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Peripheral complement interactions with amyloid β peptide ($A\beta$) in Alzheimer's disease: 2. Relationship to $A\beta$ immunotherapy

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Abstract

INTRODUCTION—Our previous studies have shown that amyloid β peptide ($A\beta$) is subject to complement-mediated clearance from the peripheral circulation, and that this mechanism is deficient in Alzheimer's disease (AD). The mechanism should be enhanced by $A\beta$ antibodies, which form immune complexes (ICs) with $A\beta$, and therefore may be relevant to current $A\beta$ immunotherapy approaches.

METHODS—Multidisciplinary methods were employed to demonstrate enhanced complement-mediated capture of $A\beta$ ICs compared to $A\beta$ alone in both erythrocytes and THP1-derived macrophages.

RESULTS— $A\beta$ antibodies dramatically increased complement activation and opsonization of $A\beta$, followed by commensurately-enhanced $A\beta$ capture by human erythrocytes and macrophages. These *in vitro* findings were consistent with enhanced peripheral clearance of intravenously-administered $A\beta$ ICs in non-human primates.

DISCUSSION—Together with our previous results showing significant AD deficits in peripheral $A\beta$ clearance, the present findings strongly suggest that peripheral mechanisms should not be ignored as contributors to the effects of $A\beta$ immunotherapy.

Keywords

Alzheimer's disease; amyloid β peptide; $A\beta$ immunotherapy; complement; complement receptor 1; macrophage; erythrocyte; blood; human

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CONFLICTS OF INTEREST: None

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1. Background

Over the last decade, the most visible strategy for the treatment of Alzheimer's disease (AD) has been amyloid β peptide ($A\beta$) immunotherapy [reviewed in 1]. Although the first efforts with $A\beta$ immunotherapy failed to complete clinical trials [1], AD transgenic mice [2,3] and human AD patients [4,5] did show significantly reduced $A\beta$ burden after treatment. Perhaps as a result, $A\beta$ immunization approaches continue to be pursued [1].

A critical, unresolved issue with $A\beta$ immunization is whether or not its presumed mechanism of action, enhanced glial clearance of brain $A\beta$ [e.g., 6–8], provides a sufficient explanation for its reported effects. For example, an $A\beta$ antibody, m266, that did not react with brain $A\beta$ deposits and appeared to have most if not all of its effect in the periphery, nonetheless reduced brain $A\beta$ levels in a transgenic AD mouse model [9]. This antibody formed immune complexes (ICs) with $A\beta$ in the peripheral circulation [10] and appeared to induce efflux of brain $A\beta$ to plasma [9,11], leading to the “peripheral sink” hypothesis [9–11]. Moreover, the penetration of $A\beta$ antibodies into the CNS remains open to debate. Levites and colleagues [12], for example, reported that only 1 fmol/mg of $A\beta$ antibody could be detected in AD transgenic mouse brain after a 500 μ g intraperitoneal injection. This brain concentration of antibody is nearly 3 orders of magnitude less than estimates of total brain $A\beta$ in the mice [12]. Cerebrospinal fluid (CSF) concentrations of bapineuzumab, a humanized monoclonal $A\beta$ antibody, are also found to be, on a molar basis, approximately three orders of magnitude less than typical CSF $A\beta$ concentrations [reviewed in 13].

The above considerations, of course, do not necessarily disallow direct CNS actions of $A\beta$ immunotherapeutics. Golde [14], for example, has cogently argued that if endogenous antibodies can have material effects on the CNS, which is clearly the case [15,16], then exogenous antibodies should be able to do so as well. On the other hand, considering that only minute quantities of peripherally-administered $A\beta$ antibodies reach the CNS, whereas they are wholly and directly exposed to circulating $A\beta$, it is difficult to understand how interactions of $A\beta$ antibodies with circulating $A\beta$ can be ignored as at least a potential, additional mechanism of action for $A\beta$ immunotherapy.

We have explored specific mechanisms by which $A\beta$ / $A\beta$ antibody immune complexes ($A\beta$ ICs) formed in blood in the course of $A\beta$ immunization might enhance clearance of $A\beta$ through enhanced interactions with the complement system. These studies were informed by the fact that major pathways for peripheral pathogen clearance in primates hinge on complement receptor 1 (CR1) [17], single nucleotide polymorphisms (SNPs) in which have been consistently identified as a significant risk factor for AD [18–22]. Compared to $A\beta$ alone, we found that the presence of $A\beta$ antibodies in the fluid phase dramatically increased virtually all steps in the major pathways for peripheral pathogen clearance in primates, including complement activation, formation of complement-opsonized complexes that are ligands for CR1, and peripheral capture and disposal of $A\beta$ through CR1-mediated erythrocyte and macrophage mechanisms. Consistent with these *in vitro* results, clearance of $A\beta$ from plasma and erythrocyte compartments *in vivo* was also robustly enhanced in non-human primates intravenously (IV) inoculated with $A\beta$ ICs. Although, as noted, these findings do not disallow CNS actions of $A\beta$ immunotherapy, they do strongly suggest that

peripheral effects should be considered as well—particularly since peripheral strategies might avoid the CNS adverse effects that have been encountered in previous AD immunotherapy trials [1,4,5].

