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**Acute Liver Allograft Antibody-Mediated Rejection: an inter-institutional study of
significant histopathological features**

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Abbreviations:

AMR, antibody-mediated rejection

aAMR, Acute-AMR score

BD, bile duct

BUMC, Baylor University Medical Center

DSA, donor-specific HLA alloantibody

MFI, mean fluorescence intensity

PV, portal vein

UPMC, University of Pittsburgh Medical Center

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ABSTRACT

Acute antibody-mediated rejection (AMR) occurs in a small minority of sensitized liver transplant recipients. Although histopathologic characteristics have been described, specific features that could be used: a) for a generalizable scoring system; and b) to trigger a more in-depth analysis are needed to screen for this rare but important finding. Toward this goal, we created a training and validation cohort from 3 high volume liver transplant programs of putative acute AMR and control cases that were evaluated blindly by 4 independent transplant pathologists. Evaluations were performed on H&E sections alone without knowledge of either serum DSA results or C4d stains. Routine histopathological features strongly correlated with severe acute AMR included portal eosinophilia, portal vein endothelial cell hypertrophy, eosinophilic central venulitis, central venulitis severity, and cholestasis. Acute AMR inversely correlated with lymphocytic venulitis and lymphocytic portal inflammation. These and other characteristics were incorporated into models created from the training cohort alone. The final **Acute-AMR (aAMR) score** (portal vein endothelial cell hypertrophy + portal eosinophilia + eosinophilic venulitis / lymphocytic portal inflammation + lymphocytic venulitis) exhibited a strong correlation with severe acute AMR in the training (OR=2.86, $p<0.001$) and validation cohort (OR=2.49, $p<0.001$). SPSS tree classification was used to select 2 cutoffs, one that optimized specificity at a score >1.75 (sensitivity = 34%, specificity = 87%) and a second that optimized sensitivity at a score >1.0 (sensitivity = 81%, specificity = 71%). In conclusion, routine histopathological features of aAMR score can be used to screen for acute AMR on routine H&E in indication liver transplant biopsies, however, a definitive diagnosis requires

substantiation by donor-specific HLA alloantibody testing, diffuse C4d staining, and exclusion of other insults.

Accepted Article

INTRODUCTION

The first evidence that antibodies can cause acute injury/rejection (antibody-mediated rejection; AMR) in human liver allografts was observed in ABO-incompatible cadaveric, brain-dead whole organ donors (1, 2). Antibody and complement deposition, platelet-fibrin thrombi, micro-vasculitis, and arteritis were typical and expected histopathological findings (1), based on previous observations in ABO-incompatible renal allografts (3) and in ABO-compatible renal allografts harboring alloantibodies (4, 5).

It was recognized early on, however, that human liver allografts were highly resistant to acute AMR from preformed HLA alloantibodies compared to kidney allografts (6). This relative resistance was attributed to: the liver's inherent "tolerogenic" properties, the difficulty detecting antibody and complement tissue deposits, the paucity of typical histopathological findings (6) and, even when damage was present, to the noticeably diminished severity of injury compared to ABO-incompatible liver transplants (7, 8). Relative hepatic resistance to AMR has been attributed to: a) secretion of soluble HLA class I molecules that form immune complexes with alloantibodies, which are then cleared by Kupffer's cells; b) Kupffer cell phagocytosis of platelet aggregates, immune complexes, and activated complement components (9); c) limited distribution of HLA class II expression in the microvasculature; d) large liver size and dual hepatic vasculature; and e) marked hepatocyte regenerative capacity after injury [reviewed in (7, 8)]. In addition, the inferior sensitivity and specificity of cell-based cytotoxic antibody detection methods impaired prior investigators abilities to find associations between HLA antibodies and adverse patient and graft outcomes (1, 7, 8).

Nevertheless, in the late 1980's and early to mid-1990's HLA class I and II antibodies, as measured in cytotoxic cell-based assays, were suspected to cause or substantially contribute to acute and chronic liver allograft rejection (7, 10-13). In addition, experimental rat studies clearly showed that extreme sensitization (14, 15) could override the liver's natural resistance and defense mechanisms. Similar observations were made in humans and risk factors for acute liver allograft AMR included high-titer pretransplant sensitization with persistence of serum alloantibodies after transplantation. When acute liver allograft AMR ensued, refractory thrombocytopenia, circulating immune complexes, and severe liver injury were then seen (7, 11).

Recent studies using more sophisticated and sensitive (16) solid phase donor-specific HLA alloantibody (DSA) detection methods have confirmed and extended earlier studies with cytotoxic cell-based assays, even though the two tests have been documented to sometimes produce substantially different results on the same serum samples (17). These confirmed findings include: 1) the liver allograft's relative resistance to AMR (18, 19) associated with the rapid disappearance of the vast majority of low to moderate MFI class I and II alloantibodies (11, 17, 18); and 2) an association of acute AMR with high-titer alloantibodies that most-often persist after transplantation and result in refractory thrombocytopenia and acute liver injury that can evolve into combined acute antibody-mediated and T-cell-mediated rejection. Inadequately treated, the end result can be chronic or ductopenic rejection (17, 20-24). Solid phase DSA analyses have also shown an association between multiple IgG subclasses, especially when alloantibodies of the IgG3 subclass are present, and chronic rejection and

diminished allograft survival (25). These newer serum assays have also facilitated a closer correlation between histopathological findings and serum DSA characteristics (18, 23, 24, 26).

Histopathological patterns of injury associated with acute liver allograft AMR include organ-specific findings such as portal edema, ductular reaction, eosinophilia, hepatocyte swelling and hepatocanicular cholestasis, and histopathological findings similar to those seen with acute AMR in other solid organ allografts such as marked (portal) microvascular endothelial cell hypertrophy and monocytic/histiocytic, eosinophilic, and neutrophilic (portal) microvasculitis. During the early stages, the constellation of findings can resemble preservation/reperfusion injury or biliary stricturing, but often quickly progress to acute “cellular” or T-cell-mediated and finally chronic rejection (8, 11, 22-24, 26, 27). Detection of microvascular complement deposition with C4d staining has been a valuable adjunct to the histopathological evaluation for acute AMR in all solid organ allografts, but C4d staining should not be used in isolation to establish an AMR diagnosis in liver allografts [reviewed in (27-29)].

