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Thrombotic microangiopathy after kidney transplantation: Expanding etiologic and pathogenetic spectra

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Abstract

Thrombotic microangiopathy (TMA) is an uncommon but serious complication that not only affects native kidneys but also transplanted kidneys. This review is specifically focused on post-transplant TMA (PT-TMA) involving kidney transplant recipients (KTRs). Its reported prevalence in the latter population varies from 0.8 to 14% with adverse impacts on both graft and patient survival. It has many causes and associations, and the list of etiologic agents and associations is growing constantly. The pathogenesis is equally varied and a variety of pathogenetic pathways lead to the development of microvascular injury as the final common pathway. PT-TMA is categorized in many ways in order to facilitate its management. Ironically, more than one causes are contributory in PT-TMA and it is often difficult to pinpoint one particular cause in an individual case. Pathologically, the hallmark lesions are endothelial cell injury and intravascular thrombi affecting the microvasculature. Early diagnosis and classification of PT-TMA are imperative for optimal outcomes but are challenging for both clinicians and pathologists. The Banff classification has addressed this issue and has developed minimum diagnostic criteria (MDC) for pathologic diagnosis of PT-TMA in the first phase. Management of the condition is also challenging and still largely empirical. It varies from simple maneuvers, such as plasmapheresis, drug withdrawal or modification, or dose reduction, to lifelong complement blockade, which is very expensive. A thorough understanding of the condition is imperative for an early

diagnosis and quick treatment when the treatment is potentially effective. This review aims to increase the awareness of relevant stakeholders regarding this important, potentially treatable but under-recognized cause of kidney allograft dysfunction.

INTRODUCTION

INTRODUCTION

Thrombotic microangiopathy (TMA) is an uncommon but serious complication that not only affects native kidneys but can also kidney allografts, with resultant graft dysfunction and graft loss. Its reported prevalence in the latter setting varies from 0.8 to 14% of kidney transplant recipients (KTRs) with adverse impacts on both graft and patient survival [1,2]. It is a highly heterogeneous condition with equally heterogeneous outcomes. TMA is not a single disease entity but rather a morphologic pattern of microvascular occlusive injury that can be seen with a variety of disease states and conditions. It has many causes and associations, and the list of these is growing steadily as new cases are being reported [3-7]. The heterogeneous etiology is reflected in a multitude of pathogenetic pathways leading to the final common pathway of occlusive microvascular injury [8,9]. This review is directed at post-transplant TMA (PT-TMA) in KTRs, which is an important cause of kidney allograft injury and loss if not treated promptly and appropriately. The main focus will be on the expanding etiologic and pathogenetic spectra with some description devoted to the pathology and diagnosis of the condition. The management and prognosis will not be dealt with in detail in this review. TMA not only involves the native kidneys but also occurs in the transplanted kidneys. The condition has many similarities as well as some differences in the two settings. This review will be confined mainly to PT-TMA in kidney transplant setting. A thorough understanding of the condition is imperative for an early diagnosis and quick treatment when the treatment is potentially effective. This review aims to increase the awareness of relevant stakeholders regarding this important, potentially treatable but under-recognized cause of kidney allograft dysfunction.

DEFINITION AND CLASSIFICATION

TMA is a clinicopathological syndrome characterized by endothelial injury and the presence of thrombi in the microvasculature (arterioles and capillaries). Thrombus formation in the vascular lumina leads to platelet consumption, damage to the red blood cells (RBCs), and occlusion of the lumina. The latter phenomenon leads to tissue ischemia and organ dysfunction, typically involving the kidneys but sometimes also other organs [10-14]. It is a potentially life-threatening condition. TMA is broadly categorized into two flagship clinical prototypes: hemolytic uremic syndrome (HUS) and thrombotic thrombocytopenic purpura (TTP). The former is characterized by severe kidney disease manifested as oliguria and uremia, but few extra-renal manifestations [11-13]. In the second form, TTP, the kidney changes are similar but milder than in HUS. However, widespread systemic manifestations, particularly, central nervous system (CNS) involvement, are highly prevalent [15, 16]. TMA syndromes are an emerging field of research and discoveries in nephrology, hematology, and rheumatology disciplines. Although many developments have taken place, much work remains to be done in genetics, molecular biology, and therapeutics to disentangle the conundrum of the relationships and the apparent differences between the different subclasses of TMA syndromes [16-18].