2. Methods

2.01 Subjects

Human erythrocytes for the various experiments were obtained from study investigators under an Institutional Review Board-approved protocol. Two male Cynomolgus macaque monkeys (25 years old, 7.0 kg weight, and 28 years old, 6.5 kg weight) received intravenous injections of A β and blood samples were taken from them at various intervals (see below). These studies were performed under an Institutional Animal Care and Use Committee-approved protocol.

2.02 Preparation of A β

Lyophilized human synthetic A β (1–42) (Genscript, Piscataway, NJ) or FITC-conjugated A β (1–42) (Bachem, Torrance, CA) was resuspended in sterile 100% DMSO (Sigma, St. Louis, MO) at 10 mg/ml, diluted to 2 mg/ml in sterile ddH₂O, and then brought to 1 mg/ml in sterile 100 mM Tris, pH 7.4. The suspension was incubated overnight at room temperature (RT), in the dark, with shaking at 450 rpm. The resulting 1 mg/ml stock solution was then diluted with 100 mM Tris to achieve the concentrations employed in the experiments. Western blots of A β solutions prepared in this manner showed the presence of A β aggregates at multiple molecular weights, an important point since the monomeric form of A β poorly activates complement, if at all, whereas A β aggregates are relatively potent activators [23].

2.03 Serum complement activation

Various concentrations of A β prepared in 100 mM Tris, as above, were incubated with either A β antibody (4G8, Biolegend, San Diego, CA) or PBS, pH 7.2, after which the solutions were mixed with normal human serum (NHS) (CompTech, Tyler, Texas) for 30 minutes at 37°C. NHS plus 10 mM EDTA (Amresco, Solon, OH) (final concentration in the serum) was employed as a control. C3a production, one of several standard measures of complement activation, was assayed by ELISA (Affymetrix, Santa Clara, CA, #BHS2089) following the manufacturer's protocol.

2.04 iC3b Western blots and densitometry

NHS was incubated with 300 μ g/ml A β alone or A β ICs for 30 minutes at 37°C to permit complement activation, generation of complement opsonins, and their covalent binding to A β . To form A β ICs, a 9:1 molar ratio of A β :4G8 antibody was employed, as this ratio gave optimal complement activation (see Section 3.01, below). As a control to block complement activation and opsonization, 10 mM EDTA was added to NHS prior to incubation with A β or A β ICs. The solutions were run under reducing/denaturing conditions on SDS-PAGE 4–15% mini-PROTEAN TGX gels (BioRad, Hercules, CA, #146-1086), transferred to PVDF membranes (BioRad, #170-4156), blotted with a biotinylated (Thermo Fisher Scientific, Waltham, MA, #21326) iC3b antibody (Quidel, San Diego, CA), and imaged on an Odyssey

Imaging System (LI-COR, Lincoln, NE). To control for any effects of endogenous Igs or iC3b, a parallel gel was also run and analyzed under the same conditions, except that NHS was pre-depleted of endogenous Ig and iC3b using A β antibody 4G8 (Biolegend), iC3b antibody (Quidel), protein A/G PLUS agarose beads (Santa Cruz Biotechnology, Dallas), and standard immunoprecipitation methods. Uncompressed images of both Western blots were analyzed using ImageJ software (<https://imagej.nih.gov>). Blot background was quantified by integrating the signal in 10 different regions where there were no visible bands on the membrane, and linear regression was used to determine the relationship between integrated area and integrated background intensity ($R^2 > 0.99$ for both blots). For each band, the signal was integrated and the background (based on the area of the band) was subtracted to give final integrated optical densities.

