways. This review focuses on three skin disorders as chronic spontaneous urticaria (CSU), angioedema, and bullous pemphigoid (BP), in which the relationships among the three systems have been investigated or their clinical consequences are relevant. Markers of thrombin generation, fibrinolysis, and inflammation have been reported to be increased in the plasma during flares of CSU and angioedema, as well as in the active phase of BP, with the marker levels reverting to normal during remission. The coagulation activation seems to be important only at local level in CSU and angioedema while both at local and systemic levels in BP which is the only condition associated with an increased thrombotic risk. The prothrombotic state in autoimmune skin diseases raises the question of the indication of anticoagulant treatment, particularly in the presence of other cardiovascular risk factors.

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INTRODUCTION

Immune system, blood coagulation and inflammation strictly interact in providing a defense against a variety of potentially injurious stimuli, such as infections and tissue damages (1). The molecular mechanisms of this interaction have been largely elucidated. Indeed, pro-inflammatory cytokines, like interleukin 1 (IL-1), IL-6, and tumor necrosis factor alpha (TNF- α), induce the expression of tissue factor (TF) the main initiator of blood coagulation whereas downregulate the natural anticoagulants such as antithrombin, protein C, and TF pathway inhibitor (2). On the other hand, the coagulant mediators (FVIIa, FXa, and FIIa) in turn act on protease-activated receptors (PAR) inducing the expression of pro-inflammatory cytokines (3, 4). The relationships between the activation or dysfunction of the immune system and the coagulation system are evident in systemic autoimmune or immune-mediated diseases including lupus erythematosus (5, 6), rheumatoid arthritis (7, 8), and inflammatory bowel diseases (9, 10) which show an increased risk of thrombosis. A few studies suggested the involvement of blood coagulation also in some immune-mediated skin disorders whose aspects will be analyzed in the present review.

Immune-mediated inflammatory skin diseases comprise a group of heterogeneous chronic disorders that share similar immune-mediated pathogenic mechanisms as well as genetic susceptibility. Although their specific etiologies remain often unknown, all are recognized to involve dysregulation of the immune system, including an over-expression of the pro-inflammatory cytokines. In autoimmune and immunemediated skin disorders, the cross-talk between inflammation and coagulation creates a self-refueling loop which amplifies and sustains the activation of both systems (3). Growing evidence suggests that this has both local and systemic implications.

The aim of the present study is to focus on several skin disorders in which the relationships between immune response, inflammation and blood coagulation have been investigated and/or their clinical consequences are relevant such as chronic spontaneous urticaria, angioedema, and bullous pemphigoid.

CHRONIC SPONTANEOUS URTICARIA

Urticaria is a common skin disease characterized by short-lived swellings, called wheals, which resolve in <24 h. Urticaria can be classified according to duration and etiology as acute or chronic (11). Chronic urticaria is defined as urticaria with or without angioedema lasting more than 6 weeks and can be further classified according to whether it is inducible or not into chronic inducible urticaria or chronic spontaneous urticaria (CSU).

Despite the great research effort of the last 20 years, etiology and pathogenesis of CSU remain largely unclear. However, there is growing evidence that different biologic systems including autoimmunity (12–14), inflammation (15–18), coagulation (19), and auto-allergy (20–23) are involved in the mechanisms leading to mast cell and basophil degranulation and hence to wheal formation (**Figure 1**).

The autoimmune mechanism is based on the presence of circulating histamine-releasing IgG autoantibodies directed against either the high-affinity IgE receptor (FcERI) on both mast cells and basophils (in most cases) or membrane-bound IgE (in a minority of patients) (12, 13, 24). In a variable proportion (30-60%) of patients with active disease, the intradermal injection of autologous serum (autologous serum skin test, ASST) causes a wheal-and-flare reaction (19). This phenomenon mirrors the activation of mast cells and basophils as well as of the complement cascade by these autoantibodies. Experimental and clinical findings have supported an autoimmune origin in about 30-50% of patients affected with CSU (12, 13). Also other autoantibodies are involved in CSU pathogenesis such as autoantibodies to CD23, the low-affinity IgE receptor (FceRII), which contribute to eosinophil activation and mast cell degranulation (14, 25). The possible autoimmune nature of CSU is further supported by the association with other autoimmune diseases, notably thyroiditis, and the increased frequency of HLA DRB1*04 (DR4) (26).