The classification of the TMAs is not only challenging but is constantly evolving. Traditionally, these were classified on the basis of clinical findings: TTP for predominant neurologic involvement and HUS for kidney-dominant disease. TMA syndromes can also be classified according to the pathogenetic processes involved in endothelial injury [19-21]. However, the ideal approach to categorize TMA is that of etiology, which, however, may not be identified in each and every case of TMA. Broadly, TMA is labeled as primary when a genetic or acquired defect is identified (as in atypical HUS [aHUS] and TTP) and secondary when it occurs in the setting of

another disease process, such as autoimmune disease, malignancy, infection, or drugs (Table 1). This subdivision is also not absolute because underlying genetic defects have been recognized in many cases of secondary TMA as well [21, 22].

POST-TRANSPLANT THROMBOTIC MICROANGIOPATHIES (PT-TMA)

TMA not only involves the native kidneys but also the transplanted kidneys. Kidney transplantation poses a challenging scenario due to multiple potential inciting factors for the development of TMA [1, 2, 23-26]. PT-TMA has many similarities with native renal TMA as well as some differences necessitating its detailed review. Like native kidney TMA, PT-TMA is caused by endothelial injury in the vast majority of cases and manifests as thrombotic occlusion of the microcirculation resulting in often clinically unexplained allograft dysfunction [27]. The endothelial injury may be caused by a myriad of injurious agents including but not limited to immunologic, genetic, and hematologic disorders and drugs either alone or in various combinations [28-30]. A kidney transplant biopsy is required for a definitive diagnosis [31]. The histopathologic diagnosis of Tx-TMA is based on the subjective interpretation of a large number of histopathologic lesions, whose nature, prevalence, and extent vary from case to case depending on many factors including the duration of the pathologic process. It also depends on the expertise and diagnostic insight of the pathologist [31]. Accurate diagnosis and classification are important for optimal treatment of the condition and favorable patient outcomes. The diagnosis can sometimes be challenging and delayed with consequent delay in the initiation of targeted treatment [32].

PT-TMA has been categorized in many ways. It can occur in a localized (L-TMA) form, limited to kidney allograft with resulting allograft dysfunction, or in a systemic form, with microangiopathic hemolytic anemia (MAHA), thrombocytopenia, and renal failure. In native kidneys, TMA is often part of the systemic illness, whereas in PT-TMA, it is often allograft kidney-limited. It can also be classified as recurrent or *de novo* PT-TMA; the latter being more common. Recurrent PT-TMA is almost invariably

complement-mediated, whereas *de novo* PT-TMA may be complement-mediated or secondary to other inciting factors (Table II). *De novo* TMA is reported in 0.8%–14% of KTRs, although the true frequency is unknown, and the incidence of a genetic complement abnormality may be underestimated [1, 2]. Differentiating between a primary complement-mediated process and one caused by secondary factors is important to minimize allograft damage since the former is non-responsive to supportive therapy and has a high risk of recurrence. However, distinguishing between the two types can be difficult, given their overlap of clinical, laboratory, and pathological features. TMA syndromes can also be classified according to the pathogenetic processes involved in endothelial injury. However, the ideal approach to categorize TMA is that of etiology, which, however, may not be identified in every case of PT-TMA (Figure 1).

