

Non-HLA Antibody Induced Agonism on the Angiotensin II Type 1 Receptor in Renal Allograft Vascular Injury

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Abstract

The rising incidence of steroid-refractory rejections is a challenging problem in renal allotransplantation. Etiological studies implicate either an overwhelming T-cell response or, more frequently, involvement of alloantibodies. Alloantibodies induce a spectrum of histologic tubulointerstitial and vascular changes paralleled with immunohistochemical positivity for C4d along peritubular capillaries. The degree of vascular involvement seems to be a more important prognostic determinant. Fibrinoid necrosis of the arteries with secondary thrombotic occlusions is C4d negative in 50% of cases and has a worst prognosis among all allograft vascular lesions. Apart from donor-specific human leukocyte antigen antibodies, non-human leukocyte antigen antibodies reacting to arterial antigens have been speculated to be responsible for rejections in some patients. We recently reported the presence of agonistic antibodies against the angiotensin II type 1 receptor (AT₁R-AA) in 16 recipients of renal allografts who had severe vascular rejection and malignant hypertension, but who did not have anti-human leukocyte antigen antibodies. The AT₁R-AA appear to be non-complement-fixing autoantibodies targeting the second extracellular loop of AT₁R. The AT₁R-AA act as allosteric activators on AT₁R and induce mediators of inflammation and thrombosis. Transfer of AT₁R-AA into rats with kidney allografts induced vasculitis and hypertension, supporting the notion that AT₁R-AA are not an epiphenomenon. Removal of AT₁R-AA by plasmapheresis in combination with pharmacologic AT₁R blockade leads to improved renal function and graft survival in AT₁R-AA-positive patients. We have shown that the analysis of the subtle diagnostic and mechanistic differences may help to identify patients at particular risk and improve outcome of steroid-refractory rejections with vascular pathology. (Trends in Transplant. 2007;1:113-20)

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Introduction

The successful developments in immunosuppressive modalities that were designed to target T-cell-mediated immune response and prevent destruction of tubular epithelia have resulted in the reduction of acute rejection episodes and improved overall allograft survival¹. On the other hand, humoral pre-sensitization emerges as one of the major risk factors for renal allograft rejection and allograft loss², despite the fact that most of the patients with antibody mediated rejection (AMR) have negative crossmatch. Generally, AMR has a worse prognosis than the T-cell-mediated rejection, which forced significant research efforts during the last decade³. Antibody mediated rejection remains a diagnostic and therapeutic challenge and can occur with all immunosuppression regimens, even in the context of profoundly depletion therapy⁴. It is not possible to distinguish AMR from T-cell-mediated rejection on simple clinical grounds. However, when AMR occurs in organ transplants, the process resists conventional treatment approaches and frequently leads to allograft loss⁵. The morphologic features apply to a wide spectrum of tubulointerstitial and vascular lesions in the allograft also, including severe changes like thrombosis, fibrinoid necrosis of the arteries, and endarteritis^{6,7}. Neutrophils in the capillaries are characteristically but not always found⁶. Among different morphologic features, vascular lesions carry the worst prognosis⁸. The association of antidonor humoral reactivity against human leukocyte antigen (HLA) class I antigens and vascular rejection has been documented in studies by Halloran, et al. more than a decade ago⁹. Donor-specific anti-HLA alloantibodies initiate rejection through complement-mediated and antibody dependent cell-mediated cytotoxicity¹⁰. The diffuse staining of the complement degradation product C4d affecting the surface of peritubular capillaries is generally regarded

as a marker for HLA antibody mediated allo-response and is associated with inferior graft survival¹¹. Nevertheless, 40-50% of rejections with severe vascular changes such as fibrinoid necrosis are C4d negative, implicating involvement of non-complement-fixing antibodies against undefined targets¹². Clinical, etiological, and histopathologic heterogeneity of AMR emphasizes the necessity for better recognition of the subtle etiologic differences between affected patients. Unknown immune targets and consecutive lack of detection methods make non-HLA AMR particularly difficult to diagnose and treat.