2.05 Erythrocyte adhesion assay

To measure erythrocyte binding to A β alone and to A β ICs, a previously-published method for assaying erythrocyte CR1 binding to opsonized CR1 ligands was employed [24]. A β 42 was prepared as described above and 50 μ L was coated on a high-binding Costar 96-well plate (#07-200-39, Fisher Scientific, Waltham, MA) for 1 hour with shaking (450 rpm) at RT. The plate was washed 3X with 2/3 TBST (10 mM Tris, pH 7.2, 100 mM NaCl, 0.05% Tween-20), blocked with 100 μ L 0.5% PEG-3350 (Sigma, # P4338-1KG) in 2/3 TBST for 1 hour, RT, with shaking, and washed 3X again before 50 μ L of various concentrations of amyloid antibody 4G8 were added to each well so as to form A β /A β ICs, as in the previous experiment on complement activation. The plate was then incubated for 1 hour at RT with shaking, washed 3X, and incubated for 30 minutes at RT with 50 μ L of NHS, diluted 1:32 in VBS++ (CompTech, #B112), as a complement source for opsonization of the A β and A β ICs. To demonstrate that A β binding is mediated by complement opsonization, parallel wells were incubated with heat-inactivated (56°C, 30 minutes) NHS or C1q-depleted NHS (Comp Tech), both of which block various stages of complement activation but at different points. Plates were washed 2X with 2/3 PBST and 1X with adhesion buffer (140 mM Dextrose, 100 mM NaCl, 450 uM CaCl₂, 170 uM MgCl₂, 8 mM Tris, pH 7.4). Packed human erythrocytes were then diluted 1:2667 in adhesion buffer, and 200 μ L each of the suspensions were incubated with all plates for 45 minutes at RT. After washing, erythrocytes that remained bound to the plate were imaged (brightfield, 100X) using an inverted Olympus IX71 microscope (Olympus, Center Valley, PA) and quantified using investigator-independent ImageJ software.

2.06 Cell culture and differentiation

As in many other published studies to investigate various effects on macrophages *in vitro* [reviewed in 25], THP-1 cells (TIB-202, ATCC, Manassas, VA), a monocytic leukemia cell line, were cultured for fewer than 30 passages following ATCC recommendations, then differentiated to cells bearing immunologic, morphologic, and functional characteristics of macrophages [reviewed in 25]. All media used fetal bovine serum (FBS, ATCC) that was heat-inactivated for 30 minutes at 56°C in order to prevent complement activation. To differentiate THP-1 monocytes, cells were collected by centrifugation at 500 x g for 5 minutes at 23°C and resuspended in fresh media with 0.01 μ M phorbol 12-myristate 13-acetate (PMA, Sigma-Aldrich #P1585, St. Louis, MO) and 1 μ M retinoic acid (RA, Sigma

#R4643). After a brief vortex, 1 mL of the suspended cells was dispensed to each well of a tissue-culture-treated 24-well plate (Costar #3526, Corning, Corning, NY), followed by incubation for 24 hours at 5% CO₂ and 37°C, at which point the cells became adherent and morphologically distinct. The media was aspirated off and replaced with 1 mL media containing 1 µM RA. The plate was then further incubated at 5% CO₂ and 37°C for 78 hours, a time when our pilot studies showed cellular expression of typical macrophage markers (e.g., CD11b, CD14, HLA-DR, CD35, CD36) and maximal expression of CR1.

2.07 Aβ capture and uptake by THP-1-derived macrophages

THP-1-derived macrophages were incubated for 1.5 hours at 5% CO₂ and 37°C with 10% NHS in RPMI-1640 (#30-2001, ATCC) containing 1 µM FITC-conjugated Aβ alone or 1 µM FITC-conjugated Aβ plus 0.1 µM Aβ antibody 4G8 to permit formation of Aβ ICs. In order to control for specificity to complement reactions, parallel wells employed heat-inactivated or C1q-depleted NHS, and Fc receptor-mediated and non-specific antibody binding were blocked in all wells by replacing the culture media with 1 µM human IgG (Sigma, #I4506) in RPMI-1640 for 30 minutes prior to assay. After incubation of THP-1-derived macrophages with Aβ alone or Aβ ICs under the various experimental and control conditions, PBS was added to each well and adherent cells were disassociated from the plate by pipetting. Cells were transferred to 1.5 mL microcentrifuge tubes and spun for 5 minutes at 500 x g at 23°C, resuspended in 2% paraformaldehyde in PBS, incubated 20 min, spun again, resuspended in PBS, and assayed for Aβ using flow cytometry. Here, the cells were incubated with fluorophore-conjugated primary antibodies directed against five common macrophage antigens: PE-conjugated mouse monoclonal anti-CR1 (Biolegend, San Diego, CA, Clone E11 PE: #333406), PE/Cy5-conjugated rat monoclonal anti-CD11b (Biolegend, Clone M1/70 APC-Cy7: #101226), Pacific Blue-conjugated mouse monoclonal anti-CD14 (Biolegend, Clone M5E2 PacBlue: #301815), APC/Cy7-conjugated mouse monoclonal anti-HLA-DR (Biolegend, Clone L243 APC-Cy7: #307606), and APC-conjugated mouse monoclonal anti-CD36 (Biolegend, Clone 5-271 APC, #336208). Incubation was for 30 minutes with rapid shaking at 4°C, after which the cells were spun 3 minutes at 1000 x g, washed once with PBS, and resuspended in PBS. Compensation controls were run with IgG antibody beads (Becton-Dickinson #552843 or 552845, San Jose, CA) appropriate for antibody host species, and received the same treatment as the cells. To determine compensation for fluorescent Aβ, fluorescent Aβ and mouse anti-human Aβ antibody (6E10, Biolegend) were added to regular mouse compensation beads. Cells were run on a Becton-Dickinson LSR II flow cytometer and analyzed with FlowJo v10.1 (Ashland, OR). Only cells that gated positive for all five of the macrophage markers were included in the final analysis. Using these stringent criteria, median fluorescence of FITC-conjugated Aβ/cell was used as a measure of Aβ capture by THP-1-derived macrophages in each condition.