Finally, although severe acute AMR is rare, unrecognized it can lead to allograft failure (23, 30, 31), as evidenced by its substantial contribution to ~ 10-20% of previously idiopathic early allograft failures (<90 days post-transplant) in sensitized patients (24). Early recognition of acute AMR can prompt plasmapheresis (32) and plasma cell-specific therapy in rare patients and may result in improved outcomes (23, 30). Toward the goal of facilitating earlier diagnosis of severe acute liver allograft AMR, this study was designed to identify and validate a limited constellation of routine histopathological features in the form of a generalizable scoring system on liver biopsy

H&E analysis that can be easily used to trigger a more thorough clinicopathological evaluation (serum DSA testing, tissue C4d staining, and exclusion of other causes of a similar type of injury) needed to establish the diagnosis with certainty.

MATERIALS AND METHODS

Case Selection & Study Design:

Previous University of Pittsburgh Medical Center (UPMC) studies examined the effect of a conventional lymphocytotoxic crossmatch on patient and allograft survival (7) and the utility of C4d staining in primary liver allograft recipients (27). As part of these prior studies a constellation of severe histopathological findings associated with acute AMR was described (27). These findings included microvascular (portal vein and portal capillary) endothelial cell hypertrophy; variable histiocytic, eosinophilic, and neutrophilic portal inflammation with microvasculitis; portal/periportal edema; cholangiolitis; centrilobular hepatocyte swelling; and hepatocanalicular cholestasis. AMR-related microvasculitis was defined as inflammatory cells adherent to, or near, the *luminal aspect of hypertrophied endothelial cells*, which differs from the *subendothelial* lymphocytic infiltration of portal and central veins seen in typical T-cell-mediated rejection. The goals of this study were: 1) to determine whether four pathologists from 3 different liver transplant centers in two continents could blindly recognize histopathological findings of severe AMR on H&E staining alone associated with diffusely C4d-positive putative AMR episodes, and 2) to develop and validate a simple, generalizable scoring system that would facilitate recognition and an earlier diagnosis of acute AMR in liver allografts. Cases of acute AMR were selected based on the

following criteria (24): 1) microvasculitis (as described above), 2) diffusely positive C4d-staining (>50% of portal tracks with positive C4d staining of the portal microvasculature with or without sinusoidal or central vein staining) , 3) elimination of other causes of a similar type of injury, and 4) DSA in serum. However, because of a lack of serum available for re-testing and the known insensitivity of cell-based cytotoxic assays, DSA was not required in the training cohort.

The training set consisted of UPMC for cause biopsies (n = 26) obtained within 21 days of primary liver transplant and divided into two groups: 1) those showing evidence of putative acute AMR with or without co-existent “cellular” rejection (n=13); and 2) an equally-sized group of control biopsies matched for Banff rejection grade severity (n = 13; indeterminate = 4; mild = 6; moderate = 3), but with negative C4d staining.

A single blinded pathologist (AJD) re-reviewed the H&E-stained slides without knowledge of the C4d results and evaluated 27 different histologic features. After the initial appraisal, several histopathological categories were combined and those with a p-value <0.3 or those with a strong pathophysiological basis for inclusion remained part of the final list of 9 variables (Table 3). Following selection of the histopathological variables, 3 additional pathologists (SMS, CB, and MAN) evaluated the training material without knowledge of the number of C4d-positive or C4d-negative cases in each group that originated outside their own institution or C4d staining results for all cases. Variables positively associated with putative AMR or mixed AMR and T-cell-mediated rejection in the training set were considered for inclusion in the numerator of the model

based on a p-value <0.2 . The denominator variables were selected for their negative association with AMR.

Multiple models were made from the training cohort variables based on the following guiding principles: 1) a scientific understanding of AMR, 2) simplicity, 3) the least inter-observer variability, and 4) the best correlation with C4d staining. The final model was selected for its lowest p-value from the training cohort data only.

Following evaluation of the UPMC training set, a separate validation cohort was created from 2 different centers: 1) Edinburgh University and 2) Baylor University Medical Center (BUMC). The Edinburgh University cases (between 2007 and 2013) were selected in a similar fashion to UPMC cases: a diagnosis of rejection within 21 days of transplant with histopathological evidence of rejection-related injury, strong and diffuse microvascular C4d staining, and fortunately a pretransplant positive cytotoxic or flow crossmatch or single antigen bead assay was available for all cases (n=5) and matched to a control group based on the Banff grade of cellular rejection with negative C4d staining and negative pre-transplant DSA testing (n=5).

The second portion of the validation cohort included all 29 HCV RNA-negative cases of biopsy-proven steroid-resistant rejection from BUMC within 60 days of liver transplantation with single antigen bead testing performed pre-transplant (from 1/1/00 to 5/31/09) (21). This approach was based on the unrealized expectation that the cohort would be enriched for recipients suffering from acute AMR (22), but only 4 stained diffusely positive for C4d and 1 showed focal positivity. C4d staining was performed at UPMC using the listed protocol. Three showing diffuse C4d positivity were also DSA positive and included in the final group; the remaining two cases: one originally

interpreted as diffuse had high background staining and one with focal C4d positivity were excluded because of equivocal C4d staining and negative DSA testing leaving a total of 27 cases. None of the DSA-negative cases had definitive diffuse C4d-positive staining.

To achieve adequate statistical power we combined the Edinburgh and BUMC cohorts into one validation cohort. The appraisal performed by all pathologists on this validation cohort was on the H&E material alone without knowledge of the C4d staining results. All biopsies were obtained before therapy for rejection was initiated.

