REVIEW

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AIM associated with the IgM pentamer: attackers on stand-by at aircraft carrier

Toru Miyazaki^{1,2,3}, Tomoko Yamazaki¹, Ryoichi Sugisawa¹, M Eric Gershwin⁴ and Satoko Arai¹

Circulating immunoglobulin M (IgM) exists in a pentameric form, possessing a polyreactive nature that responds not only to foreign antigens but also to autoantigens; thus, it is involved in both beneficial and detrimental immune responses, including protection from infection and the progression of autoimmunity. On the other hand, IgM also behaves as a carrier of the apoptosis inhibitor of macrophage (AIM) protein, storing a large amount of the inactivated form of AIM in the blood through this association. Under different disease conditions, AIM can dissociate from IgM locally or systemically to exert its function, inducing the removal of various biological debris such as excess fat, bacteria, cancer cells or dead cell debris. Most typically, upon induction of acute kidney injury (AKI), IgM-free AIM is filtered by the glomerulus in the kidney, which stimulates the clearance of intraluminal dead cells debris at the obstructed proximal tubules, thereby facilitating the repair of kidney injury. Interestingly, cats exhibit a deficiency in AIM release from IgM, which may increase their susceptibility to renal failure. Conversely, association with AIM inhibits IgM binding to the $Fc\alpha/\mu$ receptor on follicular dendritic cells at the splenic germinal center, thereby protecting the IgM immune complex from $Fc\alpha/\mu$ receptor-mediated internalization, which supports IgM-dependent antigen presentation to B cells and stimulates high-affinity IgG antibody production. The regulation of AIM–IgM binding, resulting from the discovery of reciprocal actions between AIM and IgM, could lead to the development of novel therapies against different diseases.

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INTRODUCTION

Various biological cascades respond to alerting signals against life-threatening events, such as tissue injury. In most cases, the production of effector molecules is rapidly upregulated via an increase in the transcription of mRNA for the relevant molecules. Conversely, in some cases, effectors that have already been generated and stored within cells are rapidly released in response to more acute events; this action is typically observed in immune cells secreting cytokines during infection or Langerhans β cells emitting insulin in response to hyperglycemia. However, to conserve the time and energy required for effector activation, it may be more efficient to maintain effectors on stand-by in the blood and to release them on demand in an active form against the target, similar to

fighter jets being launched from a large aircraft carrier. Interestingly, we recently found that circulating immunoglobulin M (IgM) behaves as such a carrier and that its primary aircraft is the apoptosis inhibitor of macrophage (AIM) protein.

CHARACTERISTICS OF AIM

AIM, also known as CD5-like antigen (CD5L), is a circulating protein initially identified as an apoptosis inhibitor that supports the survival of macrophages against different types of apoptosis-inducing stimuli.¹ Serum AIM levels are relatively high (\sim 5 μg/ml) in humans and mice.^{1–3} AIM belongs to the scavenger receptor cysteine-rich (SRCR) superfamily, which all share a highly conserved cysteine-rich domain of \sim 100 amino

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acids[.4](#page-9-0) The AIM protein sequence is well conserved between humans and mice, with 78% amino-acid homology, but exhibits variations in its glycosylation state.^{1,5,[6](#page-9-0)} To the best of our knowledge, human and mouse AIM proteins are functionally equivalent[.6](#page-9-0) AIM is mainly produced by tissue macrophages, including liver Kupffer cells and peritoneal resident macrophages, and is transcriptionally regulated by nuclear receptor liver X receptor/retinoid X receptor heterodimers.^{7–9} Hamada *et al.*^{[10](#page-9-0)} reported that the transcrip-
tion factor MafB is also involved in the requlation of \overline{AM} tion factor MafB is also involved in the regulation of AIM mRNA expression[.10](#page-9-0) We analyzed serum AIM levels in more than 20 000 healthy human individuals and found that AIM levels are high in young women (teens to 20s) and gradually decrease with increasing age until \sim 50 years of age, after which they are fairly steady. In contrast, AIM levels in men are fairly steady at all ages, similar to those found in women over 50 years of age.³

AIM ASSOCIATES WITH THE IgM PENTAMER IN BLOOD AND IS PROTECTED FROM URINARY EXCRETION

In 2002, Tissot *et al.*^{[11](#page-9-0)} first reported that the blood IgM
immune complex (IC) contains AIM which was determined immune complex (IC) contains AIM, which was determined using mass spectrometric analysis of co-precipitates with IgM. Indeed, when we tested mouse and human sera for AIM by immunoblotting under non-reducing conditions, although the molecular weight of AIM is 37 kDa in humans and 42 kDa in mice, the AIM signal was only observed at a position identical to that of the pentameric IgM $(>500$ kDa).² Size fractionation of mouse serum also showed that both AIM and IgM were present in the same fractions, confirming that the majority of circulating AIM is associated with the IgM pentamer. We also determined that AIM binds to the Fc region of $IgM^{2,12}$ ([Figure 1a](#page-2-0)). Even the recombinant IgM-Fc, which lacks the variable region forming the pentamer, associates with AIM. Consistent with this, the association of AIM with different monoclonal IgM pentamers produced by B-cell hybridoma cells was achieved *in vitro*.^{[2](#page-9-0)} However, it is not currently clear
how AIM and the JaM pentamer associate with each other how AIM and the IgM pentamer associate with each other, mainly due to the difficulty of structural analysis of such a large complex. Additional efforts are required to elucidate this further.