2.08 Histology

To complement the quantitative assessment by flow cytometry, uptake of Aβ and Aβ ICs in THP-1-derived macrophages was qualitatively assessed using fluorescence and confocal microscopy. Macrophages were differentiated as described above, but were grown on glass coverslips instead of culture plates. After exposure to 1 µM fluorescent Aβ or Aβ/4G8 ICs (1 µM Aβ plus 0.1 µM 4G8), as in the flow cytometry studies, the cells were stained with a

cell membrane marker (Sigma, # PKH26) and fixed with 2% paraformaldehyde (Electron Microscopy Services, Hatfield PA, #15714-5) in PBS. The glass coverslips were inverted and mounted on glass slides with DAPI-containing mounting medium (ProLong Gold with DAPI (Thermo Fisher). Images were collected on a Nikon A1 confocal microscope.

2.09 Clearance of IV-administered A β and A β ICs in non-human primates

To demonstrate functional relevance of the above mechanisms, two Cynomolgus macaque monkeys, ages 28 and 25 years old, received saphenous vein injections of either 3.0 mg A β 42 alone or 3.0 mg A β 42 that had previously been incubated with 11.1 mg A β antibody 4G8 to form A β ICs. Although the latter amounts to only a 9:1 molar ratio of A β to antibody, our earlier complement activation assays (see Section 3.01, below) indicated near-asymptotic responses to this ratio, and higher titers of antibody would have been prohibitively expensive (a request to obtain relevant doses of bapineuzumab from the license holder was denied). Femoral artery blood samples were taken in EDTA tubes 5 minutes before and 1, 5, 10, 15, 20, and 40 minutes after IV infusion of A β or A β ICs. Plasma and erythrocyte samples were processed as previously described [26], and were assayed using Wako A β 42 ELISA kits (#298-62401, Richmond, VA) following the manufacturer's recommendations throughout. To control for baseline differences in plasma and erythrocyte A β levels in the animals, the data were normalized as percent of A β recovered in the erythrocyte compartment relative to the amount available in the plasma compartment.

2.10 Statistics

Parametric statistical tests (ANOVA, repeated measures ANOVA, Pearson's R) were employed throughout. Significance levels were two-tailed.

3. Results

3.01 A β activation of complement is enhanced by formation of A β ICs

Replicating our previous work [23,26,27] and that of others [28,29], A β alone was found to activate complement in a dose-dependent fashion ($F = 203.9$, $P < 0.001$). Prior incubation of A β with A β antibody 4G8 to form A β /4G8 ICs significantly enhanced activation by as much as 34-fold compared to A β alone ($F_{\text{overall}} = 110.6$, $P < 0.001$) (Fig. 1A). Complement activation by A β ICs was significantly dose-dependent both when 4G8 concentration was equimolar to increasing concentrations of A β ($F = 95.6$, $P < 0.001$) (Fig. 1A) or when A β concentration was held constant and 4G8 concentrations were varied ($F = 76.4$, $P < 0.001$) (Fig. 1B). Notably, maximum enhancement of complement activation by A β ICs occurred at an A β :4G8 molar ratio of approximately 9:1 (Fig. 1B). This is consistent with the fact that A β aggregates are required to activate complement [23]; steric hindrance and epitope masking in A β aggregates would therefore make it unlikely that all A β moieties in an A β aggregate could simultaneously be bound by an A β Ig.

Spontaneous aggregation of antibodies *in vitro* is known to provide the multiple binding sites necessary to activate C1 [c.f., 23]. Thus, A β antibody 4G8 alone also dose-dependently activated complement ($F = 108.2$, $P < 0.001$), though, by comparison, the activation was again far less than that for A β ICs ($F = 181.7$, $P < 0.001$) (Fig. 1A). EDTA, a standard

control for specificity to complement reactions, completely abolished the effects in all conditions.