C4d Staining Protocol

Deparaffinized and hydrated slides were treated with Target Retrieval Solution, pH=9.0 (DAKO, Carpinteria, CA) in a pressure cooker for 10 min to unmask antigens. After blocking slides, rabbit polyclonal anti-human C4d antibody [1:30-1:50, distributed by Alpco Diagnostics, Salem, NH in the United States and Biomedica, Austria in Europe (BI-RC4d)] was applied and incubated at 4°C overnight. After washing with PBS 3 times, biotinylated goat anti-rabbit antibody, VECTASTAIN Elite ABC kit (Vector Laboratory, Burlingame, CA) with: AEC Chromogen (Scytek Laboratories, Logan, UT), DAB Chromogen (Vector Laboratory) or Bond™ Polymer Refine Red Detection (Leica Biosystems, Buffalo Grove, IL) and Vector Red Alkaline Phosphatase Substrate kit (Vector Laboratory) was used for visualizing C4d staining and counterstained with hematoxylin for visualization.

Pretransplant DSA Evaluation

All UPMC patients had a pretransplant T-cell cytotoxic crossmatch performed prior to liver transplantation. In crossmatch-positive patients a steroid recycle was routinely given regardless of laboratory parameters, followed by standard per protocol immunosuppression. Neither pretransplant nor post-transplant serum was available for single antigen bead DSA analysis.

All Edinburgh University patients had pretransplant DSA testing performed since 2007; however the protocol has evolved: from 2007 to 2010 a cytotoxic T- and B-cell crossmatch, from 2011 to 6/2012 a flow cytometric crossmatch, and since 7/2012 all patients are screened for anti-HLA antibodies with multi-antigen beads to class I and class II antigens, with single antigen bead testing for DSA specificities in all positive patients.

All BUMC patients had prospectively collected pretransplant serum available for retrospective analysis of preformed DSA by single antigen bead technology, where mean fluorescence intensity (MFI) ≥ 5000 was considered positive, although data was acquired and reported on all DSA with MFI ≥ 1000 . All patients and donors were typed for HLA-A, -B, -DRB1, -DRB345 and -DQ using commercially available serologic typing trays or by molecular methods (Terasaki HLA tissue typing trays and Micro SSPTM or LabType[®] SSO, respectively; One Lambda Inc., Canoga Park, CA). All sera were blindly analyzed at the Terasaki Foundation Laboratory for HLA IgG antibodies using LABScreen single antigen class I (lot 6) and II (lot 8) beads (One Lambda Inc., Canoga Park, CA) according to the manufacturer's protocol. No serum was available to perform additional testing at the time of liver biopsy for any cohort.

Statistical Analyses

Patient characteristics for the 3 cohorts are reported with median values and interquartile ranges of continuous data and percentages of categorical data where appropriate. Chi squared analyses of categorical variables and two-sample t-tests of continuous variables were performed. Univariate logistic regression was utilized to evaluate individual variables and the model's ability to predict association with C4d positive rejection.

Although our final model produced a linear score, the output was not thought to be linearly associated with the ability to predict AMR. Therefore, we employed SPSS 16.0 to determine predictive cutoffs using tree classification. This was performed on the training cohort data from all 4 blinded pathologists before the validation cohort data was available for analysis and not modified after its completion.

Inter-observer variability was assessed with the Kendall's coefficient of concordance (33). Coefficient of concordance analyses were performed for each individual variable. This measure, unlike the Kappa statistic, is for ordinal values and takes into consideration the magnitude of disagreement between evaluators. For the final model the coefficient of concordance measured their agreement on the Acute-AMR (aAMR) category (≤ 1 , >1 but ≤ 1.75 and >1.75).

Significance was always defined as a $P < 0.05$. SAS 9.1 was used for all statistical analyses except SPSS 16.0 was utilized for tree classification.

RESULTS

Patient characteristics for the 3 cohorts are presented in Table 1. The cohorts were chosen differently because of local care standards, therefore, intergroup differences existed, but because of the blinded nature of analysis, none were felt to substantially influence the results.

Table 2A shows pretransplant T-cell cytotoxic crossmatch data from UPMC cases according to C4d staining; of the 13 diffusely C4d positive cases 38% were T-cell crossmatch positive, whereas the remainder was T-cell crossmatch negative. All C4d-negative cases/biopsies had a negative T-cell cytotoxic crossmatch except one. Table 2B shows the pretransplant DSA correlation with C4d staining in the Edinburgh cases. All C4d positive cases had evidence of pre-transplant DSA by either single antigen beads [class I MFI_{SUM} of 28,500 and class II MFI_{SUM} of 27,300] (n=1), T-cell flow crossmatch (n=1), or T-cell cytotoxic crossmatch (n=3). All C4d negative controls were also DSA negative by either single antigen bead analyses (n=4) or flow crossmatch (n=1). Table 2C shows the pretransplant single antigen bead data from the BUMC cases: only 3 cases were C4d positive with DSA in serum, and each one had at least one DSA with MFI \geq 5000. The first had a single class I DSA with MFI of 11,353, the second had 2 class I DSAs with MFI_{SUM} of 13,620 and 4 low MFI (all between 1000 and 4999) class II DSAs with MFI_{SUM} of 9,066, and the third had one class I DSA with MFI 1,868 and 5 class II DSAs with MFI_{SUM} of 62,375. None of the C4d negative cases had any single DSA with a MFI \geq 5000, but nine had low MFI DSAs (between 1000 and 4999); 2 with class I only, 4 with class II only and 3 with class I and II.

Table 3 highlights the 7 evaluated histologic characteristics associated with C4d positive early rejection or putative acute AMR (Figures 1-4): eosinophilic central

venulitis, portal vein endothelial cell hypertrophy, eosinophilic portal venulitis, central venulitis severity, portal eosinophilia, hepatocyte ballooning, and cholestasis; and 2 histologic characteristics inversely associated with acute AMR in the training cohort: lymphocytic portal inflammation and lymphocytic venulitis (Figure 2). Although cholestasis was not associated with DSA injury when all 4 pathologists scores were utilized in the training cohort, when particular attention was refocused to distinguish hepatocanalicular cholestasis from centrilobular hepatocyte lipofuscin deposition, an association was found in the validation cohort. In addition, the coefficients of concordance improved significantly after learning from the training cohort was followed by evaluation of the validation cohort.