Based on the calculated molarities of blood AIM and IgM in wild-type mice, each IgM pentamer appears to harbor only one to two AIM molecules; the circulating IgM pentamer therefore has more space for AIM. This observation is substantiated by the fact that when mice were injected with $\sim 10 \,\mu$ g of recombinant AIM (rAIM), most of the injected rAIM was bound to the IgM pentamer (Hiramoto and Miyazaki, unpublished results). Intriguingly, AIM cannot associate with the IgM pentamer in the absence of the J chain, which forms a complex with the IgM pentamer and possibly stabilizes the pentameric structure.[2](#page-9-0) Furthermore, the AIM–IgM pentamer association is abrogated in serum from mice deficient in the J chain[.13](#page-9-0) It is important to note that we confirmed that AIM does not bind directly to the J chain. It is possible that the

J chain may change the structure of the IgM pentamer to expose the AIM binding site within the Fc.

Interestingly, association with IgM increases the stability of the AIM protein in blood. The association prevents the urinary excretion of AIM, and this response results in the accumulation of AIM in the blood to a relatively high concentration $(-5 \mu g)$ ml). IgM-free AIM is easily filtered by the glomerulus due to its molecular size; therefore, serum AIM is largely lost in mice lacking circulating IgM, such as recombinase activating gene 1 deficient ($RAGI^{-/-}$) mice or secreted-type IgM-deficient (Δ sµ) mice[.14](#page-9-0) The serum AIM level was also decreased in J chaindeficient mice, as AIM does not bind to the IgM pentamer in the absence of the J chain. It is noteworthy that the macrophage AIM mRNA level is comparable in various tissues in all types of mice. In line with these findings, serum AIM levels rapidly increased when a monoclonal mouse IgM was intravethe absence of the J chain. It is noteworthy that the macro-
phage AIM mRNA level is comparable in various tissues in all
types of mice. In line with these findings, serum AIM levels
rapidly increased when a monoclonal mo strong correlation between AIM and IgM levels in the blood has been found in both humans and mice^{$2,3$} [\(Figure 1b\)](#page-2-0). In contrast, it is unlikely that AIM contributes to the protein nously injected into $\Delta s\mu$ or $RAG1^{-/-}$ mice.^{2,15} Consequently, a
strong correlation between AIM and IgM levels in the blood
has been found in both humans and mice^{2,3} (Figure 1b). In
contrast, it is unlikely that AIM showed comparable levels of blood IgM. Moreover, this correlation was not observed between AIM and IgG levels in the blood, in accordance with the fact that AIM does not bind to IgG.

AIM INFLUENCES DISEASE PROGRESSION IN AN IgM-FREE MANNER

Over the past decade, many research groups have clarified various AIM functions that essentially protect against different disease types. It is most noteworthy, however, that all the effects exhibited by AIM are achieved in the IgM-free form. When associated with the IgM pentamer, AIM is protected from renal excretion and is thus stabilized; however, it is inactivated [\(Figure 2](#page-3-0)).

Unlike many other soluble proteins, AIM does not mediate signal transduction in target cells but is incorporated by several cell types through scavenger receptor-mediated endocytosis.¹⁶ In obese people, AIM is released locally or systemically in certain circumstances and endocytosed into adipocytes and hepatocytes through CD36, where it binds to and inactivates cytoplasmic fatty acid synthase[.16,17](#page-10-0) This leads to a reduction in lipid droplet-coating proteins, such as fat-specific protein 27 and perilipin, thereby decreasing triacylglycerol deposition within the cells.¹⁷ This AIM action prevents obesity and fatty cytoplasmic fatty acid synthase.^{16,17} This leads to a reduction in lipid droplet-coating proteins, such as fat-specific protein 27 and perilipin, thereby decreasing triacylglycerol deposition within the cells.¹⁷ This (HFD), body weight gain is significantly greater than that in and perilipin, thereby decreasing triacylglycerol deposition
within the cells.¹⁷ This AIM action prevents obesity and fatty
liver progression (Figure 2). In $AIM^{-/-}$ mice fed a high-fat diet
(HFD), body weight gain is sig in visceral adipose tissue mass.[16](#page-10-0) Conversely, this hyperobese phenotype is abrogated by the administration of rAIM to obese $AIM^{-/-}$ mice. Similarly, $AIM^{-/-}$ mice fed an HFD show more (HFD), body weight gain is significantly greater than that in wild-type mice, and $AIM^{-/-}$ mice show a remarkable increase in visceral adipose tissue mass.¹⁶ Conversely, this hyperobese phenotype is abrogated by the admin advanced liver steatosis than wild-type mice, with increased liver mass and liver triacylglycerol content.^{[18](#page-10-0)} Note that IgM does not colocalize with AIM that is incorporated into the cytoplasm of adipocytes and hepatocytes, corroborating the fact that IgM-free AIM enters the target cells and functions within

Figure 1 AIM binds to the Fc region of the IgM pentamer in blood. (a) Schematic structures for IgM and AIM. Like other immunoglobulins, such as IgG, the IgM monomer consists of two heavy chains (blue or green) and two light chains (yellow), each of which are covalently connected by disulfide bonds. The immunoglobulin monomer is divided into two regions: a variable region required for recognition of specific antigens and a constant region, which is identical among all the immunoglobulins with the same isotype. The fragment crystallizable (Fc) region is a tail part of the constant region including CH2-CH4 domains and is important for various immunological responses through binding to Fc receptors expressed on the surface of immune cells. Generally, IgM exists as a pentamer in blood. Each IgM monomer is linked by disulfide bonds, and a J chain indispensably joins to this pentameric association. AIM, which harbors three SRCR domains, binds to the Fc region of the IgM pentamer. Although five AIMs may theoretically bind to one IgM pentamer (illustrated as an AIM–IgM complex), it is hypothesized that one or two AIMs associate with an IgM pentamer in blood. (b) A strong correlation in serum levels of AIM and IgM in men (left, $n=120$, average age 49 years) and in women (right, $n=119$, average age 49 years) was observed. The data are modified from a previously published report.³