Inflammation is involved in the pathophysiology CSU, as demonstrated by the increase in different markers of inflammation, such as matrix metalloproteinase, serum C-reactive protein, IL-6, IL-6 soluble receptor, eosinophil cationic

protein (ECP), TNF- α , complement, and others, which are related to CU activity (15–17, 27). On the contrary, adipokines that affect immune responses and exert an anti-inflammatory effect, such as lipocalin 2, have a negative association with CU activity (18).

Coagulation involvement in the pathogenesis of CSU is supported by many lines of evidence (28-33). The original observation that the autologous plasma skin test (APST) may score positive in some ASST-negative patients (29) has given rise to investigation of the coagulation system in CSU patients. In a study that dates back over 10 years, some of us found that CSU patients have elevated plasma levels of prothrombin fragment F1+2, suggesting thrombin generation (29). In subsequent studies, we found that CSU patients show an activation of the tissue factor pathway of coagulation cascade by activated eosinophils (30). Immunohistochemical experiments showed tissue factor expression by eosinophils present in the inflammatory infiltrate of CSU skin lesions (31). These data highlight the relevance of eosinophils in CSU as a source of tissue factor, in accordance with studies showing that eosinophils store tissue factor and transfer it to their cell membrane during activation (34). Eosinophils may be activated directly by autoantibodies directed against the FcERII CD23 antigen (14) or secondarily to the activation of mast cells by anti-FceRI and anti-IgE autoantibodies (24). The activation of the tissue factor pathway of coagulation results, in turn, in the generation of thrombin which, in experimental models, has been shown to induce edema through an increase in vascular permeability directly by acting on endothelial cells (35) and indirectly by generating C5a (36, 37), by triggering mast cell degranulation (38-41) and by releasing inflammatory mediators (42, 43). Proinflammatory cytokines, such as IL-6 and TNF-a, induce the expression of tissue factor (3, 44), sealing the mutual activation of the two systems, i.e., coagulation and inflammation. The complexes between tissue factor, activated factor VII (FVIIa) and FVa+FXa also activate mast cells via PAR-2, thus amplifying the activation of these cells in CSU (40). On the other hand, mast cell-derived tryptase can induce thrombin generation through a direct activation of prothrombin (45), thus constituting an amplification loop. It must be taken into account also an opposite effect of mast cells on inflammatory response. In fact, recent studies showed that mast cells can also limit thrombin-induced immediate skin inflammatory responses by releasing substances able to reduce the activity of thrombin, for example proteases, like mast cell protease 4 (MCPT4) (46). During severe exacerbations of the disease, CSU patients show marked increase of plasmatic markers of thrombin generation, like prothrombin fragment F1 + 2, FVIIa and thrombin-antithrombin complex (47–49), as well as of fibrinolysis (24, 31, 50). The increase of D-dimer, marker of fibrinolysis, is associated with disease severity (24) and its level resulted higher when compared with that of patients affected with psoriasis (51). Interestingly, a study detected the increase of D-dimer and fibrinogen/fibrin degradation products (FDP), produced by fibrinolysis of either stabilized or non-stabilized fibrin, in patients with CSU, but not in healthy individuals or in patients with inducible type of urticaria (52). Finding that the increase in plasma markers of thrombin generation



and fibrinolysis parallels plasma C-reactive protein (CRP) levels further remarks the close link between coagulation activation and inflammation in CSU pathogenesis (53). The interplay between inflammatory and coagulation/fibrinolysis factors in CSU may lead to maintenance and amplification of urticarial inflammation. Both the activation of coagulation and fibrinolysis decrease till complete normalization during remission and biomarker levels return to normal values (31, 49, 54, 55). A limited number of studies investigated the role of platelets in CSU, with no univocal data (56). Platelet activation was found to occur in some skin inflammatory disorders, such as atopic dermatitis and psoriasis (57, 58). Based on these findings, platelets are regarded as a possible link between chronic inflammatory and pro-coagulant states (59). It may be hypothesized that platelet activation may be one of the triggering factors of the coagulation cascade in CSU too, leading to the activation of histamine-releasing effector cells. However, current evidence indicates that the simple determination of platelet indices is not reliable and lacks useful implications in the clinical practice (56).