Next, multiple models were created from the training cohort data alone, but Figure 5 shows the final Acute-AMR (aAMR) score. Numerical values are assigned based on the percentage of structures affected (None = 0, <10% = 1, 10-50% = 2, and >50% = 3). For the final model chosen, the OR was not appreciably changed from the training (OR=2.86, $P<0.001$) to the validation cohort (OR=2.49, $P<0.001$).

Next tree classification was utilized on the training cohort to optimize the specificity for one cutoff and sensitivity for the other cutoff of the aAMR score (Figure 5). Sensitivity in the validation cohort increased from 34% to 81% when the cutoff used decreased from >1.75 to >1 respectively. Specificity in the validation cohort also decreased from 87% and 71% when the cutoff used decreased from >1.75 and >1 respectively. In addition, the Kendall's coefficient of concordance between pathologists was 0.61 in the training and 0.50 in the validation cohorts.

DISCUSSION

Consensus histopathological criteria exist for the diagnosis of acute AMR in all solid organ transplants with the notable exception of the liver (5, 34-39), mostly because of its relative resistance to AMR and, consequent rarity of recognized cases. Absolute criteria for AMR in extra-hepatic organs invariably include serum DSA, microvascular endothelial cell hypertrophy and micro-vasculitis (40, 41), other tissue-specific injury patterns, and usually diffuse microvascular C4d staining. Kidney and heart allografts (41), however, allow for C4d-negative AMR when convincing microvasculitis is identified in the presence of DSA in serum.

Consensus criteria development for acute liver allograft AMR has been hampered by several issues, which, in turn, are related to the well-documented relative hepatic resistance to acute AMR: 1) in contrast to other solid organs, only a small fraction of DSA-positive liver allograft recipients develop overt histopathological evidence of injury (11, 17, 18, 24, 42); consequently 2) few programs routinely tissue type and screen for alloantibodies, or stain for C4d, mostly because they do not find it “cost effective”; and therefore, 3) only a few robust studies correlate histopathological findings, solid phase DSA testing, and C4d staining (18, 23, 24), and even fewer tissue biopsy and serum samples are simultaneously obtained.

Nevertheless, recent liver allograft studies confirmed and extended earlier observations by showing that high-titer DSA, in the presence of refractory thrombocytopenia, and diffuse microvascular C4d staining increase the probability of acute AMR (8, 23, 24, 43). The ability to correlate DSA with impaired outcomes,

however, remains suboptimal (17, 20, 21, 44) and more granular and specific histopathological criteria are needed.

Diffuse C4d positivity remains a critical component of an acute liver allograft AMR diagnosis at this time. However, C4d staining should not be interpreted in isolation (23, 24, 26, 27, 30) because C4d staining protocols for formalin-fixed, paraffin-embedded liver allograft tissue are evolving toward more sensitive techniques. In addition, correlation of staining results with liver dysfunction need improvement because even diffuse microvascular endothelial cell C4d deposits can occur with or without histopathological or serological evidence of liver injury [reviewed in (27-29, 45)]. Liver resistance mechanisms (listed above); more restricted hepatic microvasculature class II HLA expression compared to other organs; or the liver's position downstream from the intestine and complement activation by the lectin pathway, by bacterial products, and other factors all contribute to the complexities involved. Even so, most studies show a correlation between cell-based and often a stronger correlation with solid-phase evidence of DSA and tissue C4d staining [reviewed in (27-29, 45)]. A key consideration, therefore, is how to reliably recognize acute microvascular and perhaps stellate cell activation and injury from DSA in liver allografts?

In our opinion, the strong correlation between several histopathological features of microvascular activation (endothelial cell hypertrophy) and injury (microvasculitis) documented in a blinded analysis by 4 independent pathologists, as would be expected with AMR, and diffuse C4d staining and serum DSA in the validation cohort provide compelling evidence that antibodies substantially contribute to this injury pattern. The argument is further substantiated by the relative paucity of similar correlations in more

typical lymphocyte-predominant acute T-cell-mediated rejection biopsies matched for Banff grade of severity in controls.

It should be noted, however, that AMR-related microvasculitis is recognized primarily by increased *intra-luminal* inflammatory cells, some of which might be adherent to or apparently embedded within endothelial cells, and differs from the *subendothelial* lymphocytic infiltration of portal and central veins seen in otherwise typical T-cell-mediated rejection. Interestingly, some features originally attributed to T-cell-mediated rejection, such as an emphasis on a “mixed” inflammatory infiltrate consisting of activated and smaller lymphocytes, macrophages, neutrophils, and especially eosinophils (46), likely lumped together mixed T-cell-mediated and antibody-mediated effector mechanisms because of a lack of adequate tools to differentiate the two. Combined AMR and T-cell-mediated rejection is typical of many rejection episodes in all solid organ allografts. Therefore, changes attributable to AMR-related injury might be more difficult to isolate in livers simply because of convention.

Screening for acute AMR can be easily accomplished by using parameters included in the aAMR score on indication liver biopsies. We recommend using features identified in the overall score to screen for putative cases. Eosinophilia had the strongest correlation with acute AMR of any single histologic characteristic (HR = 4.37, $p < 0.001$). However, using the aAMR score in clinical practice, or specifically examining cases for AMR-associated features, should facilitate identification of the most severe cases of AMR. We opted, therefore, for high specificity and set a relatively high threshold aAMR score of >1.75 to raise significant concern for an acute AMR diagnosis. This approach is recommended because of potential consequences of AMR therapy

and to avoid over-diagnosis, which would inhibit widespread acceptance of a diagnosis that many already view with skepticism. Eventually, however, to improve sensitivity biopsies with scores >1 should be subjected to C4d staining and serum DSA testing should be carried out to substantiate or refute a putative AMR diagnosis. This will enable recognition of the entire spectrum of changes associated with AMR.