3.02 Opsonization of A β is enhanced by formation of A β ICs

Following complement activation, complement opsonins such as iC3b bind back to the activating substrates. Because this binding is covalent, the opsonins remain attached to the activator even under the reducing/denaturing conditions of conventional SDS/PAGE Western blots. When A β alone or A β /4G8 ICs were incubated with NHS to permit complement activation, A β IC solutions exhibited markedly enriched iC3b immunoreactivity on iC3b Western blots compared to A β alone (Fig. 2A), consistent with the enhanced complement activation provided by A β ICs. As expected, immunoreactivity for iC3b (observed as iC3b fragments under reducing/denaturing conditions) was evident across multiple molecular weights, since multiple aggregate species of A β can activate and bind complement opsonins [23]. EDTA abolished iC3b immunoreactivity in both A β IC and A β samples (Fig. 2A), demonstrating that the iC3b was generated by complement reactions specific to A β ICs or A β . Pre-depletion of endogenous Igs and iC3b by immunoprecipitation in a second blot had no detectable effect on these results, and the depleted blot could not be distinguished visually or by densitometry ($F = 1.12$, $P = 0.307$) from the undepleted blot, showing that the opsonization observed is the result of *de novo* complement activation and A β IC formation in the experiments. Densitometry of the Ig/iC3b-depleted and undepleted blots gave a nearly identical pattern of results, with immunoreactivity for parallel bands in the A β IC condition exhibiting from 3-fold to 5-fold increases in iC3b compared to A β alone ($F = 20.51$, $P = 0.001$) (Fig. 2B). The addition of EDTA to A β and A β IC solutions reduced densitometry values to background or less.

3.03 Erythrocyte capture of A β is enhanced by formation of A β ICs

Once opsonized by complement, the opsonized complexes can bind to CR1 expressed on the surface of primate erythrocytes. They are then ferried to the liver and stripped off by Kupffer cells for degradation [30,31]. To assay erythrocyte binding, A β or A β ICs in various concentrations were coated on 96-well plates, incubated with serum, then incubated with erythrocytes. Consistent with the ability of A β ICs to enhance complement activation and opsonization, as shown above (Figs. 1, 2), binding of erythrocytes to A β ICs was dramatically enhanced compared to A β alone ($F = 93.7$, $P < 0.001$), with increases of more than 10-fold at many A β IC concentrations (Fig. 3). Specificity to complement opsonization, as opposed to non-specific interactions with Igs or capture by non-complement-mediated mechanisms, was demonstrated by use of heat-inactivated NHS and C1q-depleted NHS, both of which abolish complement opsonization and/or classical pathway complement activation and both of which reduced adhesion to background levels (Fig. 3).

3.04 A β capture and uptake by THP-1-derived macrophages is enhanced by formation of A β ICs

THP-1-derived macrophages were characterized by flow cytometry for expression of typical macrophage markers such as CD14, HLA-DR, CD11b, CD36, and CR1 (Fig. 4A). The cells were then exposed to fluorescence-labeled A β alone or fluorescence-labeled A β ICs in the presence of NHS, heat-inactivated NHS, or C1q-depleted NHS, followed by flow cytometry

assay for capture of A β . By gating for expression of CD14, HLA-DR, CD11b, CD36, and CR1 (Fig. 4B), only cells that expressed appropriate macrophage markers were evaluated for A β . When incubated with NHS to permit complement opsonization, detection of cell-associated A β fluorescence was increased by nearly 4-fold in the A β IC condition compared to A β alone (Figs. 4C,4D). These results appeared to be mediated almost wholly by macrophage binding of complement-opsonized A β and A β ICs, as opposed to non-complement-dependent mechanisms, since THP-1-derived macrophages were pretreated with IgG to block Fc receptor-mediated and non-specific antibody binding. Moreover, A β uptake was reduced to background levels when heat-inactivated NHS and C1q-depleted NHS were employed as complement sources (Fig. 4D). Heat-inactivated NHS blocks all complement pathways, whereas C1q-depleted NHS blocks the classical pathway but should permit alternative pathway activation. These data for macrophages and the above data for erythrocytes therefore suggest that classical pathway activation is a key mechanism in THP-1-derived macrophage and erythrocyte binding of A β and A β ICs.