Figure 2 Scheme of two different states of AIM in the body: dissociation from IgM improves activity but reduces stability, and vice versa. Under physiologic conditions, AIM associates with the IgM pentamer in the blood, which prevents its urinary exclusion due to its large molecular size (>500 kDa); however, IgM-bound AIM is thought to be functionally inactivated. Once AIM is dissociated, such as during AKI or other specific circumstances, AIM demonstrates different functions, as depicted in the right panel. However, IgM-free AIM is easily filtered by the glomerular membrane into the urine, and rapidly disappears from the body. The precise mechanism that causes the dissociation of AIM from IgM is not yet known.

the cell. In humans, we have observed significant negative correlations between circulating AIM levels and body mass index, abdominal circumference and body fat percentage, 3 demonstrating that circulating AIM dissociates from IgM and regulates cellular fat deposition, thereby preventing obesity and fatty liver.

Although AIM is incorporated into hepatocytes to decrease triacylglycerol deposition, once hepatocytes have undergone malignant transformation to hepatocellular carcinoma (HCC) cells, AIM is no longer endocytosed but instead accumulates on the cell surface^{[18](#page-10-0)} (Figure 2). This phenotypic change in AIM is possibly due to defective endocytosis, which is a common characteristic of many different types of cancer cells.^{19,20} Thus, in HCC cells, the IgM-free AIM accumulates on the cell surface after AIM-CD36 binding with insufficient AIM cellular incorporation; hence, AIM distinguishes HCC cells from normal hepatocytes. Interestingly, cell surface AIM specifically stimulates HCC cell death via necrosis through specific activation of a complement cascade against cancer cells, thereby preventing poration; hence, AIM distinguishes HCC cells from normal
hepatocytes. Interestingly, cell surface AIM specifically stimu-
lates HCC cell death via necrosis through specific activation of
a complement cascade against cancer HCC tumors when fed an HFD or a high-fructose diet for a year; in contrast, wild-type mice do not develop HCC tumors.^{[18,21](#page-10-0)} Overall, AIM prevents HCC tumor development through the elimination of cancerous hepatocytes. In addition, Lozano's group reported that AIM attaches to certain bacteria to induce coagulation, which aids effective bacterial eradication by the immune cells.²²

From these findings, it can be hypothesized that AIM is stored in blood by binding to the IgM pentamer and is released on demand in an active form to remove 'biological garbage', for example, excess fat, cancer cells or bacteria, to maintain a healthy state and overall organismal homeostasis.²³ Such garbage targeted by AIM also includes dead cell debris, and the adequate removal of such debris is important for maintaining tissue homeostasis (Figure 2). Several reports indicate that insufficient clearance of dead cells disturbs recovery from injury in various tissues, including the lung, heart, mammary gland and liver, primarily due to further inflammation caused by the debris.^{[24](#page-10-0)–28} Thus, in the liver, to avoid secondary inflammation and to promote tissue recovery, the HCC cells killed by AIM need to be rapidly removed. In fact, the accumulation of AIM on the surface of necrotic HCC cells appears to serve this purpose. We have found that an AIM coating profoundly increases the preference of phagocytes, including macrophages, for necrotic cell debris. Accordingly, when we induced the expression of cell surface-bound AIM in Hepa1.6 mouse HCC cells after these cells were transplanted into the liver of AIM−/[−] mice, a massive increase in HCC cell necrosis caused by activation of the complement system was observed specifically in transplanted HCC cells, followed by a rapid clearance of dead HCC cell debris by infiltrating Kupffer macrophages.[18](#page-10-0) This AIM-stimulated clearance of dead cell debris also appears to be involved in injury recovery by different tissues, including the kidney. We recently found that the efficient release of AIM from the IgM pentamer is a critical response in the recovery from acute kidney injury (AKI), as impaired AIMdebris also appears to be involved in injury recovery by different tissues, including the kidney. We recently found that the efficient release of AIM from the IgM pentamer is a critical response in the recovery from acute dramatically impaired their recovery from kidney damage.

PATHOLOGY OF AKI

AKI is characterized by impaired kidney filtration function, caused by insults such as kidney ischemia/reperfusion (IR) injury), sepsis or nephrotoxins. AKI is associated with prolonged hospitalization and high mortality and, most importantly, predisposes patients to chronic kidney disease and endstage renal disease.^{29–33} To date, although numerous therapeutic strategies have been proposed and tested for AKI, none have reached widespread clinical use. Renal tubular obstruction by debris from dead tubular epithelial cells is an important pathological characteristic often observed in AKI.34–[36](#page-10-0) Cell death in this tissue occurs due to apoptosis and necroptosis, `particularly in the proximal tubules at the corticomedullary junction; these dead cells detach from the tubular basement membrane physically obstructing the tubular lumen. These events reduce glomerular filtration and induce the production of inflammatory mediators by injured epithelial and infiltrating hematopoietic cells, further exacerbating tubular injury.³⁵ Thus, sufficient removal of intraluminal debris is a key mechanism for restoring tubular structure and renal function after AKI.