This study evaluated acute AMR at a more granular level than prior appraisals in an effort to help recognition of the most severe form of acute AMR. However, there are several shortcomings. One, training and validation cohorts were selected differently because of local standards of care. Two, there are no current Banff criteria for acute AMR, and therefore previous descriptions were used to select cases (27). Three, in the training cohort not all recipients with diffuse C4d-positive putative AMR showed pre-sensitization based on conventional T-cell cytotoxic crossmatches, which: a) miss most class II DSA; and b) are less sensitive (16) and can show substantially different results than solid phase assays when testing the same serum (17). The validity of this training cohort selection is substantiated by our BUMC patients in the validation cohort where a strong correlation between MFI of DSA and C4d staining was found: all patients with steroid resistant rejection and at least one DSA with MFI ≥ 5000 stained C4d positive, and all patients with steroid resistant rejection with lower MFI (1000 – 4999) DSA were C4d negative. Four, unavailability of simultaneous serum DSA testing and liver biopsy tissue hindered our ability to make tighter correlations. Five, part of our validation cohort was chosen from all the early (<60 days) steroid-resistant rejections that occurred in HCV RNA negative patients with pre-transplant DSA testing; this was done based on prior data showing this approach would enrich (41%) for C4d positive rejection (22),

however, only 11% of this group had C4d positive steroid resistant rejection. Finally, the histopathological changes shown in this manuscript represent only the most severe form of acute liver allograft AMR.

We attempted to mitigate most of these shortcomings by selecting cases from 3 different institutions, evaluating all material without knowledge of C4d or DSA test results, including 4 different pathologists, creating training and validation cohorts (the latter having solid phase DSA testing for most cases) and, relying on stringent criteria, including: 1) histopathological evidence of diffuse microvascular activation, injury, and microvasculitis; 2) diffuse microvascular C4d staining; 3) serum DSA (usually high MFI); and 4) reasonable exclusion of other causes of a similar type of injury (24). However, over time our understanding of acute AMR and C4d staining protocols will improve and molecular signatures of liver allograft AMR will be developed. As these advances unfold we expect that, like renal transplant pathology, histopathological features of acute and chronic liver AMR will be even more precisely defined, and C4d negative AMR will be described.

In summary, routine histopathological features in the aAMR score can be used to suspect the most severe form of acute AMR, a diagnosis that requires further substantiation by donor-specific HLA alloantibody testing, C4d staining, and exclusion of other insults. In the future, more subtle forms of DSA induced liver allograft injury will likely be discovered and described.

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Figure Legends.

Figure 1. Composite of early acute AMR histopathological changes in an allograft that failed 18 days after transplantation because of hepatic artery thrombosis in a highly sensitized patient. A) Note the monocytic and eosinophilic “capillaritis” in the peribiliary capillary plexus (arrows) surrounding a large segmental bile duct (BD) on H&E stain (40X). B) A C4d stain of the same area and throughout the entire liver showed diffuse endothelial cell C4d positivity (red *; 40X). C) Monocytic capillaritis was also noted in the smaller portal tracts (PT; 30X) (arrow shows area shown at higher magnification in the inset (80X). D) A C4d stain (red) showed strong and diffuse portal microvascular positivity, typical of severe acute AMR.

Figure 2. Composite histopathological features of severe, acute, C4d+ antibody mediated rejection (AMR) (A-F). A) Note intense and diffuse C4d staining (red) in the portal vein (PV) and portal capillaries (*). B) Routine H&E appearance of the same portal tract as shown in A). C) Shows a C4d stain (red) of the same portal vein (PV) branch as B) at higher magnification. D) Shows the routine H&E appearance of this vein. Note the marked portal venous endothelial cell hypertrophy of a tangentially sampled vein with eosinophils and histiocytes embedded within the hypertrophied endothelial cells. E) Another C4d staining example of the portal venous changes typical of severe acute AMR with the H&E counterpart shown in figure F). A scale bar is shown at the top left of each image.

Figure 3. High magnification (60X) H&E stain of the marked microvascular endothelial cell hypertrophy and cytoplasmic eosinophilia (arrows) that is typical of severe acute AMR. Note the cuboidal or “hobnail” appearance of the endothelial cells.

Figure 4. Inflammatory arteritis was present in several of the cases diagnosed as severe acute AMR, as in other solid organ allografts (H&E; 20X). This biopsy was obtained 11 days after transplantation from a 66-year-old female who underwent liver transplantation for primary biliary cirrhosis and hepatocellular carcinoma. Solid phase DSA determination revealed several class 1 and 2 DSA at a cumulative MFI >50,000. When arteritis is detected, C4d staining (inset) and DSA determinations are recommended. Arteritis, however, was not included in the acute AMR score because it is uncommonly detected in needle biopsies. Note the presence of lymphocytes, macrophages and eosinophils within the intima of the affected artery (large arrow), the endothelial hypertrophy in a nearby capillary (*). The inset (40X) shows C4d positivity (brown staining) in a portal capillary (*) and sinusoids (small arrow).

Figure 5: (A) The Acute-AMR (aAMR) score to predict antibody-mediated rejection was developed from 4 pathologists' scores on the training cohort and validated on a separate cohort. (B) The Odds Ratio demonstrates the aAMR model's association with a diagnosis of acute AMR on the training and validation cohorts. (C) SPSS16 tree classification developed diagnostic categories on the training set that were subsequently validated. Sensitivity and specificity of 2 different cutoffs are presented for

the training and validation cohorts; the higher cutoff optimizes specificity, while the lower cutoff optimizes sensitivity.

Accepted Article

Table 1: Patient characteristics of the training cohort from University of Pittsburgh Medical Center (UPMC) and the validation cohort from Edinburgh University and Baylor University Medical Center (BUMC) are seen. Medians with interquartile ranges are presented where appropriate.