Relevance of non-HLA antibody response

Putative pathogenic antibodies that are not directed against the HLA system were considered in recipients who rejected HLA-identical kidneys more than three decades ago¹³. However, characterization of non-HLA antibodies remains very poor; many appear to be autoantibodies¹⁴. In renal allograft rejection, the presence of antibodies to non-HLA antigens has been associated with antibodies against endothelial cells, tubular epithelial cells, podocytes, mesangial cells, and monocytes¹⁴. Most of the efforts in the past were focused on anti-endothelial cell antibodies¹⁵. Anti-endothelial cell antibodies are a heterogeneous group of antibodies directed against a variety of antigenic determinants, but the existence of a common polymorphic non-HLA antigen system in endothelial cells could not be confirmed by biochemical identification of the relevant antigens¹⁶. They have been reported in a variety of autoimmune diseases featuring vasculitis as a denominator¹⁵. The most comprehensive evidence about their biologic relevance is derived from studies using immunoglobulins isolated from patients with systemic sclerosis¹⁷. Anti-endothelial cell antibodies are especially common in renal transplant recipients who are pre-sensitized

against a panel of HLA antigens¹⁸. Non-HLA-reactive anti-endothelial cell antibodies seem to recognize endothelial cell antigens that can be induced upon tumor necrosis factor- α (TNF α) and interferon- γ (IFN γ) stimulation, implicating a permissive role of endothelial activation that may be a prerequisite for the pathogenicity of anti-endothelial cell antibodies¹⁸.

Similarly to some autoimmune diseases, non-HLA antibodies may be diagnostic of disease, but they may not necessarily be an effector mechanism. For this reason, it is important to identify non-HLA antibodies, determine their antigen specificity and pathogenicity, and clarify the mechanisms by which they contribute to rejection or other forms of allograft injury.

Clinical manifestations of AT₁R-AA related vascular rejections

We reported the presence of agonistic antibodies against the angiotensin II type 1 receptor (AT₁R-AA) in 16 recipients of renal allografts who had severe vascular rejection and malignant hypertension, but who did not have donor-specific anti-HLA antibodies¹⁹. These AT₁R-AA have also been associated with preeclampsia and malignant hypertension^{20,21}. Pregnancies complicated by preeclampsia and graft rejection bear some immunologic similarities²². The decision to seek and isolate AT₁R-AA was instigated by the observation that the first patient we studied developed accelerated vascular rejection refractory to steroids and anti-lymphocyte antibody preparations in a “zero-mismatch” kidney. As this patient developed malignant hypertension with seizures during the rejection process, the clinical picture was so reminiscent of eclamptic crisis in pregnancy, a condition that she had developed two decades before transplantation, that we start-

ed to prospectively look for patients with similar clinical features. We detected our further 15 patients primarily based on severe vascular pathology, absence of donor-specific antibodies, hypertensive crisis accompanied by seizures in three other patients, and lack of response to steroids or anti-lymphocyte preparations. All patients in our study had primary graft function.

Clinicopathologic features of AT₁R-AA-related process in patients were noticed between day 5 and 14 posttransplantation, with minor interindividual differences concerning temporal occurrence of maximal allograft injury and increase in blood pressure. Most of patients (13/16) did not have hypertension before vascular rejection occurred, implying that the posttransplant hypertension was most likely secondary to rejection. Some of the patients developed thrombopenia together with other signs of microangiopathic hemolysis such as an elevated lactic dehydrogenase and presence of schistocytes. Reactivation or *de novo* cytomegalovirus infection was excluded in all patients. Autoimmune and hereditary causes of thrombophilia were also ruled out.