AIM DISSOCIATES FROM IgM AND IS EXCRETED IN THE URINE DURING AKI

During AKI, AIM is systemically released from the IgM pentamer in the blood, 37 resulting in urinary excretion of IgM-free AIM and allowing AIM to exert its ability to scavenge garbage. In sera from mice and humans with AKI, a substantial amount of IgM-free AIM $(<$ 40 kDa) was detected via immunoblotting. In contrast, healthy donors showed only a high molecular weight IgM-bound AIM ([Figure 3a](#page-5-0)). In human AKI patients, serum levels for IgM-free AIM increase in correlation with serum creatinine (Cre) levels. In accordance, the strong correlation between whole AIM and IgM levels in blood, which is observed in healthy human individuals, was absent in AKI patients. The IgM-free AIM appeared in the urine of AKI mice and humans, as determined by ELISA (enzyme-linked immunosorbent assay). AIM is not detectable in urine samples from healthy mice and humans but increases markedly in urine samples from individuals with an AKI attack. During the recovery phase, urinary AIM levels subsequently decline in both mice and humans. Consequently, the AIM excreted in urine coats the intraluminal debris that obstructs the proximal tubules. Immunohistochemistry analysis of the kidneys of AKI mice showed prominent accumulation of AIM on dead cell debris within the proximal tubules ([Figure 3b\)](#page-5-0). Similarly, in the kidneys of human AKI individuals, marked AIM accumulation on intraluminal debris was observed ([Figure 3b\)](#page-5-0). IgM did not colocalize with AIM at the intraluminal debris in mice and humans with AKI, confirming that the AIM deposited on the debris was IgM-free. In addition, AIM accumulation was not detected at the intraluminal debris in Δsμ mice in contrast to wild-type mice. This observation strongly supports the need for AIM to be stored in blood, through association with circulating IgM pentamer, to respond to emergency situations.

ABROGATED INTRALUMINAR DEBRIS REMOVAL AND IMPAIRED RECOVERY FROM AKI IN AIM-DEFICIENT MICE **ABROGATED INTRALUMINAR DEBRIS REMOVAL AND**
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shown by comparative analysis of wild-type and $AIM^{-/-}$ mice
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The role of AIM in the removal of intraluminal debris has been after the induction of AKI by IR^{37} On day 1 after IR, the structure of the proximal tubular cells was severely impaired in both wild-type and $AIM^{-/-}$ mice, and many proximal tubules located at the corticomedullary junction were obstructed with necrotic cell debris, which stained strongly with periodic acid-Schiff. In wild-type mice, the periodic acid-Schiff-positive intraluminal debris was reduced at day 3 post injury, whereas located at the corticomedullary junction were obstructed with
necrotic cell debris, which stained strongly with periodic acid-
Schiff. In wild-type mice, the periodic acid-Schiff-positive
intraluminal debris was reduced a at day 7. In parallel, there was no recovery of the brush border in the proximal tubules, a sign of healthy epithelial cells, which was apparent in wild-type mice at day 3. The differences in $AIM^{-/-}$ mice, the clearance of debris was not apparent even
at day 7. In parallel, there was no recovery of the brush border
in the proximal tubules, a sign of healthy epithelial cells, which
was apparent in wild-type m precisely in the acute tubular necrosis score, a standard method to evaluate the severity of kidney injury during AKI. In addition, as mentioned above, insufficient debris clearance caused sub-
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as mentioned above, insufficient de higher mRNA levels of various proinflammatory genes were to evaluate the severity of kidney injury during AKI. In addition,
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stitial areas in $AIM^{-/-}$ mice than in wild-type in
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Such a delay in the clearance of intraluminal debris, and the consequently, persistent inflammation and progressive fibrosis, mRNA levels of various fibrogenic genes were persistent in the kidneys of $AIM^{-/-}$ mice even at day 28 post injury.
Such a delay in the clearance of intraluminal debris, and the consequently, persistent inflammation and pro wild-type mice were subjected to IR, serum blood urea nitrogen and Cre levels, markers of renal dysfunction, peaked at day 1 and recovered thereafter. The general health status of the mice was lowest on day 1 and improved thereafter. The overall mortality of wild-type mice at day 7 was only 6.3% ([Figure 3c](#page-5-0)). In contrast, in and Cre levels, markers of renal dysfunction, peaked at day 1 and
recovered thereafter. The general health status of the mice was
lowest on day 1 and improved thereafter. The overall mortality of
wild-type mice at day 7 w levels, as well as their general health status, were comparable to wild-type mice on day 1, these parameters continued to worsen after day 1 and remained elevated at day 7. Ultimately, after IR, $AIM^{-/-}$ mice, although the serum blood urea nitrogen and Cre
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after day 1 and remained elevate levels, as well as their general health status, were comparable to wild-type mice on day 1, these parameters continued to worsen after day 1 and remained elevated at day 7. Ultimately, after IR, more than 40% of $AIM^{-/-}$ mi wild-type mice on day 1, these parameters continued to worsen
after day 1 and remained elevated at day 7. Ultimately, after IR,
more than 40% of $AIM^{-/-}$ mice died by day 3, and the mortality
at day 7 was 67% (Figure 3c). T serum Cre and blood urea nitrogen levels, as well as animal mortality in response to IR, were also markedly greater in the circulating IgM-less Δsμ mice than in wild-type mice. Thus, the storage of sufficient AIM by the IgM pentamer in blood and its release and subsequent translocation of AIM in urine are required for efficient repair of AKI. The cellular and molecular mechanisms of how AIM accumulation induces the removal of intraluminal debris were clarified but are not described here (see Arai et al^{37} al^{37} al^{37}). We would like to mention, however, that the administration of rAIM (in an IgM-free form) can facilitate the clearance of of how AIM accumulation induces the removal of intraluminal debris were clarified but are not described here (see Arai *et al.*³⁷). We would like to mention, however, that the administration of rAIM (in an IgM-free form well as in wild-type mice that have undergone severe AKI but in which the level of endogenous AIM was not sufficient [\(Figure 3d](#page-5-0)).

