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Intracranial injection of Gammagard, a human IVIg, modulates the inflammatory response of the brain and lowers Aβ in APP/ PS1 mice along a different time-course than anti-Aβ antibodies

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Abstract

Gammagard[™] IVIg is a therapeutic approach to treat Alzheimer's disease currently in phase 3 clinical trials. Despite the reported efficacy of the approach the mechanism of action is poorly understood. We have previously shown that intracranial injection of anti-A antibodies into the frontal cortex and hippocampus reveals important information regarding the time-course of events once the agent is in the brain. In the current study we compared IVIg, mouse pooled IgG and the anti-A antibody 6E10 injected intracranially into the frontal cortex and hippocampus of 7 month old APP/PS1 mice. We established a time-course of events ranging from 1 day to 21 days postinjection.

IVIg and pooled mouse IgG both significantly reduced A deposition to the same degree as the 6E10 anti-A antibody, however, the clearance was much slower to occur, happening between the 3 day and 7 day time-points. In contrast, as we have previously shown, A reductions were apparent with the 6E10 anti-A group at the 1-day time-point. Also, neuroinflammatory profiles were significantly altered by the antibody treatments. APP/PS1 transgenic mice at 7 months of age typically exhibit an M2a inflammatory phenotype. All antibody treatments stimulated an M2b response, yet anti-A antibody was a more rapid change.

Because the neuroinflammatory switch occurs prior to the detectable reductions in amyloid deposition, we hypothesize that the IVIg and pooled mouse IgG act as immune modulators and this immune modulation is responsible for the reductions in amyloid pathology.

Introduction

Alzheimer's disease (AD) is characterized by deposition of amyloid plaques, composed of aggregated amyloid-beta (A) peptide, and neurofibrillary tangles, composed of aggregated tau protein (Hyman et al., 2012). Anti-A immunotherapy has been shown to improve cognition while lowering brain A in numerous mouse models (Morgan et al., 2000; Wilcock et al., 2004c; Maier et al., 2006). It has been shown that anti-A immunotherapy lowers brain A through both central and peripheral mechanisms (Wilcock and Colton, 2009). We previously showed that anti-A antibodies activate microglia, and the microglial activation is important for the clearance of compact amyloid deposits (Wilcock et al., 2003; Wilcock et al., 2004a). Antibodies can also catalytically disaggregate amyloid deposits in vitro (Solomon et al., 1997). When administered systemically, anti-A antibodies can enter

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the CNS and activate microglia (Wilcock et al., 2004c), but they also result in the efflux of A from the brain via a mechanism called the peripheral sink (DeMattos et al., 2001).

Adverse cerebrovascular events continue to plague the anti-A immunotherapy field. These adverse events include microhemorrhages and vasogenic edema (termed ARIA-H and ARIA-E, respectively) (Wilcock et al., 2004b; Sperling et al., 2011). We have previously published data that suggests activation of matrix metalloproteinases (MMPs) MMP2 and MMP9 are associated with the increase in microhemorrhage by anti-A immunotherapy (Wilcock et al., 2011a).

Intravenous immunoglobulin (IVIg) is a term applied to the pharmaceutical preparation of pooled human IgG derived from plasma. IVIg is used to treat immune deficiency disorders including idiopathic thrombocytopenic purpura and hypogammaglobulinaemia (Hartung et al., 2009). The described immunomodulatory effects of IVIg include downregulation of antibody production, inhibition of B-cell mediated IL-6 production and inhibition of nitric oxide secretion, among many other immunological effects (reviewed in (Dodel et al., 2010). After early reports of IVIg being tested in AD patients (Dodel et al., 2004), an open label dosing of Gammagard, a commercially available IVIg, reported good tolerance and increased plasma A following infusion. MMSE scores also increased in the small number of patients receiving the IVIg (Relkin et al., 2009). Phase 3 trials are ongoing after passing the futility analysis in January 2012. Despite the early indicators of a positive clinical benefit of IVIg in AD patients, the underlying mechanism(s) of action are unknown. Some have suggested the presence of anti-A antibodies accounts for the effects of IVIg (Magga et al., 2010), while others suggest immune modulation effects (Dodel et al., 2010).