	UPMC*	Edinburgh*	BUMC**
Number	26	10	27
C4d positive	13; 50%	5; 50%	3; 11%
Male Gender	46%	10%	59%
Age	55 (42-59)	55.5 (50-59)	53 (38-57)
Model for End-Stage Liver Disease	18 (14-23)	15.5 (11-16)	18 (14-24)
Cold Ischemia Time (hours)	9.8(8.1-12.4)	9.9 (7.7-13.2)	9.3 (5.6-11.1)
Hepatocellular Carcinoma	8%	30%	14%
HCV RNA Positive	31%	10%	0%
Recipient Race	Caucasian	96%	73%
	African-American	4%	10%
	Other	0%	17%
Donor Race	Caucasian	69%	62%
	African-American	27%	14%
	Other	4%	24%
Donor age	53 (41-74)	56 (49-67)	54 (39-61)
Induction	0%	0%	24%
Calcineurin [^]	100%	100%	81%
Steroids [^]	100%	100%	52%
Sirolimus [^]	0%	0%	26%
Mycophenolate [^]	38%	0%	44%

*C4d positive cases of rejection within 21 days of transplant were matched by Banff grade to C4d negative cases of rejection.

**All HCV RNA negative patients with steroid resistant rejection within 60 days of transplant from 1/1/00 to 5/31/09 who had a pre-transplant sample tested for donor-specific antibodies were stained for C4d.

[^] Immunosuppression at the time of rejection.

Table 2: Comparison of pre-transplant serological and C4d staining data for patients from (A) University of Pittsburgh Medical Center (UPMC); (B) Edinburgh University; and (C) Baylor University Medical Center (BUMC) (See Materials and Methods: case selection and study design). C4d staining was considered positive only when it was diffuse (>50% of portal tracts).

(A)

		UPMC
Total Patients/biopsies		26
C4d positive	Positive by cytotoxic T-cell crossmatch	5
	Negative by cytotoxic T-cell crossmatch	8
C4d negative	Positive by cytotoxic T-cell crossmatch	1
	Negative by cytotoxic T-cell crossmatch	12

(B)

		Edinburgh
Total Patients/biopsies		10
C4d positive	None	0
	Positive by class I & II SAB	2
	Positive T-cell by flow cytometry crossmatch	1
	Positive by T-cell cytotoxic crossmatch	2
C4d negative	Negative by Single Antigen Beads	4
	Negative by flow cytometry crossmatch	1

(C)

		BUMC*
Total Patients/biopsies		27
C4d positive	None	0
	Class I	1
	Class II	1
	Class I & II	1
C4d negative	None	24
	Class I	0
	Class II	0
	Class I & II	0

* Only cases with at least one individual donor-specific HLA Alloantibody with a MFI >5000 were considered positive.

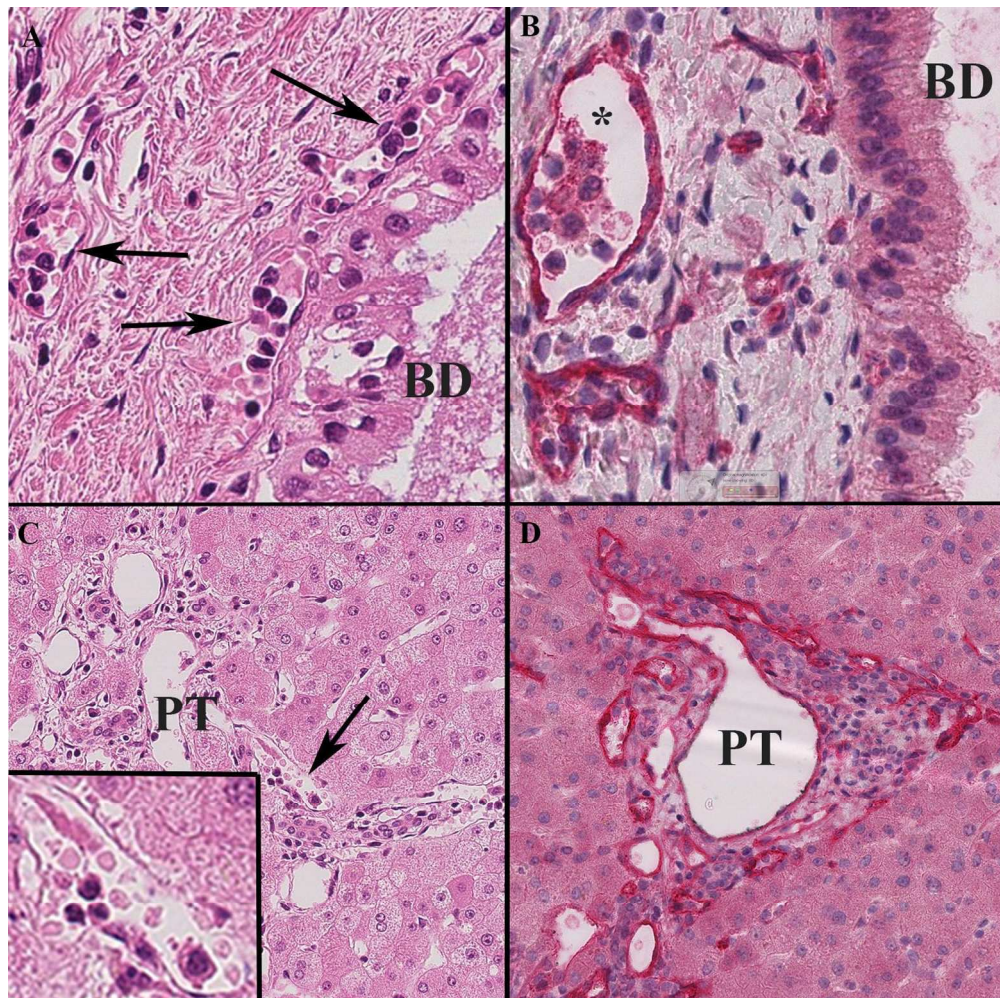
Table 3: In the (A) University of Pittsburgh Medical Center (UPMC) training cohort and (B) Edinburgh University and Baylor University Medical Center (BUMC) validation cohort, the odds ratios for all 4 pathologists (blinded to C4d results) for the 9 variables with the highest positive or inverse correlations with C4d positive rejection are displayed. Kendell's coefficient of concordance displays the inter-observer variability.