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Figure 3 AIM in AKI. (a) Immunoblotting in non-reducing conditions for AIM involving sera from healthy and AKI-induced wild-type mice $(n=2 \text{ each})$, as well as healthy individuals $(n=3)$ and AKI patients $(n=5)$. IgM-bound AIM and IgM-free AIM are indicated by arrows. IgM-free AIM is observed only in AKI conditions, and IgM-bound AIM is reduced with AKI in mice. (b) Immunohistostaining of the kidney specimens of healthy and AKI mice (upper panels), as well as a healthy individual and an AKI patient (lower panels) for AIM. Signals were visualized by horseradish peroxidase/3'-diaminobenzidine (HRP/DAB). Scale bars: 50 μm. Modified from a previously published report.^{[2](#page-9-0)} (c) Survival of wild-type (AIM^{+/+}) and AIM^{-/-} mice after IR was analyzed using the Kaplan–Meier method. P: statistical significance between AIM^{+/+} and AIM^{-/-} mice was calculated by using the log-rank test. Modified from our previous report.^{[2](#page-9-0)} (d) Schema for the involvement of AIM in AKI pathogenesis. (Left) When sufficient IgM-free AIM dissociates from IgM pentamer during AKI (this usually occurs in wild-type mice and most humans), AIM is excreted in the urine, accumulating on the lumen-obstructing necrotic cell debris and thereby enhancing its clearance by injured proximal tubular epithelial cells (indicated by red cells), leading to the regeneration of epithelial cells (indicated by blue cells) and AKI recovery. (Middle) In the absence of AIM (for example, AIM-deficient mice), the debris removal is deficient. (Right) In cats, due to the strong binding of feline AIM to IgM, AIM is unable to dissociate from IgM during AKI, abolishing its excretion in urine; the intraluminal debris therefore cannot be removed efficiently. Intravenous rAIM (which is IgM-free) administration could be therapeutically applied for AKI treatment.

FELINE AIM NEVER DISSOCIATES FROM IgM

The importance of AIM release from IgM in facilitating recovery from AKI is more directly highlighted in cats. The number of pets is increasing worldwide due to the

recently decreasing birth rate and increasingly greater age of the human population, and cats are the most popular pet in the vast majority of populations.^{38,39} Cats are profoundly more susceptible than other animals to, and more often die

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from, chronic kidney disease. $40-43$ However, the reason for their susceptibility to renal disease, which is one of the most pressing questions in veterinary medicine, has remained unclear. Based on the important involvement of AIM in the resolution of AKI described above, we evaluated feline AIM to assess whether it is related to the susceptibility of cats to renal disease. resolution of AKI described above, we evaluated feline AIM to
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cats might be geneti

In fact, before the start of this study, we had expected that however, they do possess AIM. After feline AIM cDNA was isolated, feline AIM protein was produced, and several antibodies specifically recognizing feline AIM were generated, serum AIM levels were assessed by immunoblotting in cats of different strains[.44](#page-10-0) The average AIM level in cats was 21.2 μg/ ml (in $>$ 30 cats), which is markedly higher than that in mice $(2-3 \mu g/ml)$ and humans $(\sim 5 \mu g/ml)$. In blood, feline AIM associates with IgM pentamers at the Fc region, and serum levels of AIM and IgM are positively correlated, as in humans and mice. $2,3$

Surprisingly, however, when cats are subjected to IR for experimental induction of AKI,[45](#page-10-0) all IR cats showed no IgMfree AIM in the serum, in contrast to humans and mice in which a substantial level of IgM-free AIM was detected during AKI via immunoblotting. This was also the case when sera from several hospitalized cats with spontaneous AKI caused by various events, such as renal stones, were analyzed; no IgM-free AIM was observed in sera from these cats. In humans and mice, AIM is not detectable in urine samples from healthy individuals, and its levels increase during AKI.³⁷ Consistent with the serum results above, AIM was not detectable in the urine of IR AKI cats, indicating no urinary AIM excretion during AKI. Furthermore, histological analysis demonstrated no AIM accumulation in the intraluminal debris of cats after IR, in contrast to humans and mice with AKI wherein AIM accumulation is observed in the debris. Thus, in cats, AIM does not dissociate from the IgM pentamer upon AKI induction, and is therefore not excreted in the urine.