We have previously shown that anti-A antibody administration systemically results in significant changes in inflammatory markers in the brain (Wilcock et al., 2011b). We are using the macrophage phenotypes of M1, M2a, M2b and M2c to characterize the neuroinflammatory phenotype of the brain. The use of the multiple markers of neuroinflammation allows us to better characterize the inflammatory state of the brain beyond examining typical microglial cell surface markers of "activation". In the current study we use a time-course to examine the effects of IVIg and mouse IgG on neuroinflammation and A load, and contrast these effects with anti-A antibody.

Materials and Methods

Animals

APP/PS1 transgenic mice (C57BL6 mice carrying human APPSwe and PS1-dE9 mutations; Jankowsky et al., 2004) were bred in house and aged to 7 months. 120 seven month old APP/PS1 mice were assigned to one of four injection groups; saline, anti-A antibody $(6E10, A_{3-8}, IgG1, Covance, Princeton, NJ), IVIg (trade name Gammagard, Baxter,$ Deerfield, IL composed primarily of IgG1 and IgG2 (Schwab and Nimmerjahn, 2013)) or mouse IgG (Sigma-Aldrich, St Louis, MO, composed primarily of IgG1 and IgG2a (Sant'Anna et al., 1985)). The anti-A antibody, IVIg and mouse IgG were diluted to a final concentration of 1mg/ml in sterile saline. Six mice in each group (3 female and 3 male) were euthanized at each time-point (1d, 3d, 7d, 14d and 21d). The study was approved by the University of Kentucky Institutional Animal Care and Use Committee and conformed to the National Institutes of Health Guide for the Care and Use of Animals in Research.

Bilateral intracranial injection

On the day of surgery mice were weighed, anesthetized with isoflurane, and placed in a stereotaxic apparatus (51733D digital dual manipulator mouse stereotaxic frame, Stoelting Co. Wood Dale, IL). A midsaggital incision was made to expose the cranium and four burr

holes were drilled with a dental drill mounted in the stereotaxic frame over the frontal cortex and hippocampus to the following coordinates: frontal cortex, anteroposterior, +1.7mm, lateral +/- 2.0mm; hippocampus, anteroposterior -2.7mm; lateral, +/- 2.5mm, all taken from bregma. A 26 gauge needle attached to a 10ml Hamilton syringe (Hamilton, Reno, NV) containing the solution to be injected was lowered 3.0mm ventral to bregma, and a 2μl injection was made over a 2 min period. The incision was cleaned and closed with surgical staples for the 1-7 day time points, and sutures for the 14 and 21-day time points. Sutures were removed at 10 days.

Tissue processing and histology

After injection with a lethal dose of pentobarbital, mice were perfused intracardially with 25ml normal saline. Brains were rapidly removed and bisected in the mid-sagittal plane. The left half was immersion fixed in freshly prepared 4% paraformaldehyde. The right half was dissected with the frontal cortex and hippocampus being isolated, flash frozen in liquid nitrogen and stored at -80°C. The left hemibrain was passed through a series of 10, 20 and 30% sucrose solutions as cryoprotection and 25μm frozen horizontal sections were collected using a sliding microtome and stored floating in PBS containing sodium azide at 4°C.

Section spaced 300μm spanning the estimated injection site were initially mounted and stained by Cresyl Violet to identify the injection site. For all subsequent histology and immunohistochemistry six sections spanning the injection site, spaced approximately 100μm apart were selected and analyzed. Free floating immunohistochemistry for CD45 (Rat monoclonal, Thermo Scientific, Rockford, IL) and A (Rabbit polyclonal A 1-16, Life Technologies, Carlsbad, CA) were performed as described previously (Wilcock et al., 2003).