(A)

	Odds Ratio [CI]	P-value	Coefficient of Concordance
Eosinophilic Central Venulitis	1.93 [1.25-2.96]	0.003	0.49
Portal Vein Endothelial Cell Hypertrophy	1.89 [1.19, 2.99]	0.007	0.42
Eosinophilic Portal Venulitis	2.48 [1.24-4.96]	0.01	0.30
Central Venulitis Severity	2.26 [1.2-4.25]	0.02	0.33
Lymphocytic Portal Inflammation	0.59 [0.34-1.03]	0.06	0.40
Portal Eosinophilia	1.43 [0.92-2.21]	0.11	0.40
Lymphocytic Venulitis	0.78 [0.49-1.24]	0.3	0.32
Hepatocyte Ballooning	1.14 [0.76-1.72]	0.53	0.44
Cholestasis	1.00 [0.71-1.42]	1	0.29

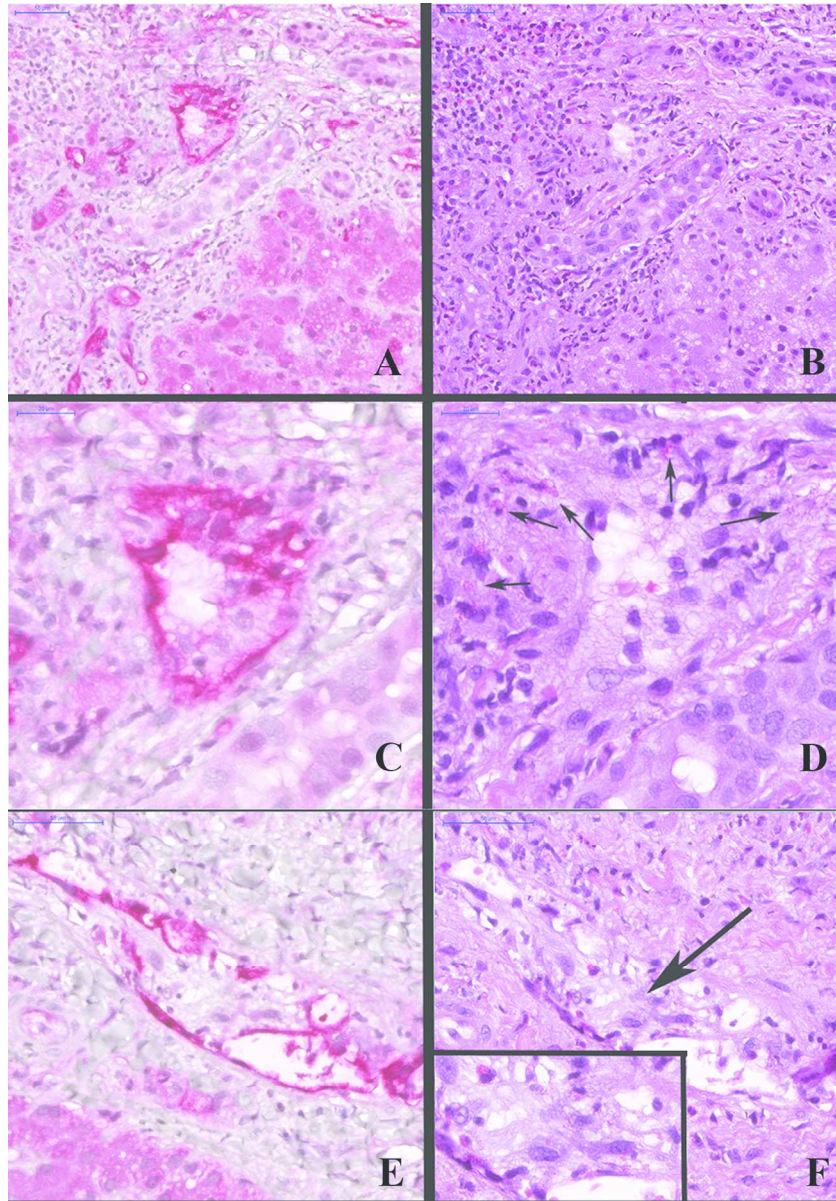
(B)

	Odds Ratio [CI]	P-value	Coefficient of Concordance
Eosinophilic Central Venulitis	2.48 [1.37-4.49]	0.003	0.63
Portal Vein Endothelial Cell Hypertrophy	2.88 [1.83-4.55]	<0.001	0.62
Eosinophilic Portal Venulitis	3.05 [1.96-4.69]	<0.001	0.38
Central Venulitis Severity	2.44 [1.47-4.06]	<0.001	0.63
Lymphocytic Portal Inflammation	1.33 [0.79-2.22]	0.3	0.58
Portal Eosinophilia	4.37 [2.54-7.51]	<0.001	0.61
Lymphocytic Venulitis	1.65 [1.05-2.58]	0.03	0.42
Hepatocyte Ballooning	2.00 [1.35-2.95]	<0.001	0.63
Cholestasis	2.09 [1.35-2.95]	<0.001	0.65