The activation of the coagulation cascade in CSU pathogenesis may have clinical effects not only in eliciting skin lesions, namely wheal eruption, but also at systemic level. The most important potential clinical consequence of the hypercoagulable state in CSU patients is an increased thrombotic risk (25). In spite of this, patients with CSU are not reported to have an

increased risk for thrombotic events. A possible explanation is that the activation of coagulation occurs mostly extravascularly and can be efficiently counteracted by coagulation inhibitors and fibrinolysis. Other potential clinical implications of the recognized involvement of coagulation in the pathogenesis of CSU are the possible therapeutic efficacy of pharmacological agents that interfere with the coagulation pathway. A few reports indicate that heparin, an anticoagulant which potentiates the effect of antithrombin, may be effective in treating CSU (60, 61). An anticoagulant/anti-fibrinolytic therapy, namely nadroparin and tranexamic acid, has been found effective in some patients with refractory CSU and elevated D-dimer levels (62). Moreover, warfarin, an oral anti-vitamin K drug, induced a good response in patients with CU unresponsive to antihistamines in a double-blind, placebo controlled study (63). With reference to further clinical implications of the activation of coagulation and fibrinolysis, elevated D-dimer plasma level has been found to be associated both with a poor response to antihistamines (64) and with a limited response to cyclosporine treatment (65). It is worthy of note that D-dimer plasma levels were observed to parallel the clinical response to omalizumab treatment, dropping in responders and remaining unchanged in non-responders (20, 66). Furthermore, elevated D-dimer plasma levels seem to be a predictive marker of prompt and complete response to the anti-IgE monoclonal antibody (67). It may be assumed that

in a subset of CSU patients, the activation of the coagulation cascade has a key role in the pathogenesis of this disorder. In a recent study, the treatment with omalizumab induced a significant decrease in WBC count, platelet count, neutrophil count, ratio of platelet to lymphocyte (PLR), ratio of neutrophil to lymphocyte (NLR) and CRP level and a significant increase in mean platelet volume (MPV) and eosinophil count (68). Taken together, these data suggest that in CSU patients omalizumab may be effective not only antagonizing IgE but also affecting a number of other pathways (69). In particular, it can exert inhibitory effects on inflammation and coagulation cascade, further confirming the strict interaction between immunity, inflammation, and coagulation.

The "autoallergic" mechanism, which has been recently identified, is mediated by specific IgE to different allergens, including double-strained DNA (21), thyroperoxidase (22), IL-24 (23), tissue factor, and thyroglobulin (20). IgE anti-autoallergens seem to cause "autoallergic" mast cell degranulation. In particular, the elevated specific IgE antibodies to tissue factor have been demonstrated to be functionally able to mediate the release of leukotriene C4 by tissue factor-stimulated peripheral basophils (20). Tissue factor, the main initiator of coagulation, is overexpressed in CSU lesional skin (32) where it can activate coagulation and is also accessible to mast cell-bound IgE. Thus, IgE anti-tissue factor could cause "autoallergic" mast cell degranulation, linking the autoallergic and coagulation activation pathways. The presence of autoreactive IgE against several targets in patients with CSU might explain, at least in part, the clinical success of the humanized monoclonal anti-IgE antibody omalizumab.