ELISA measurement

Protein was extracted for A analysis from the right frontal cortex using a two-step extraction method. First, the brain was homogenized in PBS containing a complete protease and phosphatase inhibitor (Pierce Biotechnology Inc. Rockford, IL), These samples were centrifuged at 16,000 \times g at 4 \degree C for 1 hour. The supernatant was removed and became the "soluble" extract. The resulting pellet was homogenized in 100μl 70% formic acid and centrifuged at 16,000×g at 4°C for 1 hour. The supernatant was removed and neutralized 1:20 with 1M Tris-HCl and became the "insoluble" extract. Protein concentration for both the soluble and insoluble extracts was determined using the bicinchonic acid (BCA) protein assay according to manufacturer's instructions (Thermo Scientific, Rockford IL). We used the Meso-Scale Discovery multiplex ELISA system to measure A 38, A 40 and A 42 (MSD, Gaithersburg MD). ELISA kits were run according to the manufacturer's instructions. We took the soluble extract remaining after A analysis and measured levels of MMP9, tissue inhibitor of metalloproteinases 1 (TIMP1), MMP2 and TIMP2 by ELISA according to manufacturer instructions (R&D Systems, Minneapolis, MN). We also measured the levels of mouse and human IgG to detect the injected antibodies. We used the Easy-Titer IgG assay kits (Thermo scientific, Rockford, IL) for mouse IgG and human IgG.

Quantitative real-time RT-PCR

RNA was extracted from left hippocampus using the Trizol Plus RNA Purification System (Life Technologies, Grand Island NY) according to the manufacturer's instructions. RNA was quantified using the BioSpec Nano spectrophotometer (Shimadzu North America, Columbia, MD) and cDNA was reverse transcribed using the cDNA High Capacity kit (Applied Biosystems, Foster City CA) according to the manufacturer's instructions. Realtime PCR was performed using the 384-well microfluidic card custom Taqman assays containing TaqMan Gene Expression probes for our genes of interest (Life Technologies,

Carlsbad, CA). All gene expression data were normalized to 18S rRNA expression. Foldchange compared was determined using the \wedge Ct) method.

Analysis

Data are presented as mean \pm standard error of mean (SEM). Statistical analysis was performed using the JMP statistical analysis program (SAS, Cary NC). Statistical significance was assigned where the P-value was lower than 0.05. One-way ANOVA and two-way ANOVA were used, where appropriate, to detect treatment differences and differences within treatment groups along the time-course.

Results

Measurement of mouse IgG in the protein extracted from the right frontal cortex showed that mice injected with mouse IgG and anti-A IgG had significantly increased levels of IgG in the brain 1 day after injection (Figure 1A). This was reduced dramatically by the threeday time point and was down to the levels of mice receiving saline or human IVIg by the 7 day time point. At no time point did we observe a difference between the mouse IgG and anti-A antibody injected mice for the amount of IgG present suggesting the clearance kinetics of the IgG was identical (Figure 1A). Similarly, quantification of human IgG showed significant increases only in the mice receiving the human IVIg at the 1 day time point. This was reduced but still significantly elevated at 3 days and was undetectable at the 7 day time point (Figure 1B).

Total A load measured by immunhistochemistry in the frontal cortex and hippocampus of the saline injected mice was as previously shown for this mouse model (Figure 2A-C and (Jankowsky et al., 2004)). We found that at the earliest time point, 1 day post injection, the anti-A antibody had significantly reduced the brain A load in both frontal cortex and hippocampus (hippocampus shown Figure 2D, quantification Figure 2M). There was a slight further reduction in A with anti-A antibody between 1-3 days (Figure 1D, E and M) and then no further reductions were observed. In fact, A levels began to steadily increase from the 7-21 day time-points (Figure 2M). In contrast, both reductions in A load were not observed with IVIg or mouse IgG until the 7 day time-point, indicating that this reduction occurred between the 3 and 7 day time-points (Figure 2H-I, K-L and M). At no time-point did we observe a significant difference between the IVIg and mouse IgG treatment groups.