Causes of end-stage renal disease (ESRD) were not primarily attributed to hypertensive nephrosclerosis, but instead to a variety of tubulointerstitial or glomerular diseases. None of the patients in our study was reported to have ESRD due to renal involvement in a systemic autoimmune disease or hemolytic uremic syndrome/thrombotic thrombocytopenic purpura. We believe that the specific cause of ESRD was not relevant for the development of AT₁R-AA in our patients. The frequency of AT₁R-AA-positive rejections was equal between both female and male renal transplant recipients. The AT₁R-AA-related vascular rejections occurred during the first week after transplantation. These rejections had significantly shorter allograft survival and more severe histology, irrespective of the

treatment, compared to donor-specific anti-HLA-antibody associated rejections¹⁹.

Morphologic features of AT₁R-AA-related vascular rejections

The majority of AT₁R-AA-positive patients detected in the initial study developed Banff type III rejection with fibrinoid arterial wall necrosis and secondary thrombotic occlusion with allograft infarction. In some of the patients, first biopsies were presented with Banff type II rejection, transmural arteritis (transplant endarteritis), and transplant glomerulitis. Apart from arterial and capillary changes, we also noticed tubulitis and interstitial infiltrates, characteristic for acute cellular rejection. Thus, some biopsies seem to fall into the category of "mixed rejection". However, unlike so-called "mixed AMR"²³, our patients did not respond to aggressive T-cell depletion therapies. In few patients, hyperacute rejection manifested as fibrinoid arterial wall necrosis, and interstitial hemorrhage with frank necrosis of tubular epithelia developed during the first three posttransplant days. Thus, AT₁R-AA-positive rejections tend to share many morphologic features with both "pure" and "mixed" HLA-antibody mediated AMR. Patients with AT₁R-AA-positive rejections had worse vascular scores and Banff grade compared to those with HLA-mediated AMR. In contrast to HLA antibodies, AT₁R-AA seem to operate through complement-independent mechanisms, as C4d was detected in biopsy specimens from only five of our 16 patients¹⁹.

Diagnostic and therapeutic implications

A very short period from transplantation to rejection episode implicated a possible relevance of preformed antibodies. Retrospec-

tive analysis of historic sera from our patients obtained before transplantation showed positivity for AT₁R-AA, which confirms that preformed and not *de novo* produced AT₁R-AA were likely responsible for vascular rejection. The AT₁R-AA are low-titer IgG1 and IgG3 (complement-fixing subclass) antibodies, yet they do not form immunocomplexes with the antigen. Detection of AT₁R-AA activity initially relied on the bioassay that measures the chronotropic responses to AT₁R-IgG-mediated stimulation of cultured cardiomyocytes coupled with receptor-specific antagonists. The dose/response relationship between AT₁R-AA concentration and the chronotropic response is linear. High costs and a time-consuming test setting precluded screening of larger patient cohorts by the bioassay at the time when the initial study was performed. Only patients with suggestive clinical features and biopsy findings, and not all patients with allograft dysfunction, underwent bioassay test.

In our initial study, seven of 16 patients with AT₁R-AA were treated with a combination of plasmapheresis, intravenous immune globulin infusions, and the AT₁R-blocker, losartan (100 mg daily). This combination treatment led to improved renal function and graft survival, compared to the outcomes amongst patients with AT₁R-AA who received standard treatment for AMR and rapidly lost their allografts¹⁹. None of the patients received angiotensin converting enzyme (ACE) inhibitors or AT₁R blockers prior to rescue protocol, as the practice of our transplant center was not to use them in the posttransplant period. Among seven treated patients, four remained rejection free and AT₁R-AA negative (longest follow-up, seven years). They still continuously receive 50 mg losartan daily. One patient died six months after successfully treated AT₁R-AA rejection with functioning graft due to herpes simplex virus encephalitis. Two patients who were initially rescued from AT₁R-AA rejection developed *de novo* anti-HLA class I directed against A29 and A8 loci of the donor-

specific antibody and C4d positive glomerular type of humoral rejection without affection of arterial vessels after a four- and six-month period of stable allograft function, respectively. They required rituximab rescue therapy due to plasmapheresis and intravenous immunoglobulin resistance.