Enhanced uptake of A β ICs by THP-1-derived macrophages could also be demonstrated qualitatively using fluorescence microscopy (Figs. 5A–5C). Here, THP-1-derived macrophages were incubated with FITC-conjugated A β /4G8 ICs (1 μ M A β plus 0.1 μ M 4G8) (Fig. 5A) or FITC-conjugated A β alone (1 μ M) (Fig. 5B) in the presence of NHS to permit complement activation. A β fluorescence was clearly more intense in A β /4G8 IC-treated macrophages, and was evident in many more cells compared to A β alone. Heat inactivation of serum, which abolishes complement activation, reduced A β detection to background (Fig. 5C). Evaluation by confocal microscopy (Fig. 5D) strongly suggested that A β and A β ICs were not only bound by THP-1-derived macrophages, but also were phagocytosed by the cells.

3.05 Functional effects of A β ICs on peripheral clearance of A β in non-human primates

In order to study the effects of A β IC clearance *in vivo*, two Cynomolgus macaque monkeys received saphenous vein injections of either A β alone or A β that had previously been incubated with A β antibody 4G8 to form A β ICs. In both the A β IC and A β alone conditions, levels of IV-injected A β 42 in the plasma and erythrocyte compartments rose steeply within 1 minute of infusion, the earliest time point that was possible to sample, and declined to near baseline over the next 15 minutes, similar to the kinetics we [26] and others [32] observed in a previous study with A β 40. Compared to A β alone, erythrocyte capture of A β ICs from the plasma compartment was significantly increased overall (by repeated measures ANOVA, $F = 1088.5$, $P < 0.001$) and by more than 6-fold at the earliest time points (Fig. 6).

4. Discussion

The present experiments follow up and are completely consistent with our prior findings [26] that blood A β is subject to peripheral clearance by a major pathway for circulating pathogens in primates, immune adherence [30,31]. CR1-mediated erythrocyte immune adherence, as well as CR1-mediated macrophage capture of pathogens, requires complement activation and opsonization. Although some pathogens, including A β [26–29], can, by

themselves, stimulate complement activation and opsonization through binding of the pathogen to C1q, these processes are markedly enhanced when the pathogens form immune complexes, which bind with higher affinity to C1q. Thus, in the present experiments, the formation of A β ICs, as would occur with active and passive A β immunotherapy, increased A β activation of fluid phase complement by as much as 34-fold. This enhancement was predictably followed by multi-fold increases in complement opsonization, erythrocyte adherence, and macrophage phagocytosis. *In vivo*, capture of A β through the erythrocyte clearance pathway was also robustly increased for A β ICs compared to A β .

Whether generated in the blood by active immunization or infused into the blood by passive immunization, A β antibodies are, first and foremost, directly exposed to circulating A β , permitting the formation of A β ICs. Theoretically, there is, in fact, enough A β in the plasma compartment to form ICs with all the bapineuzumab infused into the average human being after a typical 1 mg/kg dose (assuming an average body weight and blood volume of 80.7 kg and 4.7 liters, respectively, and an average plasma A β level of 500 pg/ml [33–36]). Although, in practice, antibodies do not bind antigens with 100% efficiency (the K_D for bapineuzumab is reported to be 89 nM for soluble A β 40 [37]), the amount of A β antibody available to enter the brain should still be substantially limited by formation of A β ICs in the blood and their subsequent clearance by the mechanisms studied in this report. Antibodies surviving these processes would then also have to penetrate the blood-brain barrier. It may not be surprising, therefore, that the concentrations of A β antibodies that reach the brain are estimated to be <0.1% of the amount administered peripherally, and many orders of magnitude less than the molar amount of A β present in the AD brain [12–14].

Data on plasma A β levels after A β immunotherapy [reviewed in 38] and after experimental infusion of A β antibodies in AD transgenic mice [9,11] have been reported in some cases. Although these studies typically find increased plasma A β after treatment, they do not necessarily contradict the present results, and, in fact, may support them. Efflux of CNS A β into the plasma compartment after treatment appears to be rapid and extensive, and, after treatment, virtually all of the A β in plasma appears to be in the form of A β ICs [9]. However, if these high levels of A β ICs were simply “sequestered” in the circulation, as presently assumed, they would almost certainly lead to immune complex disorders, where high levels of uncleared immune complexes lodge in various organs and cause significant pathology [reviewed in 39]. Subsequent infusions of A β antibody, as in AD immunotherapy, would further compound this problem. Thus, the data on increased plasma A β after treatment with A β antibody antibodies actually demands an active peripheral clearance mechanism such as immune adherence in order to avoid immune complex disease in the recipients. The A β ICs cannot simply persist and build up. A more accurate representation would therefore be that the enhanced efflux of CNS A β into the plasma seen after A β infusion is interactive with the enhanced clearance of peripheral A β shown in the present experiments. AD immunotherapy studies on plasma alone do not (and cannot) measure this latter contribution, whereas our experiments, using infusions of A β ICs (as opposed to only the antibodies themselves) do so.