Measurement of A levels by ELISA showed similar patterns as the immunohistochemistry. The soluble pools of A 1-40 and A 1-42 were significantly reduced by anti-A antibody at 1 day compared to saline, and a further slight reduction was observed at the 3 day time point. Interestingly, soluble A 1-40 increased dramatically between the 3 and 21 day time point, reaching the levels observed in mice receiving saline injection (Figure 3A). While a similar pattern was observed with soluble A 1-42, the increase between 3 and 21 days was less dramatic (Figure 3B). Both mouse IgG and IVIg injection reduced soluble A 1-40 and A 1-42 between the 3 and 7 day time-points and showed a gradual increase between the 7 and 21 day time points, although this increase was less dramatic for the A 1-40 than the anti-A antibody group (Figure 3A and B). In contrast to the soluble pools of A , insoluble A 1-40 was reduced between the 1 and 3 day time-points by the anti-A antibody, while insoluble A 1-42 was already reduced at the 1 day time point. Mouse IgG and IVIg both showed significant reductions in insoluble A 1-40 and A 1-42 between the 3 and 7 day time points (Figure 3C and D). Of note, is the apparent increased efficacy of IVIg to lower insoluble A 1-40 compared to the reduction seen by anti-A antibody or mouse igG (Figure 3C).

To determine the role of neuroinflammation we first examined CD45 immunohistochemistry. Saline injected mice showed low levels of microglial activation around plaques in the frontal cortex (images not shown) and hippocampus (Figure 4A-C). One day following injection a small number of CD45 positive cells could be seen associated with the injection site (Figure 4A) but these were not detectable at the 3 day time point and beyond (Figure 4B and C). In contrast, diffuse microglial activation could be seen throughout the dentate gyrus, CA3 and CA2 regions of the hippocampus 1 day following injection (Figure 4D). This activation then localized to the granule cell layer of the dentate gyrus and the pyramidal cell layer of the CA3 and CA2 regions (Figure 4E). We have previously shown this similar pattern of microglial activation following anti-A antibody injection along this same time-course (Wilcock et al., 2003). Some microglial activation in these cell layers remained at 7 days and the activation resolved close to saline injected levels by 21 days (Figure 4F, M and N). While IVIg and mouse IgG showed some increased CD45

expression at 1 and 3 days, it was not until 7 days after injection that we observed significant microglial activation that resembled the anti-A antibody injected mice (Figure 4G-L). Quantification of the CD45 immunohistochemistry showed that microglial activation peaked 3 days following injection with anti-A antibody, but 7 days after injection with IVIg and mouse IgG in both the frontal cortex and hippocampus (Figure 4M and N).

To better characterize the neuroinflammatory response to the injected antibodies, we isolated RNA from the right hippocampus and performed real-time RT-PCR for genes relatively specific for the four phenotypes of macrophages; M1, M2a, M2b and M2c. We found that all M1 associated genes (IL-1 , Marco and TNFa) were significantly increased at 3 days following anti-A injection and quickly resolved by the 7 day time-point (Figure 5A). Mouse IgG and IVIg showed significant increases at the 7 day time-point but this increase was slower to resolve (Figure 5A). In contrast to the M1 markers, we found that M2a associated genes (Arg1, IL-1Ra and YM1) were significantly decreased at the 3 day timepoint with anti-A antibody and at the 7 day time-point with mouse IgG and IVIg (Figure 5B). Again, the M2a markers were slower to reach control levels with mouse IgG and IVIg than the anti-A antibody group, much like the M1 markers (Figure 5B). Of particular interest is the M2b phenotype, which is known to be initiated by immune complexes. Markers of the M2b phenotype were significantly increased at the 1 day time-point following anti-A injection, and for two of the three markers (CD86 and Fc R1) peaked at three days before resolving rapidly at the 7 day time-point (Figure 5C). In contrast, Significant increases in M2b markers were observed at three days for the IVIg and mouse IgG injected mice and, for the CD86 and Fc R1, peaked at 7 days before resolving at the 14 day time-point (Figure 5C). Of note for the M2c phenotype, TGF was significantly elevated at the 7 day time-point following anti-A injection but only a small increase was seen in the IVIg and mouse IgG injected mice at the 14 day time-point (Figure 5D). The individual time-point values with statistical significances are shown in table 1.