Subsequent serum samples were obtained from five of eight patients who were not treated with our losartan-including rescue protocol. Two of these patients became AT₁R-AA negative approximately two months after transplant nephrectomy (removal of antigen). However, three other patients remain AT₁R-AA positive despite transplant nephrectomy. Interestingly, all these three patients previously lost two transplants due to undefined accelerated vascular rejections. We speculate that these patients may have altered plasma cell memory repertoire in terms of long-lived plasma cells that are not responsive to removal of antigen compared to other patients.

We are aware that the small number of patients in our initial study may limit the degree to which the results can be generalized, and that bioassay testing under a suggestive indication provided left-censorship bias. However, we believe our findings, even with their limitations, are highly significant. Due to high costs and our time-consuming bioassay, larger studies were initially not feasible. We have now established and validated a cell-based ELISA in collaboration with biotech partners for detection of AT₁R-AA in serum²⁴. The ELISA currently has 100% specificity and 88% sensitivity. Variability between assays is 12%²⁴. Pretransplantation screening for AT₁R-AA detects a subset of ESRD patients who are similar but not identical to patients with anti-HLA-panel reactivity. Pretransplantation screening of recipients for AT₁R-AA may help to improve individual risk assessment and offer patients with AT₁R-AA preemptive specific treatment. Whether AT₁R-AA acts as a progression factor during native renal dis-

ease, or as an independent factor of cardiovascular comorbidity, remains to be determined. Whether all AT₁R-AA-positive patients who will be continuously treated with AT₁R blockade will develop milder or no vascular rejection is the subject of current studies. Some transplant nephrologists are the only remaining clinicians skeptical about the use of anti renin-angiotensin system (RAS) drugs due to the concern of interference with renal allograft perfusion. According to reported beneficial effects of blockade of RAS on early outcomes of renal transplants, this view seems to be outdated²⁵. Moreover, AT₁R antagonists may exert a clinically relevant immunomodulatory role by blocking IFN γ production by T-cells²⁶. Pharmacologic action of AT₁R antagonists is based on inverse agonism, which implicates that reactive upregulation of AT₁R on target cells may, in case of therapy discontinuation, increase detrimental responses²⁷. Given this consideration, perioperative discontinuation of ACE inhibitors or AT₁R blockers may thus predispose for AT₁R-AA-related pathologies. For example AT₁R-AA-positive patients who receive continuously AT₁R blockers or ACE inhibitors together with intensified immunosuppression (depletional antibody induction, tacrolimus, mycophenolate mofetil, and steroids) and are recipients of living-donor kidneys seem not to be prone to development of fulminant AT₁R-AA-related rejection²⁸.

Pathophysiologic consequences of antibody mediated AT₁R stimulation

The AT₁R-AA seem to induce vascular and tubulointerstitial pathology via mechanisms independent from complement activation that are distinct from those in patients with HLA antibodies. We raised and confirmed the hypothesis that AT₁R-AA may act in similar manner as an endogenous agonist for the AT₁R, angiotensin II, and exert direct effects