To determine whether this inability to dissociate is due to feline AIM but not feline IgM, we generated transgenic mice expressing feline AIM under the regulation of the mouse AIM promoter on a mouse AIM−/[−] background, thereby felinizing AIM in mice. The transgenic feline AIM protein associated with mouse IgM pentamers, and the serum levels of feline AIM, in AIM-felinized mice, were comparable to those of murine AIM in wild-type mice $(\sim 2 \,\mu\text{g/ml})$ when assessed by immunoblotting. As expected, IgM-free AIM was not detected after IR in AIM-felinized mice, as observed in cats. Consistent with this result, immunohistochemical analysis of the kidneys from IR AIM-felinized mice revealed that the intraluminal debris did not stain for AIM, in sharp contrast to the AIMpositive debris in wild-type mice, clearly demonstrating that AIM did not appear in urine after IR in AIM-felinized mice. Therefore, feline AIM has a defect that disturbs its release from IgM upon AKI induction.

A STRONG BINDING AFFINITY OF FELINE AIM WITH IgM

We used surface plasmon resonance analysis to quantitatively compare the binding affinity constant of IgM-Fc pentamers in feline and murine AIM. The resulting dissociation–association rate (K_D) was 5.97 × 10⁻⁹ M in feline AIM/IgM-Fc, whereas in mouse AIM/IgM-Fc, it was 5.82×10^{-6} M; thus, in cats, AIM associates with IgM \sim 1000-fold more strongly than in mice. The binding affinity constant between feline AIM and murine IgM-Fc pentamers was also high, comparable to that in feline AIM/feline IgM-Fc, further suggesting that the strong association is caused by feline AIM, not by feline IgM. The strong binding of feline AIM with IgM likely plays a role in the increasing serum AIM levels observed in cats.

Recently, we found that the third SRCR domain (SRCR3) is responsible for IgM binding, since its deletion abrogated the association with IgM in murine, human and feline AIM[.15,44](#page-10-0) Of note, three-dimensional mapping of the charge distribution of amino acids revealed that the SRCR3 domain of feline AIM possesses a specific cluster of positively charged amino acids on the surface of the molecule that is not observed in mouse and human AIM, and this feline AIM-specific cluster appeared to be involved in the increase in its binding affinity with IgM. Indeed, using a mutant feline AIM in which feline-specific arginine residues constituting the positively charged aminoacid cluster were substituted with the corresponding amino acids of a mouse sequence, the mutant feline AIM was found to exhibit decreased binding to the Fc pentamers, which was verified biochemically via immunoblotting.^{[44](#page-10-0)} This finding highlights the crucial involvement of the positively charged amino acids of complement C1q in association with IgM-Fc.⁴⁶ It is possible that additional parameters defining the binding affinity between AIM and IgM are present, as the binding potential decreased only partially in the mutant feline AIM. Further studies including structural assessments of the AIM/ IgM-Fc complex are required to clarify the complete nature of the AIM–IgM association.

SAVING CATS FROM KIDNEY DISEASE USING AIM

The apparent inability of feline AIM to dissociate from IgM appears to be an important cause of the susceptibility of cats to renal failure. When wild-type and AIM-felinized mice were subjected to an IR protocol to induce AKI, serum Cre levels peaked at day 1 and recovered thereafter in wild-type mice, and 75% of mice survived to day 5. In complete contrast, serum Cre levels in AIM-felinized mice increased to similar levels as in wild-type mice at day 1 after IR but continued to increase thereafter, resulting in the death of 100% of mice by day 3.[44](#page-10-0) By histology, on day 3 after IR, there was significantly less intraluminal debris at the proximal tubules located at the corticomedullary junction in wild-type mice than in AIM-felinized mice. Similarly, recovery of the brush border in the proximal tubules was apparent on the same day in wild-type mice but not in AIM-felinized mice. The overall acute tubular necrosis score at day 3 was significantly worse in AIM-felinized mice than in wild-type mice. Overall, as expected, AIM-felinized mice showed impaired recovery from

Figure 4 AIM retains IgM immune-complexes on splenic FDCs. (a) Immunohistostaining of specimens of WT mouse spleens for AIM (green), IgM (blue) and FDC (red). Nuclei were also stained by DAPI (4',6-diamidino-2-phenylindole) (light blue). FDC areas are indicated by white arrows. The data are modified from our previously published report.² (b) Schema describing the role of AIM in retention of IgM-IC on the surface of FDCs. For the affinity maturation of GC B cells to plasma cells producing high-affinity IgGs, it is necessary that ICs are retained on the surface of FDCs for a sufficient period of time to present antigens to GC B cells. Although IgM-IC is retained by complement receptors (CRs), in the absence of AIM, it is internalized via Fcα/μR into FDCs and rapidly disappears from the surface, resulting in the failure of affinity maturation (right). When AIM is associated with IgM-IC, AIM prevents its internalization, supporting the retention of IgM-ICs on FDCs, which enables affinity maturation of GC B cells recognizing autoantigens that are presented by IgM-ICs (left).

AKI, similar to $AIM^{-/-}$ mice. Cats that underwent experimental IR (described above) exhibited poor recovery from AKI, as determined by the serum Cre levels and blood urea nitrogen. As implier to $AM^{-/-}$ mice. Cats that underwent
perimental IR (described above) exhibited poor recovery
pm AKI, as determined by the serum Cre levels and blood
ea nitrogen.
As in $AM^{-/-}$ mice, the administration of rAIM fac

AKI recovery in AIM-felinized mice. We injected mouse rAIM intravenously into AIM-felinized mice on days 1, 2 and 3 after IR. The survival of rAIM-injected mice at day 3 was markedly improved to 80% from 0%[.44](#page-10-0) The serum Cre level in rAIMinjected felinized mice decreased by day 5, whereas in nontreated AIM-felinized mice, it continued to increase until the

mice died. Using histology, the acute tubular necrosis score of rAIM-injected AIM-felinized mice was improved at day 5. We also confirmed using immunohistochemistry that the injected rAIM accumulated at the intraluminal debris. These observations strongly suggest the therapeutic potential of rAIM in cats ([Figure 3d\)](#page-5-0).