We have previously shown that MMP2 and MMP9 activation is increased in mice that develop microhemorrhages due to anti-A immunotherapy. Although we did not observe microhemorrhages in the current study, as expected due to the young age of the mice and the intracranial administration paradigm used, we did examine gene expression for the MMP2 and MMP9 systems in order to determine whether IVIg and mouse IgG have different effects on the systems than anti-A antibody. We found that MMP3 (the activator of MMP9) and MMP9 gene expression were significantly elevated 1 and 3 days following injection of anti-A antibody (Figure 6A). The increase in mice injected with IVIg or mouse IgG were peaked at the 7 day time point, however, the increase in expression of MMP3 and MMP9 was significantly less than that observed with anti-A antibody (Figure 6A). The expression of TIMP1, the endogenous inhibitor of MMP9, was increased at 3 days in all groups with no significant difference between any of the antibody injected groups (Figure 6A). Protein

levels of MMP9 and TIMP1, measured by ELISA, matched the time-course of the RNA (Figure 6B). Expression of MMP14 (the activator of MMP2) and MMP2 was significantly increased at the 1 day time-point in mice receiving anti-A antibody injection (Figure 6C). We saw modest increases in MMP14 with IVIg and mouse IgG at the 3 day time-point (Figure 6C). MMP2 gene expression was also increased with IVIg and mouse IgG at the 3 day time-point (figure 6C). TIMP2, the endogenous inhibitor of MMP2, was significantly increased at the 1 day time-point following anti-A antibody injection, and the 3 day timepoint following IVIg and mouse IgG injection (Figure 6C). Interestingly, expression of all genes in the MMP2 system was significantly reduced to almost 0 at the 7 day time-point with anti-A antibody injection before rebounding back to control levels at 14 and 21 days (Figure 6C). This was a phenomenon we did not observe with the IVIg or mouse IgG injection mice. Protein for MMP2 and TIMP2 matched the temporal patterns observed in the gene expression studies, with the exception of the lack of a reduction at the 7 day time-point with anti-A antibody.

Discussion

Intravenous immunoglobulin (IVIg) is a therapeutic approach currently in clinical trials for the treatment of AD. Despite early indicators of some clinical benefit (Relkin et al., 2009), the mechanism(s) of action are unknown. There has been much speculation on the mechanisms, ranging from the presence of circulating anti-A antibodies in IVIg (Dodel et al., 2002), and therefore acting in much the same way as anti-A immunotherapy (Magga et al., 2010), to immune modulation, a known mechanism of action of IVIg utilized in the treatment of autoimmune disorders (Dodel et al., 2010). In the current study we applied a study design previously used to establish mechanism(s) of action of anti-A antibody once it is in the brain (Wilcock et al., 2003). We intracranially administered IVIg, mouse IgG or anti-A antibody, all at the same concentration and volume, bilaterally into the frontal cortex and hippocampus of APP/PS1 transgenic mice and euthanized the mice 1 day, 3 days, 7 days, 14 days and 21 days following injection. Mouse IgG was tested in addition to the human IVIg to determine whether there were any differences between these two groups. Mouse IgG should not contain antibodies to human A protein and, therefore, if differences were observed between the mouse IgG and IVIg then we could conclude that there were unique antibodies in the IVIg contributing to its efficacy. We attempted to detect anti-A antibodies in the mouse IgG preparation by either ELISA or decoration of plaques on tissue sections and were unable to achieve signal.