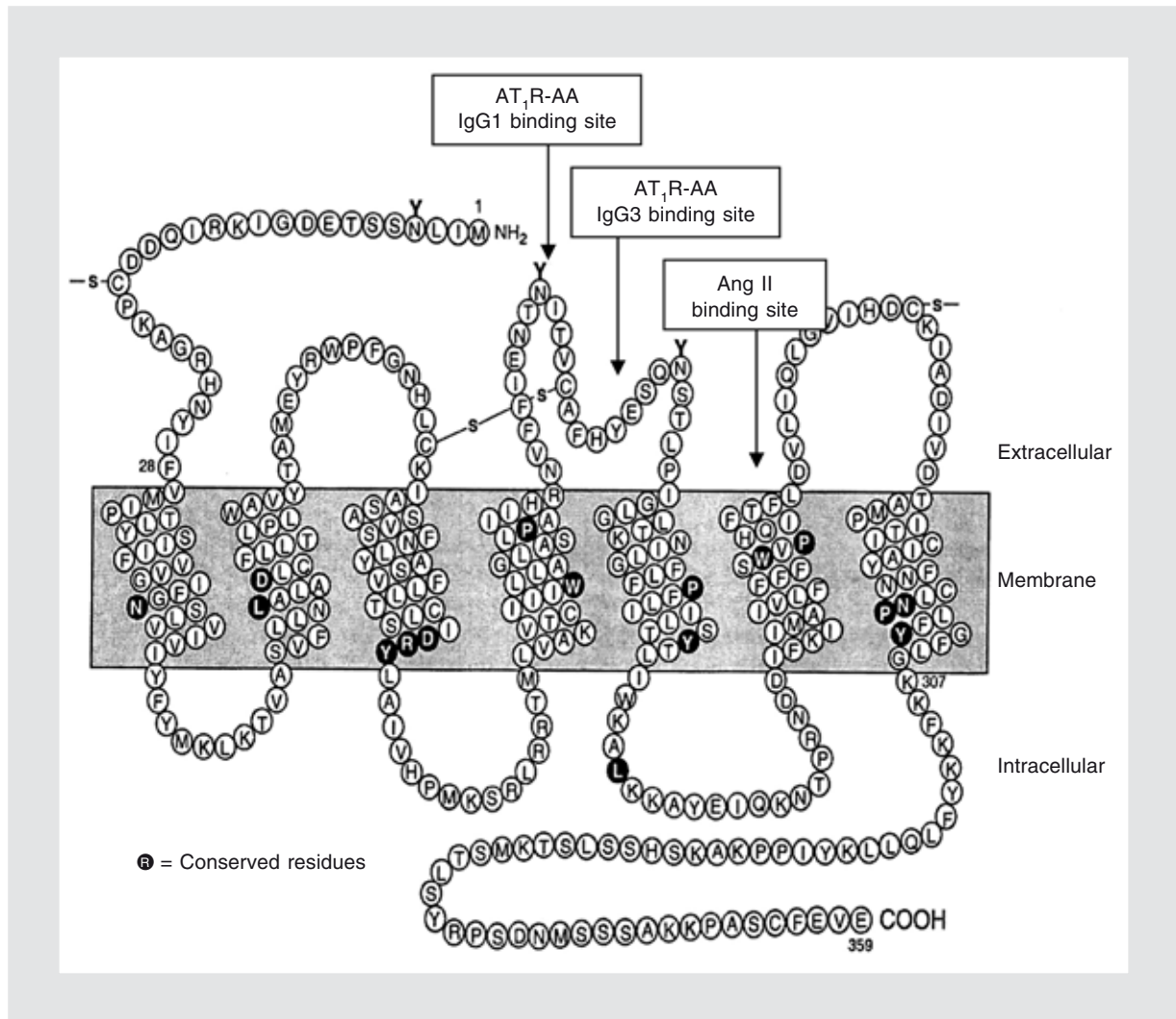


Figure 1. Secondary structure and consensus sequence of the mammalian angiotensin II type 1 receptor. The amino acid sequence shown is based on the derived sequences of five individual cloned mammalian AT₁ receptors. The amino acid residues that are highly conserved among G-protein coupled receptors are indicated by bold letters. The positions of the three extracellular carbohydrate chains, and of the two extracellular disulfide bonds, are also indicated. IgG1 and IgG3 AT₁R-AA bind to the second extracellular loop of the receptor, while a natural agonist angiotensin II binds to transmembrane “pockets” close to the plasma membrane.