ANGIOEDEMA

Angioedema refers to a circumscribed, non-itchy edema of the subcutaneous tissues mostly involving lips, face, neck and extremities and/or submucosal tissues affecting oral cavity, larynx and gut (70). It typically lasts from many hours to 3 days before the tissue returns to normal. Angioedema is lifethreatening if it occurs in the upper airways and can be very painful if it occurs in the gastrointestinal tract (70). Wheals and angioedema often coexist but may also present alone (71). Most of these conditions are mainly mediated by histamine and other pro-inflammatory mediators released by mast cells and basophils (72). Overall, angioedema can be caused by allergies, inherited or acquired deficiencies of C1-inhibitor protein, or drug reactions. In many cases its pathophysiology remains undetermined (54). Among the vasoactive agents involved in mediating swelling attacks in angioedema, histamine, released by mast cells or basophils, has a prominent role, especially in allergic reactions and in cases of associated urticaria (73). Antihistamines are effective agents in these forms. However, there are some subtypes of angioedema that do not respond to antihistamine, such as angioedema due to C1-inhibitor deficiency, angioedema due to ACE inhibitors and a rate of idiopathic angioedema (54). Bradykinin is a recognized mediator of the increased vascular permeability in angioedema. Bradykinin is a vasoactive peptide acting on specific receptors such as B2 receptors that are constitutively present on the endothelium and B1 receptors that are expressed during inflammatory reactions. Bradykinin receptor signaling induces peripheral vasodilatation, enhancement of vascular permeability and subsequent vascular leakage. Bradykinin is the end-product of the contact activation system. This enzymatic cascade circulates in the plasma and consists of factor XII (FXII), plasma prekallikrein (PPK), and high molecular weight kininogen (HK). This system is linked to the intrinsic coagulation system via factor XI (FXI). Animal models showed that production of bradykinin in plasma is \sim 50% dependent on FXII (74). This suggests the presence of alternative pathways generating bradykinin *in vivo*. The endothelial cell-derived factors *heat shock protein 90* (HSP90) and prolylcarboxypeptidase may be involved in FXII-independent release of bradykinin (75).

Bradykinin was first identified as a mediator for hereditary angioedema (HAE) (76, 77). The genetic form of C1-inhibitor deficiency is due to mutations in one of the two alleles of the C1-inhibitor gene, which leads to either reduced plasma protein levels [hereditary angioedema [HAE] type I] or normal levels but reduced protein function (HAE type II) (78). HAE type III is a rare subtype of HAE that is not connected with C1-inhibitor deficiency but with a dysregulation of the contact (plasma kallikrein-bradykinin) system (79). In a subset of patients with HAE with normal C1-inhibitor, a gain-of-function mutation in FXII has been identified (80). The acquired form of C1-inhibitor deficiency is known as acquired angioedema and is due to C1inhibitor consumption associated with the presence of anti-C1inhibitor autoantibodies and/or lymphoproliferative disorders (81). Bradykinin involvement in angioedema pathogenesis is not limited to hereditary forms. Bradykinin is a key mediator in patients with acquired C1-inhibitor deficiency due to underlying auto-immune or lymphoproliferative diseases (82) as well as in those treated with anti-hypertensive drugs that inhibit bradykinin breakdown, such as angiotensin-converting enzyme inhibitors (83). It is also possible that bradykinin may play a supportive role in forms of angioedema that are currently classified as "histaminergic" (84, 85). C1-inhibitor deficiency involves different biological systems that interplay during angioedema attacks. In fact, C1 inhibitor is a serine protease inhibitor (serpin) that blocks the activity of (i) C1r and C1s in the complement system, (ii) factor XII and kallikrein in the contact system, (iii) factor XI and thrombin in the coagulation system, and (iv) tissue plasminogen activator and plasmin in the fibrinolytic system (86). A deficiency in C1 inhibitor results in the hyperactivation of the contact system (87, 88), which leads to the generation of bradykinin. Furthermore, C1-inhibitor deficiencies activate the complement (89) as well as coagulation (90, 91) and fibrinolysis systems (92, 93). Thus, the unregulated activation of coagulation leads to the generation of thrombin, which can potentiate the vasoactive effect of bradykinin both directly (35, 94) and by releasing fibrinopeptides, which enhance the effects of kinins (95). It may be argued that in angioedema due to C1-inhibitor deficiency, thrombin acts synergistically with other vasoactive substances released by the concomitant activation of contact system, complement, or mast cells leading in turn to increased vasopermeability. Plasmin has been assumed to act as a main trigger for contact system activation and bradykinin production in the pathogenesis of most forms of