Overall, we found that IVIg and mouse IgG lower A levels in brain by comparable degrees as that seen with anti-A antibody. However, this reduction occurs over a slower timecourse. We also found that the modulation of the neuroinflammatory response occurs over a slower time-course than anti-A antibody and appears, in some instances, to be longer lasting. These differences in time-course are not accounted for by the presence of the IgG in brain, since all the injected antibodies showed the same clearance kinetics as one another. Our data suggest that IVIg can lower A as effectively as anti-A immunotherapy, but the two therapies likely have distinct mechanisms of action due to the different time-courses. However, it is also feasible that the same mechanisms are engaged with both anti-A antibodies and IVIg/mouse IgG but simply delayed. Immune modulation is one likely mechanism of action suggested by our data. In addition, isotype differences may account for some of the differences between the IVIg/mouse IgG and anti-A antibody groups. Mouse mouse IgG and IVIg compose primarily IgG1 and IgG2 in IVIg and mouse IgG compared to IgG1 only in the anti-A antibody.

Anti-A antibody injection rapidly reduced A burden in the brains of APP/PS1 transgenic mice, with the greatest reduction seen at the 1 day time-point, as previously shown in

Tg2576 mice (Wilcock et al., 2003). In contrast, we did not see significant reductions in A with either the IVIg or mouse IgG until the 7 day time-point, indicating the clearance occurs between the 3 and 7 day period. The difference between the anti-A antibody and the IVIg/ mouse IgG groups suggests that there may be distinct mechanisms of action. We focused on inflammation as an explanation for the differences in A removal because we have previously found a relationship between microglial activation and A clearance by anti-A antibodies injected intracranially (Wilcock et al., 2003; Wilcock et al., 2004a). More recently, we reported that systemically administered anti-A antibody results in altered neuroinflammatory phenotype in the brains of Tg2576 mice (Wilcock et al., 2011b). Most amyloid depositing mice show an M2a phenotype primarily (Colton et al., 2006) and this is what we observed in our saline injected mice. When we injected anti-A antibodies we found increases in M1 genes and M2b genes at the 1 day time-point, peaking at 3 days. There was also a concomitant reduction in M2a gene expression at the 3 day time-point. Both IVIg and mouse IgG showed similar increases in M1 and M2a gene expression, but along a slower time-course, peaking at 7 days.

While macrophages were once broadly grouped into M1 and M2 based on their expression of IL-12 and IL-10 (M1; high IL-12 and low IL-10, M2; low IL-12 and high IL-10), studies found other phenotypes that did not fit this description (Gerber and Mosser, 2001). The M2b macrophage was described in the periphery by Mosser and colleagues several years ago (Edwards et al., 2006). Called the type-II activated macrophage, the group elegantly showed that activation of the Fc receptors by immune complexes switches the macrophage phenotype to this unique state characterized by elevations in some M1 markers, particularly IL-1 , TNF and IL-6, while there was increased IL-10 and no induction of IL-12 (Mantovani et al., 2004; Mosser and Edwards, 2008). There were also some markers that were identified to be more specific for an M2b phenotype, including CD86 (Edwards et al., 2006). CD86 is expressed on antigen presenting cells and is a ligand for CD28 and CD152 on T-cells, leading to T-cell activation (Sansom et al., 2003). We have recently shown that in the AD brain samples, both early stage and late stage, the phenotype that is not observed is M2b (Sudduth et al., 2013a). In the current study, we show that the increased presence of IgG in the brain by intracranial administration, be it IVIg, mouse IgG, or specific anti-A IgG, leads to the modulation of the inflammatory response away from M2a and into an M2b phenotype. We cannot be sure whether the increase in IL-1 , TNF and marco are due to an M2b or M1 phenotype since the primary distinguishing factor between these is the balance of IL-10 and IL-12, and neither are consistently and reliably detectable in the mouse brain tissue. The concomitant increase in M2b specific genes suggests that at least some of the increase in these M1 genes is a part of the M2b response.