Although it is unquestionable that AD is a CNS disease, there is ample precedent for peripheral influences on brain disorders (e.g., effects of diabetes on the CNS) [reviewed in

40]. Whether considered in light of the “peripheral sink hypothesis” [9–11] or some other mechanism [32], impaired peripheral clearance of A β , as shown for AD subjects in our previous research [26], could not be favorable for brain concentrations of A β , whereas enhanced peripheral clearance by A β antibodies, as shown by the present data, should be beneficial. Accordingly, whether or not the A β antibodies provided by A β immunotherapy actually penetrate to the brain in sufficient amounts to assist removal of brain A β , the peripheral effects of such antibodies should not be ignored. Likewise, whether or not normal erythrocyte and macrophage peripheral clearance pathways are sufficient to deal with the increased plasma A β brought about by AD immunotherapy [38], approaches to enhance peripheral, complement-mediated removal of circulating pathogens have been reported and might be considered for A β . For example, a bispecific antibody that bound both the pathogen and the erythrocyte receptor for complement-opsonized complexes, CR1, has been shown to substantially enhance clearance of the pathogen [41].

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RESEARCH IN CONTEXT

Systematic Review

Although amyloid β peptide ($A\beta$) immunotherapy for Alzheimer's disease (AD) continues to be pursued, critical questions still remain with respect to its mechanism and sites of action. This is particularly true given the incomplete penetration of peripherally-administered $A\beta$ antibodies into the CNS compared to the full penetration of the antibodies in the peripheral circulation. **Interpretation:** Our findings demonstrate a heretofore unexplored mechanism wherein $A\beta$ immune complexes (ICs) dramatically enhance virtually all steps in complement-mediated peripheral clearance of $A\beta$ compared to $A\beta$ alone. These enhanced effects of $A\beta$ ICs apply to erythrocyte clearance of $A\beta$ (immune adherence), as well as to macrophage clearance of $A\beta$. They therefore warrant consideration as additional mechanisms by which $A\beta$ immunotherapy may act. **Future Directions:** Complement receptor 1 (CR1) plays a pivotal role in erythrocyte and macrophage clearance mechanisms, and polymorphisms in CR1 are a known risk factor for AD. Demonstrating that these polymorphisms predictably alter peripheral $A\beta$ clearance, a task we are pursuing, would further indicate the potential importance of peripheral mechanisms in $A\beta$ pathogenesis and therapeutics.

Fig. 1. Complement activation by Aβ and Aβ/4G8 ICs

A) Aβ₄₂ was incubated with equimolar Aβ antibody 4G8 to form Aβ/4G8 ICs, then reacted with NHS as a complement source. Significant dose-dependent complement activation ($P < 0.001$), as measured by C3a generated, was observed for Aβ ICs even at low nM concentrations. Aβ alone and 4G8 alone also dose-dependently activated complement ($P < 0.001$), but required μM concentrations to stimulate C3a production above background levels. EDTA abolished these effects, showing that they are specific to complement mechanisms. **B)** Complement activation by Aβ ICs could also be shown to be dose-dependent with respect to the amount of Aβ antibody available ($P < 0.001$). Here, a constant amount of Aβ (2215 nM) was incubated with varying concentrations of Aβ antibody 4G8, from low to equimolar concentrations relative to Aβ. Note that the X-axis is log-scaled in these graphs to include the wide range of concentrations. For both figures, each data point represents the mean of triplicate samples. Error bars denote standard error of the mean. Standard error for replicates is not shown when they are smaller than the symbols for a particular concentration.

Fig. 2. Complement opsonization of Aβ and Aβ/4G8 ICs

A) After exposure to NHS to permit complement activation, Aβ alone and Aβ IC solutions were run on SDS reducing/denaturing Western blots to detect the complement opsonin iC3b. EDTA treatment (lanes 2 and 4 of the iC3b blot) abolished complement activation. The absence of iC3b immunoreactivity after EDTA treatment therefore shows that the iC3b detected in Aβ alone and Aβ IC conditions (lanes 1 and 3 of the iC3b blot) was specifically generated by Aβ and Aβ IC complement activation, and not by endogenous levels of iC3b. Compared to Aβ alone, iC3b immunoreactivity was markedly enhanced by Aβ ICs, consistent with enhanced complement activation by Aβ ICs in our earlier experiments. A parallel Western blot for Aβ (6E10 detection antibody) using the Aβ solution employed in the experiment is shown in the left-most lane. Note that bands for iC3b and Aβ aggregate forms typically coincide, as would be predicted given that iC3b is covalently bound to Aβ aggregate forms that can activate complement. **B)** Densitometry of the blot in panel A, as well as a second blot, prepared in the same way but pre-depleted of endogenous Igs and iC3b, gave a similar pattern of results that did not differ significantly from blot to blot ($P = 0.307$), with both showing highly-enriched iC3b immunoreactivity in Aβ IC compared to Aβ alone samples ($F = 71.65$, $P = 0.001$). For example, integrated optical density of the molecular weight bands designated by the four circled numbers on panel A were increased by 3-fold to 5-fold in the Aβ IC condition.