HAE and specific forms of non-hereditary angioedema (96, 97). Indeed, several clinical observations supported the relevance of plasmin as a natural FXII activator and evidence for plasmindependent bradykinin generation as a cause of angioedema during treatment with fibrinolytic agents is accumulating (73). Consistent with this hypothesis, complexes of plasmin with its inhibitor a2-antiplasmin are elevated during attacks of HAE due to C1-inhibitor deficiency, as are the levels of markers of ongoing fibrinolysis, like D-dimer (98). Similarly to CSU, in patients with angioedema there is lack of prothrombotic features (55, 99). Therefore, it may be assumed that in angioedema, factor XII-driven contact system starts inflammatory mechanisms via the bradykinin-producing kallikrein-kinin system, without procoagulant effects (35). Alternatively, it has been proposed that the extreme vascular leakage may move the plasma coagulation factors into the extravascular space, triggering coagulation in the absence of vascular injury or intravascular thrombi (73). Therapy targeting the contact system has been successful in HAE, strongly supporting that angioedema is mediated via bradykinin production (73). Anti-fibrinolytic therapy, mainly tranexamic acid, has been used as prophylactic therapy for HAE attacks for some decades (100).

BULLOUS PEMPHIGOID

Bullous pemphigoid (BP) is an autoimmune blistering disease that mainly affects the elderly and carries a high risk of death (101, 102), mostly due to sepsis and cardiovascular events (103). Either comorbidities or immunosuppressive treatments seem to concur on BP mortality (104). Albeit rare, it is the most common autoimmune blistering disease in Western countries (105). It involves the skin and rarely the mucous membranes and is characterized by the presence of tense blisters usually surrounded by erythematous-edematous, urticaria-like plaques (106, 107). BP has a chronic relapsing evolution, with spontaneous exacerbations and remissions (108). The pathophysiology of BP is linked to the presence of circulating IgG autoantibodies directed against two antigens, BP180 and BP230, which are components of junctional adhesion complexes called hemidesmosomes that promote dermoepidermal cohesion (109, 110) with the contributory role of other pathomechanisms (Figure 2). BP180 (type XVII collagen) autoantibodies have been demonstrated to be directly pathogenic by triggering an inflammatory cascade. Inflammatory cells, particularly autoreactive T cells and eosinophils, produce and release a number of cytokines and soluble factors that amplify and maintain tissue damage, which ultimately leads to subepidermal blister formation (111). During acute phase of BP, autoreactive T helper (Th) 1, Th2, and Th17 lymphocytes cooperatively play a role in the development of the disease process (112-114). Although the role of Th17 cells in human BP is not completely defined, recent experimental data showed that IL17Adeficient mice are protected against autoantibody-induced BP, and pharmacological inhibition of IL-17A reduces the induction of BP (115). Moreover, a dysfunction of both T and B regulatory cells, whose immunosurveillance action is critical in

preventing autoimmunity, was observed in lesional skin of BP patients (116, 117). The involvement of many cellular players in the BP inflammatory process is supported by elevated serum levels of activation markers, including molecules released by T cells upon their stimulation, molecules involved in B-cell maturation, granule proteins liberated from activated mast cells, neutrophils and eosinophils and adhesion molecules indicative of neutrophil and platelet activation. Complement activation is considered to be critical for blister formation too. Indeed, complement activation by autoantibodies ex vivo as measured by the complement-fixation assay in serum was correlated with the clinical disease activity in patients with BP (118). Experimental data in murine models demonstrate that complement-dependent and -independent mechanisms coexist in blister formation. On the basis of knock out mice and pharmacological inhibition studies, the activated 5th component of complement (C5a) appears to be involved, and its receptor 1 (C5aR1) seems to be important during the early phase of the disease while its receptor 2 (C5aR2) seems to be protective. Once the skin inflammation has fully developed, release of reactive oxygen species and proteases from neutrophils and macrophages may become independent of complement (119). Several studies showed that the coagulation cascade is activated in BP and correlates with the severity of the disease (25, 120-123). In particular, coagulation activation markers have been found to be increased at both local level, i.e., in blister fluid, and systemically, i.e., in plasma (123). It has been hypothesized that the activation of blood coagulation is induced by the inflammatory response underlying BP pathogenesis. In this regard, eosinophils are highly represented in the inflammatory infiltrate of the lesional skin and their levels are often increased in peripheral blood (123-125). Elevated concentrations of secretory granules, such as eosinophil cationic protein (ECP) (126, 127) as well as increased levels of IL5, the main cytokine involved in eosinophil biology (128, 129), and IL16, a chemotactic factor for eosinophils (130), in blister fluid and sera of patients with BP confirm the involvement of eosinophils in the pathogenesis of this disease. Eosinophils are recognized to concur in BP elicitation by producing and releasing matrix metalloproteinase (MMP)-9, which plays a key role in degrading BP180 and cleaving the dermal-epidermal junction (131). Release of elastase and gelatinase, namely 92 kDa gelatinase, may further contribute to tissue damage and blister formation (132, 133). Eosinophil degranulation was demonstrated not only in fully developed blisters, but also at the earliest stages of blister development and urticarial lesions of BP (134, 135). Moreover, eosinophils have been postulated as potential intermediates between anti-BP180 IgE autoantibodies, whose pathogenic role in BP has increasingly recognized in recent years, and dermo-epidermal junction separation (136, 137). High-affinity IgE receptors (FceRI) that are highly expressed on eosinophils in BP patients could enhance the capability of eosinophils to bind IgE and thus influence their subsequent degranulation (138). In recent reports, the monoclonal anti-IgE antibody omalizumab resulted effective in treating patients affected with BP, despite ongoing high levels of anti-skin IgG antibodies (139, 140). This provides further evidence of an independent role for autoreactive IgE-mediated