Data from mouse models of disease, including neurodegenerative diseases, suggest that altering neuroinflammatory phenotype results in changes in progression of pathology. For instance, Gomez-Nicola et al showed recently that changing the microglial phenotype from M1 to M2a in a mouse model of prion disease slowed neurodegeneration and disease progression (Gomez-Nicola et al., 2013). We have previously shown that lowering of the M2a response while increasing M1 markers is associated with reductions in A with immunotherapy (Wilcock et al., 2011b), however, it is now possible that this was an M2b phenotype as opposed to an M1 based on our current data. More recently, we showed that lithium treatment increased M2a markers and decreased M1 markers in APP transgenic mice and was associated with increased amyloid deposition (Sudduth et al., 2012). Contrasting results were found with the PPAR agonist, pioglitazone, where a switch from M1 to M2 was observed along with decreased plaque load (Mandrekar-Colucci et al., 2012). The contrasting data found in animal studies may be accounted for by non-immunological effects of the drugs being used and suggests a clear need for better understanding of the various neuroinflammatory phenotypes in the brain. An elegant review recently discussed

macrophage diversity in atherosclerosis, and there it was suggested that it is not necessarily the polarization to one phenotype or another, but rather a balance of phenotypes, that is necessary for the progression, or lack thereof, of the pathological process (Mantovani et al., 2009). It is possible that this is the case, too, for the diseased brain.

While microhemorrhages were not observed in the current study due to the age of the mice, and the intracranial administration route selected, they do continue to be a major concern in the development of immunotherapeutics for AD. We have previously shown that MMP2 and MMP9 activation is significantly increased in mice that develop microhemorrhages due to anti-A immunotherapy (Wilcock et al., 2011a). MMP2 and MMP9 are heavily implicated in the pathogenesis of hemorrhagic transformation following ischemic stroke (Klein and Bischoff, 2011) and blood-brain barrier disruption in vascular dementia (Candelario-Jalil et al., 2011). We have also recently shown that MMP2 and MMP9 systems are activated in a mouse model of vascular dementia based on hyperhomocysteinemia (Sudduth et al., 2013b). To begin to assess the potential capacity for IVIg to induce microhemorrhages we assessed the MMP2 and MMP9 systems in the current study by qRT-PCR and ELISA measurement of components of these systems. We found that IVIg, mouse IgG and anti-A antibodies activated both systems, however, a greater increase in the MMP9 system was observed with anti-A antibodies.

In summary, we have shown that IVIg and mouse IgG, when present in the brain, lower A deposition similarly to anti-A antibodies, but with a different time-course. We believe that the immunomodulatory functions of IVIg are active in the brain, turning on an M2b phenotype that may be involved in the reduction of the A deposition. Overall, our data suggests that immunomodulation by IVIg may be beneficial in the treatment of AD.

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Figure 1.

IgG levels in the brain are rapidly decreased between 1 and 3 days following injection. Panel A shows the mouse IgG levels in the right frontal cortex over the time-course from 1 to 21 days. Panel B shows human IgG levels in the right frontal cortex over the time-course form 1 to 21 days. ** indicates P<0.01 compared to saline.

Figure 2.