AIM RETAINS THE IgM-IC ON SPLENIC FOLLICULAR DENDRITIC CELLS

Thus far, we have discussed the importance of the efficient take-off of the aircrafts (AIM) from the aircraft carrier (IgM) to

protect against various diseases, particularly AKI. We will now turn to describing the potential influence of the aircrafts (AIM) on the aircraft carrier (IgM). It is well known that a large proportion of natural IgM is polyreactive, not only to foreign antigens but also to phylogenetically conserved autoantigens, and this is believed to be important for the progression of autoimmunity[.47](#page-10-0)–⁴⁹ The IgM-IC with an antigen, also containing complement component C3, is deposited on splenic follicular dendritic cells (FDCs) through the interaction between the activated C3 and the complement receptor (CD21/CD35) on $FDCs^{50-54}$ $FDCs^{50-54}$ $FDCs^{50-54}$ where the antigen is presented to germinal center (GC) B cells, inducing the development of plasma cells that produce high-affinity IgGs.⁵⁴ Therefore, increased IgM results in more chances to present autoantigens, thereby stimulating the autoantibody responses responsible for the progression of autoimmunity.

We asked whether the association of AIM might influence this process. Consistent with the association between AIM and the IgM pentamer, the accumulation of both AIM and IgM was specifically observed in FDCs within splenic GCs in mice² [\(Figure 4a](#page-7-0)). When we injected rAIM, alone or in combination with IgM, into Δsμ mice, rAIM accumulation in FDCs was only observed when rAIM was injected in combination with IgM, suggesting that the AIM accumulation in FDCs is IgMdependent.

Although the protein stability of IgM was not affected by the absence of AIM in blood, the AIM association influences the holding state of the IgM-IC at FDCs. We intravenously injected IgM alone or in association with rAIM (IgM/AIM) into Δsμ AIM−/[−] doubly deficient mice, and the presence of IgM at the FDC cell surface was tested kinetically. The IgM level at FDCs was profoundly increased when mice were injected with IgM/ AIM compared with IgM alone. Similarly, when the 2,4,6 trinitrophenyl antigen, which was shown to be bound to IgM,⁵⁵ FDC cell surface was tested kinetically. The IgM level at FDCs was profoundly increased when mice were injected with IgM AIM compared with IgM alone. Similarly, when the 2,4,6-trinitrophenyl antigen, which was shown to be trinitrophenyl antigen on FDCs was maintained longer in wildwas injected into wild-type and $AIM^{-/-}$ mice, the level of 2,4,6-
trinitrophenyl antigen on FDCs was maintained longer in wild-
type mice than in $AIM^{-/-}$ mice. Thus, the binding of AIM to IgM increases the retention of IgM-IC on FDCs.

AIM DISTURBS THE BINDING BETWEEN IgM AND THE Fcα/μ RECEPTOR

The Fc α/μ R, the Fc receptor for both IgM and IgA,⁵⁶ mainly found on FDCs,^{55,[56](#page-10-0)} induces internalization of IgM, thereby reducing the retention of IgM on the cell surface.⁵⁶ Indeed, **Fcα/μ RECEPTOR**
The Fcα/μR, the Fc receptor for both IgM and IgA,⁵⁶ mainly
found on FDCs,^{55,56} induces internalization of IgM, thereby
reducing the retention of IgM on the cell surface.⁵⁶ Indeed,
 $Fca/\mu R^{-/-}$ mice de FDCs when challenged with the 2,4,6-trinitrophenyl antigen.⁵⁵ These facts led to the hypothesis that the association between AIM and IgM-Fc might affect the binding of IgM to the Fcα/ μR at FDCs, which was found to be true. When HEK293T cells overexpressing Fcα/μR were incubated with a monoclonal IgM with or without rAIM association, the presence of AIM profoundly decreased the binding of IgM to $Fc\alpha/\mu R^2$ $Fc\alpha/\mu R^2$ Similarly, Fcα/μR-expressing HEK293T cells incubated with serum from wild-type mice showed reduced surface staining for IgM compared with cells incubated with AIM−/[−] mouse serum. Consistent with these binding results, internalization of IgM by

HEK293T cells expressing Fcα/μR was markedly decreased by the rAIM association. Thus, AIM interferes with the binding of IgM to Fc α/μ R, and its internalization through Fc α/μ R. Such physical disturbance of the IgM/ Fcα/μR association by AIM was also observed in the activation of the complement cascade against HCC cells by AIM that accumulates on the HCC cell surface. On the cell surface, multiple regulators of complement activation (including CD55, complement regulator complement receptor 1-related gene/protein-y, complement factor H and CD59) protect cells from complement-dependent attack by suppression of C3 activation or formation of the membrane attack complex[.57](#page-10-0)–⁵⁹ Therefore, AIM interferes with the complement regulatory activity of the regulators of complement activation through direct association with regulators of complement activation, thereby triggering complement activation and leading to specific necrotic death of the HCC cells^{[18](#page-10-0)} [\(Figure 2](#page-3-0)).