soluble factors responsible for the recruitment and activation of eosinophils. In lesional skin, eosinophils produce and release metalloproteinases, elastase, and gelatinase which contribute to tissue damage. Moreover, eosinophils strongly express tissue factor, which is the main initiator of the coagulation cascade (factors VII, X, VIII, V, and prothrombin) leading to generation of thrombin. This last increases the permeability of blood vessels, amplifying the inflammatory network. Neutrophils contribute in the pathogenesis of bullous pemphigoid by releasing reactive oxygen species (ROS) and proteases. Finally, complement is activated upon binding of the pathogenic autoantibodies to their autoantigens.

inflammation in BP skin lesion development. Eosinophils are also involved in itch induction, mainly due to the release of IL-31 (141, 142). IL31 exerts a chief role in itch, by activating endothelin-1 responsive neurons and by increasing the release of brain natriuretic peptide (BNP), a central mediator of itch (143, 144). With specific reference to the activation of the coagulation cascade in BP, it correlates with eosinophilia other than severity of the disease, thus indicating that eosinophils play a pivotal role in this process (123). Immunohistochemistry showed that eosinophils strongly express TF in lesional skin, as confirmed also by co-localization studies (120, 123). Eosinophils are a major intravascular location for TF storage and exposure. On the other hand, TF facilitates the early transendothelial migration of eosinophils (34). Consistent with these findings, it may be supposed that extrinsic blood coagulation initiated via TF expressed by eosinophils contributes to local inflammation and blister formation (145). In fact, TF plays an important role in the inflammatory process, in addition to its welldocumented prothrombotic properties (34). In BP patients, increased skin expression of adhesins and MMPs has been considered an effect of TF action (146). TF is a recognized factor connecting the immune system with coagulation system. Pro-inflammatory cytokines, such as TNF and IL-6, which are increased in BP, induce TF expression (25), as found also in other, very heterogeneous inflammatory conditions (147-149). Activation of the extrinsic coagulation pathway generates thrombin that increases the permeability of blood vessels (150, 151). The presence of thrombin may play a