Brain A is reduced by anti-A antibodies, IVIg and mouse IgG. Panels A-C show the hippocampus of mice receiving saline 1 (A), 3 (B) and 7(C) days following injection. Panels D-F show the hippocampus of mice receiving anti-A antibody 1 (D), 3 (E) and 7 (F) days following injection. Panels G-I show the hippocampus of mice receiving IVIg 1 (G), 3 (H) and 7 (I) days following injection. Panels J-L show the hippocampus of mice receiving mouse IgG 1 (J), 3 (K) and 7 (L) days following injection. In Panel A, the dentate gyrus (DG), cornu ammonis 3 (CA3) and cornu ammonis 1 (CA1) regions are indicated to provide orientation. Asterisks in panels A-L indicate the injection site within the hippocampus. Panels M and N show quantification of A across the full time-course, from 1 to 21 days, in the frontal cortex (M) and hippocampus (N). ** indicates P<0.01 compared to saline for all points under the asterisk. # indicates P<0.05 for IVIg and mouse IgG compared to anti-A antibody.

Figure 3.

Both soluble and insoluble A 1-40 and A 1-42 are reduced by anti-Ab antibodies, IVIg and mouse IgG. Panels A and B show ELISA measurement of A 1-40 (A) and A 1-42 (B) in our soluble protein extract. Panels C and D show ELISA measurement of A -140 (C) and A 1-42 (D) in our insoluble, formic acid protein extract. * indicates P<0.05, ** indicates P<0.01 compared to saline for all points under the asterisk. # indicates P<0.05 for IVIg and mouse IgG compared to anti-A antibody. ^ indicates P<0.05 for IVIg compared to both mouse IgG and anti-A antibody.

Figure 4.

Microglial cells are activated by anti-A antibodies, IVIg and mouse IgG. Panels A-C show the hippocampus of mice receiving saline $1(A)$, $3(B)$ and $7(C)$ days following injection. Panels D-F show the hippocampus of mice receiving anti-A antibody 1 (D), 3 (E) and 7 (F) days following injection. Panels G-I show the hippocampus of mice receiving IVIg 1 (G), 3 (H) and 7 (I) days following injection. Panels J-L show the hippocampus of mice receiving mouse IgG 1 (J), 3 (K) and 7 (L) days following injection. In Panel A, the dentate gyrus (DG), cornu ammonis 3 (CA3) and cornu ammonis 1 (CA1) regions are indicated to provide orientation. Asterisks in panels A-L indicate the injection site within the hippocampus. Panels M and N show quantification of CD45 immunoreactivity across the full time-course, from 1 to 21 days, in the frontal cortex (M) and hippocampus (N) . ** indicates P<0.01 compared to saline for the data point(s) directly under the asterisk. $\#$ indicates P<0.05 the difference between IVIg and mouse IgG compared to anti-A antibody.

Figure 5.

Neuroinflammatory phenotypes are altered by anti-A antibodies, IVIg and mouse IgG. Figure shows relative gene expression for genes representative of the M1 (A), M2a (B), M₂b (C), M₂c (D) phenotypes. Data are shown as fold change relative to APP/PS1 mice receiving saline injection at the given time-point. Due to the multiple statistical differences the statistical significance of the differences is highlighted in table 1.

Figure 6.

Both the MMP9 and MMP2 systems are activated by anti-A antibodies, IVIg and mouse IgG. Panels A and C show relative gene expression of components of the MMP9 system (A) and the MMP2 system (C). Gene expression is shown as fold change compared to mice receiving saline injection at the given time-point. Panels B and D show protein expression of MMP9 and TIMP1 (B), and MMP2 and TIMP1 (D) measured by ELISA. * indicated P<0.05, ** indicates P<0.01 for the data point shown compared to all other data sets at that time-point. Where multiple groups are together, the statistical significance is compared to saline.

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Table 1

Individual data points for the neuroinflammatory marker gene expression. Bolded font shows points that are statistically significantly greater than saline Individual data points for the neuroinflammatory marker gene expression. Bolded font shows points that are statistically significantly greater than saline injected mice at the same time-point, Boxed time-points illustrate the peak change for that gene. Italicized font shows points that are statistically injected mice at the same time-point, Boxed time-points illustrate the peak change for that gene. Italicized font shows points that are statistically significantly lower than saline injected mice at the same time-point. significantly lower than saline injected mice at the same time-point.