OBESITY-ASSOCIATED INCREASES IN CIRCULATING AUTOANTIBODIES AND THEIR PREVENTION IN AIM−/[−] MICE

It is possible, therefore, that the IgM-dependent antigen presentation to GC B cells at splenic FDCs is less efficient in **AUTOANTIBODIES AND THEIR PREVENTION IN AIM⁻¹-
MICE**
It is possible, therefore, that the IgM-dependent antigen
presentation to GC B cells at splenic FDCs is less efficient in
 $AIM^{-/-}$ mice than in wild-type mice due to a nalization of IgM-IC via Fcα/μR ([Figure 4b](#page-7-0)). This was demonstrated in the obesity-associated autoantibody production process. Accumulating etiological and clinical studies in humans have shown a strong correlation between obesity and autoimmune diseases. These are largely accompanied by increased levels of autoantibodies such as diabetes-associated antibodies against pancreatic β-cell antigens, including insulin, glutamic acid decarboxylase and protein tyrosine phosphataselike protein (IA2); chronic thyroiditis-associated anti-thyroid peroxidase or anti-thyroglobulin antibody; or infertilityassociated anti-sperm antibody.^{60–64} Winer *et al.*^{[65](#page-11-0)} also recently demonstrated the production of pathogenic JgG antibodies demonstrated the production of pathogenic IgG antibodies, including a unique profile of autoantibodies in obese humans and mice. Such antibody production was shown to be important in the acceleration of insulin resistance.

When mice are fed an HFD, the increased blood fatty acids stimulate the cell surface Toll-like receptor 4 on splenic marginal zone B cells, a major producer of natural IgM , $66-68$ and subsequently induce high amounts of polyclonal IgM production in an antigen-independent manner[.69,70](#page-11-0) Owing to its self-reactive nature, the augmentation of natural IgM stimulates IgG autoantibody production in obese mice. As expected, the serum from mice fed an HFD for 12 weeks contained significantly increased levels of IgG autoantibodies to more than 30 variable autoantigens related to DNA, U1RNP, histone, SSA/SSB and the cell matrix compared with lean mice when evaluated using a proteome microarray containing 70 contained significantly increased levels of IgG autoantibodies to
more than 30 variable autoantigens related to DNA, U1RNP,
histone, SSA/SSB and the cell matrix compared with lean mice
when evaluated using a proteome micr fed an HFD for the same period revealed markedly lower levels of IgG antibodies against most of the autoantigens to which wild-type mouse serum responded.^{[2](#page-9-0)} Consistent with these observations, the number of long-lived plasma cells (intracellular-IgG+/B220 [−]/IgM−/CD5−/Mac1[−])[,74](#page-11-0) which produce high-affinity IgGs, was decreased in the bone marrow (intracellular-IgG⁺/B220⁻/IgM⁻/CD5⁻/Mac1⁻),⁷⁴ which produce high-affinity IgGs, was decreased in the bone marrow of obese $AlM^{-/-}$ mice compared with obese wild-type mice.² Thus the absence of AIM abrogates t Thus, the absence of AIM abrogates the IgM-dependent maturation of high-affinity IgG-producing plasma B cells and thus prevents obesity-associated autoantibody production. It is well known that cats possess a number of autoantibodies and are therefore susceptible to various autoimmune diseases[.75](#page-11-0) It is possible that the extremely strong binding of feline AIM to IgM might be involved in the increased autoimmune nature of cats, which may be worth further assessment.

FUTURE DIRECTIONS

In this review, we have illustrated the significance of the association of AIM with the IgM pentamer in blood in terms of disease protection, particularly in kidney disease. AIM harbors a number of unique characteristics. First, AIM is a circulating protein that can be delivered throughout the body. Second, AIM possesses a 'sticky' nature, which enables it to efficiently accumulate on the surface of widely varying 'biological garbage'. This appears to be dependent on its cysteine-rich domains and is therefore a common feature shared by the members of the SRCR superfamily.4[,76](#page-11-0),[77](#page-11-0) Third, AIM is incorporated via endocytosis into cells through scavenger receptors, and does not mediate signal transduction, which is advantageous for the promotion of efficient garbage clearance. The storage of a relatively large amount of AIM in blood through binding to the IgM pentamer is likely beneficial for these AIM roles. The level of AIM in blood is high, as opposed to the levels of soluble signaling proteins, which are far lower.

To the best of our knowledge, the dissociation of AIM from IgM occurs during specific diseases; AKI causes the most prominent release. However, if we could release the stored AIM on a regular basis, it might flush away accumulating garbage that is still small enough not to cause any serious symptoms in different tissues, possibly preventing multiple chronic, progressive diseases, 78 even aging. To achieve this, we need to clarify the release mechanism, which must occur along with the massive dissociation of AIM from the IgM pentamer. We speculate that materials that can effectively control the release of AIM from IgM may represent a novel therapeutic concept and tool for various diseases in the future, although significant work is necessary to achieve this goal. One could expect that forcing the dissociation of AIM from IgM might also prevent autoimmunity via abolishing autoantigen presentation at the splenic GCs. Thus, it is likely that humans may gain considerable benefit from AIM-releasing materials. For cats, regretfully, this might be not the case, as feline AIM did not dissociate even under AKI; the binding between AIM and IgM appears to be too robustly covalent. We may need to use AIM itself or seek alternative treatments for cats. We do, however, propose that there will be significant relationships between AIM and human immunopathology. We note, for example, that patients with primary biliary cholangitis (PBC) have elevated serum IgM. This is relatively unique in human autoimmunity. It is more common to have polyclonal B-cell activation with subsequent

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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