direct role in the pathogenesis of BP by increasing vascular permeability, thus favoring the transendothelial migration of inflammatory cells and their accumulation in the skin. Activated proteases, in turn, act on PARs and induce the expression of various pro-inflammatory cytokines, and this cross-talk between inflammation and coagulation amplifies and maintains the activation of both systems (123). It is noteworthy that BP patients have high levels of coagulation activation markers, such as prothrombin fragment F1+2 (indicating thrombin generation) and D-dimer (indicating fibrin degradation), in plasma samples other than in blister fluid (120, 121, 123). During the disease remission, blood concentration of coagulation activation markers returns to normal (49, 145). Moreover, the concentration of the prothrombin fragment F1+2 correlates with the concentration of immunoglobulins directed against the BP180 antigen (120). All together, these findings clearly indicate that in BP patients there may be an activation of coagulation also at systemic level. A recent study has shown that plasma levels of prothrombotic markers are higher in BP patients when compared with patients affected with CSU (49). In pathological states, especially in inflammatory disorders, the balance between the coagulation and fibrinolysis might be deteriorated (152, 153). It is well known that the inflammatory response inhibits fibrinolysis. The results of a previous study on a group of patients with active BP showed that fibrinolysis is inhibited, due mainly to an increase in the plasma levels of plasminogen activator inhibitor type 1 (PAI-1) activity and antigen (154). The most important clinical consequence of the hypercoagulable state related to both

coagulation activation and fibrinolysis inhibition in BP patients is an increased thrombotic risk. It has been consistently reported that the risk of thrombosis is increased in patients with BP (155, 156) and we have found an annual incidence of venous thrombosis of 8% (120), undoubtedly higher than that observed in the general elderly population (157). Langan et al. investigated the main acute medical conditions associated with BP, finding an increased risk for pulmonary embolism (158). A recent multicenter cohort study has shown that the risk of developing venous thromboembolism (VTE) is increased 4-fold in patients with BP as compared with the general population of same age and sex (159). More specifically, the VTE risk increased up to 15 times during the acute phase of the disease, thus indicating a close link between the inflammatory state occurring in active BP and thrombosis. Interestingly, the VTE risk dropped to 1.5 times during clinical remission. This latter finding indicates that a good control of the disease also impacts the occurrence of thrombotic events. This study highlighted in a large cohort of BP patients the close connection between the inflammatory state and risk of VTE. This further emphasizes the tight interplay between inflammation and coagulation cascade. In general, the cardiovascular risk in BP patients may be influenced not only by the inflammatory state but also by the treatment with systemic corticosteroids and immunosuppressive agents (160).

The high thrombotic tendency found in patients with BP, especially during the acute phase of the disease, implies some practical considerations. Firstly, one may wonder whether adding VTE prophylaxis to the immunosuppressive treatments could affect the thrombotic risk of BP patients. Clinical trials on efficacy and safety of antithrombotic drugs administered in the acute phase of the disease could provide some insight into their clinical relevance.

CONCLUSIONS

Coagulation and fibrinolysis activation markers are elevated in many inflammatory conditions, including not only a wide spectrum of systemic diseases of different pathogenesis such as

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rheumatoid arthritis, inflammatory bowel diseases, and sepsis but also autoimmune or immune-mediated cutaneous disorders. The activation of coagulation in autoimmune skin disorders may have two main consequences: on one hand, a local pathogenic role in inducing the skin lesions and on the other hand a systemic role in increasing the thrombotic risk. Concerning the pathophysiology of the skin lesions, thrombin contributes to increase endothelial vascular permeability, thus amplifying the inflammatory network in CU, angioedema and BP. In the skin microenvironment of CU and BP, eosinophils express tissue factor, which can activate coagulation right there, with the generation of vascular permeability mediators. Concerning the systemic implications of the coagulation involvement, several data indicate an increased thrombotic risk in BP. In contrast, no data are available concerning the incidence of thrombotic complications in patients with CU and angioedema. The retrospective evaluation of a large cohort of patients with BP has shown an increased incidence of venous thrombosis. Although most BP patients are elderly, the incidence of venous thrombosis in this kind of patients appears to be significantly increased as compared to the age-matched population of otherwise healthy subjects. BP is the prototype of autoimmune blistering disorder mediated by autoantibodies but in which an important physiopathological role is played by eosinophils. The intense skin inflammatory infiltrate is in fact characterized by the presence of eosinophils together with autoreactive CD4+ T lymphocytes and a few other inflammatory cells. The increased incidence of thrombosis in autoimmune skin diseases raises the question of the indication of anticoagulant treatment particularly in the presence of other cardiovascular risk factors.

AUTHOR CONTRIBUTIONS

MC, AM, and AB designed and drafted the manuscript. MC, AM, and AB made the literature search. MC, AM, AB, and SG discussed the topics of the manuscript and approved the final version.

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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