ETIOLOGY OF PT-TMAS

The etiologic spectrum of PT-TMA is expanding and evolving with ever-increasing transplant activity. The etiology of PT-TMA not only includes all those causes that are seen in native kidney disease but also many additional causes unique to the transplant setting (Figure 1). The presence of a causal factor in isolation, such as ADAMTS13 deficiency or a complement mutation (the first hit), may not manifest clinically until a condition, such as an inflammatory disorder, surgery, or pregnancy (the second hit), precipitates an acute TMA episode. In fact, in PT-TMA, often more than one acquired factors are implicated in the causation of the disorder, leading to a proposal by some researchers of three-hit mechanism. It is important to identify all the predisposing factors in order to optimally treat the condition [33-38]. It is, however, often impossible to pinpoint to a single etiologic factor in an individual patient.

The role of immunosuppressant drugs posttransplantation in the development of PT-TMA deserves some attention. The two main groups of immunosuppressants used in all forms of transplantation, i.e., calcineurin inhibitors (CNIs) and mammalian target of rapamycin (mTOR) inhibitors (mTORi) such as sirolimus, can both trigger the

development of PT-TMA [39, 40]. The etiologic role of CNIs in causing PT-TMA is well established. However, the role of mTOR inhibitors is still largely controversial [40]. In vitro studies have suggested that sirolimus causes endothelial cell (EC) injury only when used in combination with tacrolimus. When used as single-agent, it does not lead to EC injury [41]. In clinical studies too, there is increasing evidence showing that sirolimus and everolimus, either alone or in combination with cyclosporine—can be associated with the pathogenesis of de novo PT-TMA. Some studies suggest that the impact of mTORi may be even higher in the development of TMA than that of CNIs. A USRDS-based study has demonstrated that there is a higher incidence of TMA in patients on initial maintenance therapy with sirolimus (18.1 episodes/1000 person-years) compared with those on CNIs (5.0 episodes/1000 patient-years) [42]. Some other studies have shown that replacing tacrolimus with rapamycin may improve PT-TMA. Thus, the exact mechanisms and roles of immunosuppressant drugs are still largely incompletely understood and need further research.

PATHOGENESIS

The pathogenesis of TMA is understandably as diverse as its etiological spectrum. The final common pathway in all forms of injury is endothelial damage with resultant activation of the thrombosis cascade. Within the TMA syndromes, two principal mechanisms participate: (1) endothelial injury and activation and (2) excessive platelet aggregation and activation. Among these, the endothelial injuries take precedence in HUS, whereas platelet aggregation and activation appear to be the main driving event in the TTP. Many different etiological precipitating factors have been described for the development of PT-TMA, such as ischemia-reperfusion-injury, use of immunosuppressive drugs, infections, and many more [43, 44].

Endothelial injury

A variable degree of EC injury and activation is the hallmark of all TMA syndromes and in many TMA syndromes, constitutes the final common pathway of microvascular injury. The endothelium is a highly active and dynamic tissue responsible in part for regulating vascular tone, coagulation, and inflammation [45, 46].

All types of TMA are characterized by a common phenotype of activated, prothrombotic ECs. This EC phenotype arises from various distinct types of injurious agents: complement activation, autoimmune diseases, infections, drug toxicity, or malignancy. For most types of TMAs, the exact intracellular mechanisms of EC injury are not well understood.

In typical or classic HUS, the initiating factor for endothelial injury and activation is usually a Shiga-like toxin, whereas, for atypical and inherited forms of HUS, excessive or inappropriate activation of complement is the main triggering event. Many other injurious agents and conditions can sometimes precipitate a HUS-like condition, probably also by damaging the endothelial layer. The EC injury in HUS causes platelet activation, aggregation, and thrombus formation within the lumina of the microvasculature. Previous research has found that reduced production of prostaglandin I₂ and nitric oxide by ECs contributes to intraluminal thrombosis. The reduced production of the above two factors and increased production of EC-derived endothelin also promote vasoconstriction, accentuating the hypoperfusion of organs [47-49].