on endothelial and vascular smooth muscle cells²⁹. The responses elicited by AT₁R stimulation are context dependent and specific for target-cell lineage³⁰. According to our working concept, AT₁R-AA bind to the second extracellular loop of AT₁R (Fig. 1) and act as an allosteric receptor agonist. The AT₁R-AA/AT₁R interaction initiates signal transduction cascades by inducing extracellular signal-regulated kinase 1/2 phosphorylation in endothelial and vascular smooth muscle cells. Consequent increasing DNA binding activity

of transcription factors activated activator protein 1 (AP-1) and nuclear factor- κ B (NF κ B) is responsible for increased expression of their target genes involved in inflammatory responses and coagulation. Increased synthesis of chemokines MCP-1 and RANTES may probably explain intravascular inflammatory cell infiltration, while augmented activity and expression of tissue factor may account for thrombotic angiopathy. Although we have documented that AT₁R-AA belong to complement-fixing IgG1 and IgG3 antibodies, our

findings suggest that genes regulated by AT₁R-triggered transcription factors and not complement-directed cytotoxicity act as an effector pathway of the vascular injury. The illustrative example is that AT₁R-AA enhanced promoter activity of tissue factor, an initiator of extrinsic coagulation pathway and a target gene for AP-1 and NFκB *in vitro*. Tissue factor mediates clotting abnormalities associated with hyperacute and xenograft rejection, as well as in antiphospholipid antibody syndrome³¹. Accordingly, renal transplant biopsy specimens obtained during an AT₁R-AA-mediated rejection episode revealed intense diffuse tissue factor staining of epithelial, endothelial, and mesangial cells in absence of complement activation. Binding of AT₁R-AA to AT₁R expressed on target cells is a critical step for activating the downstream cascade and inducing damage to the allograft. However, we have not yet proven whether AT₁R-AA function only through pro-coagulatory and chemotactic activity, or whether they also act by means of innate and specific immune responses and increased vascular reactivity. Direct effects of AT₁R-AA on immune response are likely, since human T-cells are fully equipped with functioning components of the renin-angiotensin system and express AT₁R on their surface³². Our current working hypothesis is that factors surrounding the organ transplantation process may lead to increased expression of AT₁R and thereby affect the overall reactivity of the vascular cells to AT₁R-AA. Passive transfer of human IgG containing AT₁R-AA induced a transmural arteritis similar to the human situation and led to increased blood pressure in otherwise non-rejecting and normotensive transplanted animals. These findings provided further evidence that AT₁R-AA may have a causative role. Similar to stimulation of the AT₁R by its natural ligand, angiotensin II²⁹, agonistic receptor activation mediated by AT₁R-AA could play a key role in the initiation and amplification of pathobiological events that lead to transplant vasculitis and hypertension.

Unresolved questions

We have not explained whether or not AT₁R-AA-related pathology represents a “true-rejection” or an autoimmune phenomenon that becomes overt in dependence of permissive factors related to allogeneic environment and not yet elucidated factors related to the transplant procedure itself. An allogeneic background, brain death-associated “cytokine storm”, reperfusion injury to the transplant, and/or use of calcineurin inhibitors or steroids are probably permissive factors responsible for an increased AT₁R density on target cells. For example, in heart transplantation, systemic upregulation of AT₁R could be found in donors with spontaneous intracerebral hemorrhage that was associated with subsequent development of cardiac vasculopathy³³. However, the relative individual contribution of considered permissive factors needs to be further elucidated in order to better understand and prevent AT₁R-AA-related clinical syndrome. Another important question is, why do AT₁R-AA develop in patients with preeclampsia and ESRD and what is the role of antigen mimicry (cross-reactivity with bacterial or viral antigens) or genetic predispositions?

Conclusion

We provide a novel concept in the pathogenesis of accelerated vascular rejection process where autoimmune-mediated receptor activation is linked to a severe vascular pathology in the situation of allogeneic transplantation. At present we believe that pre-transplantation testing of recipients for AT₁R-AA may help to improve individual risk assessment of patients with AT₁R-AA-preemptive specific treatment. We are also just beginning to learn more about pathophysiologic mechanisms and optimization of diagnostic and therapeutic modalities for AT₁R-AA-positive patients.

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