Fig. 3. Erythrocyte capture of A β and A β ICs

In erythrocyte adhesion assays, erythrocytes bound to A β -coated and A β /4G8 IC-coated plates in a dose-dependent manner ($P < 0.001$) and binding was significantly greater for A β ICs ($P < 0.001$). Abolition of effects by heat-inactivated NHS at the highest dose of A β ICs and A β alone (HINHS and $\circ$ symbol at bottom right of graph) demonstrated specificity to complement reactions as opposed to erythrocyte binding by mechanisms other than complement. Abolition of effects by C1q-depleted serum (-C1q and Δ symbol at bottom right of graph) confirmed this finding and, in addition, strongly suggested that erythrocyte capture of A β is primarily mediated by the classical, rather than the alternative complement pathway. Each data point represents the mean of duplicates. The vast majority of data points exhibited standard errors that were too small to show clearly. For the A β alone condition, the average standard error of the mean was 111.3 bound erythrocytes/field (range of 7.8 to 422.5). For the A β IC condition, the average standard error of the mean was 321.0 bound erythrocytes/field (range of 9.9 to 996.7).

Fig. 4. Macrophage capture of Aβ and Aβ ICs

A) Flow cytometry characterization of THP-1 monocytes and their differentiation to THP-1-derived macrophages revealed a transition to increased expression of inflammatory markers such as CD36, a mediator of macrophage phagocytosis [25], and, especially, CR1, the macrophage receptor for the major complement opsonins in primates [34,35]. **B)** To insure assessment of the macrophage-specific phenotype in subsequent experiments, all cells were gated for criterion expression of five different macrophage markers, CD14, HLA-DR, CD11B, CD36, and CR1. **C)** Fluorescence intensity distributions for samples pre-incubated with NHS. Distributions designated NHS and 4G8 represent fluorescence of the macrophage markers on which the cells were gated. Fluorescence distributions for gated, FITC-labeled Aβ and Aβ IC samples appear in the right-hand portion of the graph. A total of 100,000 cells were evaluated for each sample. **D)** Median fluorescence intensity of Aβ captured by THP-1-derived macrophages increased nearly 4-fold for Aβ ICs compared to Aβ alone. Only background readings were obtained for the cells when heat-inactivated NHS or C1q-depleted serum were substituted for NHS as a complement source.

Fig. 5. Macrophage uptake of A β and A β ICs

A–C) Representative fluorescence micrographs (10X) of THP-1-derived macrophages (DAPI nuclear counterstain) (blue) after exposure to 1 μ M FITC-conjugated A β plus 0.1 μ M A β antibody 4G8 to form A β ICs (**A**) or 1 μ M FITC-conjugated A β alone (**B**). A β IC and A β alone solutions were previously incubated with NHS to permit complement activation. The number of cells co-localized with A β (green), as well as their fluorescence intensity, was clearly enhanced by A β ICs. Heat inactivation of the serum reduced staining to background (**C**). **D)** Representative high power (40X objective) confocal micrograph of a macrophage after exposure to FITC-conjugated A β plus 4G8 and NHS. PKH26 (red) was used to stain the cell membrane and DAPI (blue) was used to stain the nucleus. Localization of the FITC label for A β (green) was primarily intracellular, suggesting that the macrophages had phagocytosed the A β ICs.

Fig. 6. Erythrocyte uptake and clearance of A β and A β ICs in non-human primates

Relative to the amount of A β available in the plasma compartment, erythrocyte uptake and clearance of A β inoculated into two non-human primates was significantly greater for A β ICs compared to A β alone ($P < 0.001$). Because erythrocyte and plasma levels at baseline differed in the animals, the data in the figure have been normalized as percent of A β recovered in the erythrocyte compartment relative to the amount available in the plasma compartment. Clearance into the erythrocyte compartment was rapid over the first 15 minutes after A β or A β IC injection, a characteristic of immune adherence [41]. By 20 minutes, A β IC values had returned to baseline, but remained elevated by some 20% relative to baseline for A β alone. Samples at each time point in each animal were assayed in duplicate. Standard error bars for each time point for each animal were smaller than the symbols at each time point and are not shown. For A β alone, mean standard error was 0.13% (range of 0.01 to 0.39). For A β IC, mean standard error was 0.82% (range of 0.10 to 2.81).