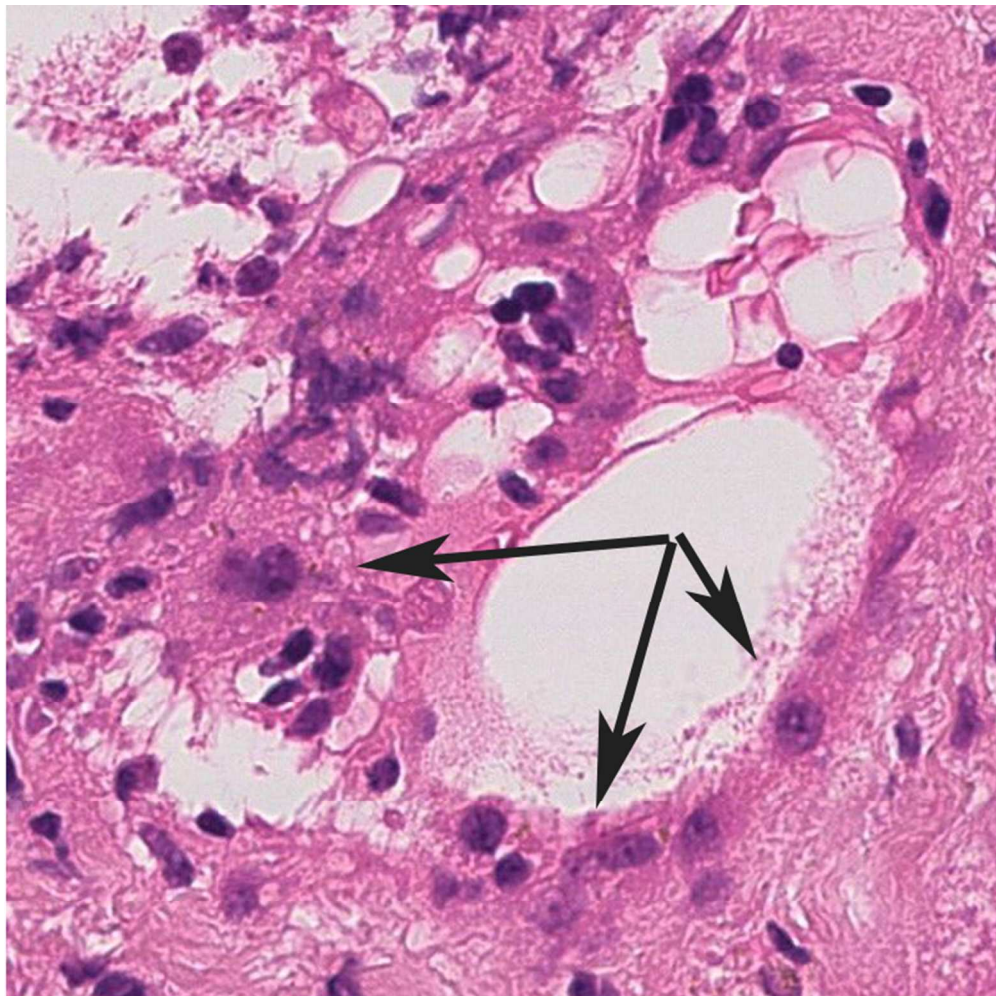
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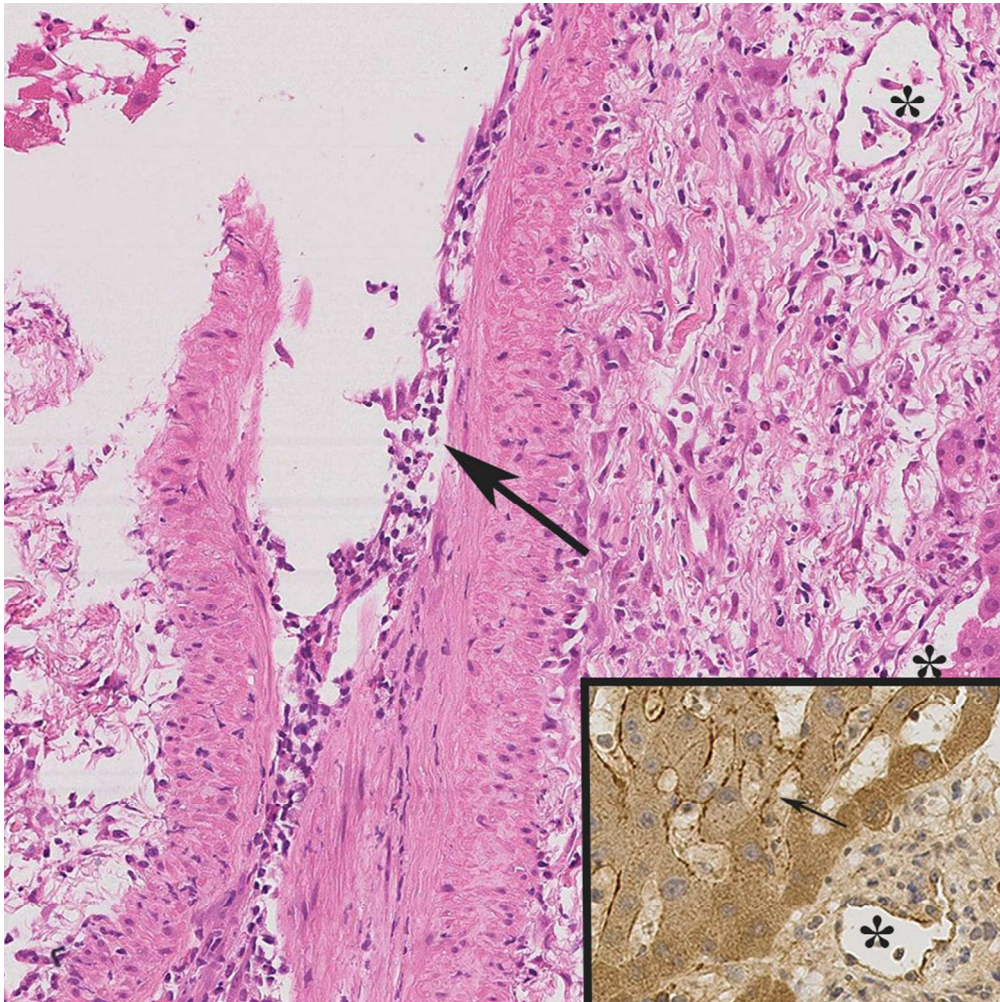
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A) Acute-AMR (aAMR) Score

Portal Vein Endothelial Cell Hypertrophy + Portal Eosinophilia + Eosinophilic Central Venulitis

Lymphocytic Portal Inflammation + Lymphocytic Venulitis

Number	% Affected Portal Tracts
None	0
1	<10%
2	10-50%
3	>50%

(B)

	OR	P-value
Training	2.86	<0.001
Validation	2.49	<0.001

(C)

		>1.75	>1
Training	Sensitivity	38% (20/52)	67% (35/52)
	Specificity	94% (49/52)	81% (36/52)
Validation	Sensitivity	34% (11/31)	81% (26/32)
	Specificity	87% (102/116)	71% (82/116)