Platelet aggregation

This is the second main pathogenetic pathway of thrombus formation in TMA syndromes, manifesting clinically as TTP. In this scenario, endothelial structure and function are relatively intact. In this pathway, the initiating event is the platelet aggregation induced by ultra-large multimers of vWF, which accumulate to a deficiency of ADAMTS13, a plasma protease that degrades vWF multimers into smaller fragments. The deficiency of ADAMTS13 is most often functional caused by autoantibodies that inhibit its function. This form of TTP is referred to as acquired or immune TTP (iTTP)

and accounts for 95% of cases. Rarely, an inherited deficiency of ADAMTS13 Leads to a chronic relapsing and remitting form of TTP. This pattern of disease is labeled as inherited or congenital TTP (cTTP) and is rare [48-50].

PATHOLOGY OF PT-TMA_s

A large variety of morphological lesions can be found on kidney allograft biopsies in cases of PT-TMA. The lesions may involve glomeruli, arterioles, and rarely small arteries in variable combinations and with varying degrees of severity (Table III). Their nature varies according to the duration of the disease process and may be categorized as acute, chronic, or acute-on-chronic [51-55]. The morphological features in various types of TMA syndromes are indistinguishable and vary mainly according to the age of the lesion than the cause of TMA. The glomeruli in active disease may show many non-specific changes such as marked congestion, bloodless appearance, capillary collapse, mild to moderate cellular proliferation, crescent formation, and rarely, complete infarction. ³ Disruption of the mesangial matrix and damage to the mesangial cells may result in mesangiolysis and aneurysmal dilatation of the capillary loops. More specific features include the thickening of the capillary walls by expansion of the subendothelial zones, intraluminal thrombi, and the presence of red cell fragmentation and extravasation into vessel walls (Figure 2). The glomerular lesions vary from case to case and from glomerulus to glomerulus.

The arterioles and small arteries in acute PT-TMA show intraluminal thrombosis and subendothelial edema resulting in marked narrowing of the lumina. Red blood cell fragmentation and extravasation in the walls of arterioles may be observed. Medial necrosis, fibrinoid necrosis, and intramural thrombosis may be seen in severe cases.

Chronic TMA lesions are commonly observed in patients with aHUS and manifest as lesions

emanating from continued endothelial injury and attempts at repair. The glomeruli are mildly hypercellular and show thickened capillary walls with double contours or tram-

tracking, producing mesangiocapillary pattern of injury. The double contours result from reduplication and formation of neobasement membrane because of persistent injury to the endothelium. In vessels, medial hypertrophy (onion-skinning) may be seen in lesions of longer duration. Mucinous intimal thickening with marked narrowing of lumina is characteristically observed in chronic TMA lesions involving the arterioles and small arteries (Figure 3).

Immunofluorescence shows deposits of fibrin in glomeruli and arterioles. There may be weak nonspecific positivity of IgM in the glomeruli and arterioles with less frequent C3 and IgG. Fibrin is invariably present in the fibrin thrombi.

Electron microscopy shows separation of the endothelium from the underlying glomerular basement membrane (GBM) by electron-lucent zone filled with fluffy electron-lucent material during early phase of the disease. Within this space also lie scattered fine fibrils, occasional stands of fibrin, fragments of red blood cells and platelets and cytoplasmic processes of mesangial and endothelial cells. No electron dense deposits are found. A newly formed basement membrane is found below the endothelial layer. Mesangial changes may be marked on ultrastructural level [51-55].

Till recent past, the diagnostic criteria were not standardized for the clinical or pathological diagnosis of PT-TMA. Moreover, the histopathologic diagnosis is a subjective task. The Banff Working Group (BWG) on TMA was formed in 2016 under the auspices of the Banff Foundation for Allograft Pathology, with the aim of standardizing the diagnostic criteria of TMA and formulating recommendations [31]. A survey conducted in January 2016 among the BWG participants, showed considerable heterogeneity among pathologists, using a variety of known TMA features with imprecise or subjective definitions. Therefore, the first objective of the BWG was to provide the nephropathology community with a standardized set of minimum diagnostic criteria (MDC) for PT-TMA. A secondary objective, identified during the study, was to scrutinize specific lesions that could potentially determine specific etiologies of PT-TMA. Diagnosis of TMA in the renal allograft is not merely a

morphologic task; clinical and laboratory information is also critical for diagnosis and needs to be standardized in phase II of the study. The Delphi approach was used by the BWG, for the first time in the Banff classification, to generate consensus, among an expert panel [31]. The group generated consensus on 24 criteria, provided a list of eight differential diagnoses, and identified areas of diagnostic difficulty. According to the authors this work is a starting point in the process of diagnosing PT-TMA in KTRs [31].

DIAGNOSIS OF PT-TMAS

Overall, there is a lack of international consensus criteria for the diagnosis of PT-TMA. Moreover, the clinical and laboratory features of the condition are non-specific and protean. This is reflected in the wide variation in the reported incidence of PT-TMA. An algorithmic approach to diagnosis, classification and treatment is presented in Figure 4. Recently, the BWG on TMA has published the results of phase I of the consensus process for MDC for the pathologic diagnosis PT-TMA in KTRs. The other main group of PT-TMA relates to patients undergoing hematopoietic stem cell transplantation (HSCT). Different diagnostic criteria are used in the hematology discipline. There is no uniformity in the approach to diagnosis and investigation in these two broad groups of PT-TMA. There is a clear need for unified, objective, and organ-specific criteria to help in the timely diagnosis of TMA in clinical practice and for use in future clinical trials.

MANAGEMENT AND PROGNOSIS

Management of the condition is challenging and still largely empirical. It varies from simple maneuvers, such as plasmapheresis, drug withdrawal or modification, or dose reduction, to lifelong complement blockade by eculizumab, which is very expensive approach (Figure 5). Careful donor selection and proper recipient preparation, including complete genetic screening, would be a more rational approach. Novel

targeted therapies are being actively researched but are still in the experimental phase and are not yet available in clinical practice [56-59].

The prognosis of *de novo* or recurrent TMA in kidney allografts is generally guarded and varies according to underlying causes [60-68]. With better understanding and characterization of the disease, the patient and allograft outcomes are improving steadily.

PREVENTIVE/PROPHYLAXIS MEASURES

These measures or strategies can only be applied in cases of aHUS for possible risk of recurrence after kidney transplantation (Figure 5). The risk of recurrence depends on the type of mutation in complement regulatory proteins and can be calculated before transplantation. Recurrence usually occurs very early in the posttransplant period and may be precipitated quickly by an ischemia-reperfusion-induced endothelial injury. However, the time between kidney transplantation and aHUS recurrence varies considerably. Due to the severity of aHUS recurrences and the unpredictable time of onset, the KDIGO workgroup recommends the prophylactic use of eculizumab for KTRs who are at high risk of recurrence based on the patient's genetic background. Eculizumab has been used both before and after transplantation. An analysis of the Global aHUS Registry showed that pretransplant use of eculizumab resulted in better allograft function than posttransplant initiation. Other preventive measures include pretransplant plasma exchange (PE), use of induction therapy and low doses of CNIs. For some complement regulatory gene mutations, use of liver-kidney transplantation has been used successfully [69]. This procedure is controversial because of potentially severe postoperative complications but the use of PE or a single dose of eculizumab until graft liver function is adequate greatly improved outcomes for the patient. However, this type of transplant should only be performed in centers with proven expertise, after a careful risk-benefit analysis.

CONCLUSION

CONCLUSION

PT-TMA is an important but underestimated cause of kidney allograft dysfunction and loss. Its etiologic spectrum and associated pathogenetic pathways are expanding steadily. Its early diagnosis and treatment are challenging. Recently attempts have been made to standardize the pathologic diagnostic criteria for its accurate diagnosis so as to optimize treatment approaches. There is a need to adopt a unified and international consensus-based approach across all the relevant specialties involved for standardizing and optimizing TMA diagnosis and management.

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