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## A T cell-binding fragment of fibrinogen can prevent autoimmunity

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### Abstract

The C-terminal domain of the fibrinogen  $\gamma$  chain ( $\gamma$ C) has been shown to bind to the integrins  $\alpha$ IIb $\beta$ 3,  $\alpha$ M $\beta$ 2 and  $\alpha$ V $\beta$ 3. It has also been reported that a peptide derived from the  $\alpha$ M $\beta$ 2-binding site of  $\gamma$ C can suppress an animal model of multiple sclerosis, experimental autoimmune encephalomyelitis (EAE). Here we have truncated  $\gamma$ C at position 399 to remove the prothrombotic  $\alpha$ IIb $\beta$ 3-binding site. We show that this truncated version of  $\gamma$ C, termed  $\gamma$ C399tr, can bind to activated T cells. In addition, T cells incubated with  $\gamma$ C399tr secreted less IFN- $\gamma$  when stimulated with antigen and APC; however, cytokine secretion was unaltered when T cells were stimulated non-specifically with a mixture of anti-CD3 and anti-CD28 antibodies. Thus, only antigen-dependent T cell activation is inhibited by  $\gamma$ C399tr. When administered intraperitoneally,  $\gamma$ C399tr potently inhibited actively induced EAE and reversed ongoing disease. We hypothesize that the ability of  $\gamma$ C399tr to inhibit autoreactive immune responses is a result of its ability to bind integrins. This activity was not solely dependent on the  $\alpha$ M $\beta$ 2 integrin-binding site. When polyalanine was substituted for the  $\alpha$ M $\beta$ 2-binding site, the resulting  $\gamma$ C390polyA was still able to inhibit EAE. To our knowledge, this is the first demonstration that T cells can bind to fibrin(ogen), an important extracellular matrix protein that is deposited at sites of inflammation. Our results also identify  $\gamma$ C399tr as a novel therapeutic molecule.

### Keywords

autoimmunity; experimental autoimmune encephalomyelitis (EAE); MOG; multiple sclerosis

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## Introduction

The fibrinogen  $\gamma$  chain has a conserved globular domain at its C-terminus, designated  $\gamma$ C. This region encompasses fibrinogen residues 151-411. According to the crystal structure of the human  $\gamma$ C domain [1],  $\gamma$ C has a C-terminal fibrin-polymerization domain, a single calcium-binding site, and a deep binding pocket. Although this structure lacks a classical integrin-binding motif,  $\gamma$ C has the ability to bind a variety of integrins including,  $\alpha$ IIb $\beta$ 3,  $\alpha$ M $\beta$ 2 and  $\alpha$ V $\beta$ 3. Elimination of residues 400-411 effectively removes the prothrombotic  $\alpha$ IIb $\beta$ 3-binding site. This truncated version of  $\gamma$ C is designated  $\gamma$ C399tr and corresponds to fibrinogen residues 151-399.

Integrins are a family of cell adhesion receptors that recognize extracellular matrix ligands, including fibrin(ogen) as well as many cell surface ligands [2]. Integrins are transmembrane  $\alpha$   $\beta$  heterodimers and at least 18  $\alpha$  and 8  $\beta$  subunits are known. It has been well-established that, after binding to its ligand, an integrin can transduce signals from the outside to the inside of a cell. The activation state of an integrin molecule depends on its conformation as well as the clustering of individual integrin subunits. The activation state can be modulated by signals from inside the cell, a process called “inside-out” signaling [2]. For example, when a T cell is activated integrins on its surface may undergo a conformational change, assuming a more “on” position, extending out from the cell surface. The known integrins that bind to  $\gamma$ C:  $\alpha$ IIb $\beta$ 3 on platelets,  $\alpha$ V $\beta$ 3 on endothelial cells, and  $\alpha$ M $\beta$ 2 on leukocytes, play an important role in thrombus formation, angiogenesis, and inflammation, respectively.

Many integrins are important for normal immune function. Integrin  $\alpha$ 4 $\beta$ 1 binds to the vascular cell-adhesion molecule-1 (VCAM-1) and facilitates extravasation of lymphocytes from the peripheral circulation to sites of inflammation. Integrin  $\alpha$ L $\beta$ 2 is important for the formation of the immunologic synapse. It localizes to the pSMAC portion of the synapse, playing an important role in the ability of a T cell to recognize MHC-bound peptide antigens.

Biologics are a new class of medications that target key components of the immune response and, in some cases, target integrins. For example, natalizumab is directed against the integrin  $\alpha$ 4 $\beta$ 1, present on the surface of lymphocytes. This medication is FDA approved and used to treat patients with multiple sclerosis and inflammatory bowel disease [3,4]. Efalizumab is a biologic directed against the integrin LFA-1 ( $\alpha$ L $\beta$ 2) that until recently was used to treat psoriasis [5]. Given the proven therapeutic potential of integrin-binding biologics, we sought to identify an endogenous integrin-binding molecule that possesses anti-inflammatory properties. Specifically, we focused on  $\gamma$ C because of its ability to bind several integrins, including some important for immune function. Takada and colleagues identified the  $\alpha$ V $\beta$ 3-binding site within  $\gamma$ C [6]. Although  $\alpha$ V $\beta$ 3 is an arginine-glycine-aspartic acid (RGD)-binding integrin, it can also recognize  $\gamma$ C, which lacks a classic integrin-binding motif. The ability of  $\gamma$ C to bind this integrin was of interest because osteopontin also binds to  $\beta$ 3 integrins and this interaction is important in inducing Th1 immune responses [7]. In addition to the  $\alpha$ V $\beta$ 3-binding site, an  $\alpha$ M $\beta$ 2-binding site has also been described within  $\gamma$ C [8]. This integrin is expressed on the surface of microglia and macrophages and it plays a role in their activation and migration.

We hypothesized that the truncated version of  $\gamma$ C,  $\gamma$ C399tr, would be able to bind to leukocytes via their surface integrin molecules and alter the leukocytes' effector function. Here we characterize the cell-binding properties of  $\gamma$ C399tr and demonstrate its ability to inhibit autoimmunity. Specifically, we found that  $\gamma$ C399tr was able to: bind to T cells, inhibit EAE and reverse ongoing autoimmunity.

## Methods

### Peptides

The encephalitogenic myelin oligodendrocyte glycoprotein (MOG) peptide p:35-55 (MEVGWYRSPFSRVVHLYRNGK) and the ovalbumin (OVA) peptide p:323-339 (ISQAVHAAHAEINEAGR) were purchased from A @ A (San Diego, CA). Peptides were purified by HPLC and purity was confirmed by mass spectroscopy.

### Mice

C57BL/6 mice were purchased from the Jackson Laboratory (Bar Harbor, ME) and housed in the UC Davis AAALAC-approved vivarium. OVA-specific TCR Tg OTII mice were also obtained from the Jackson Laboratory. Mice used in all experiments were age-matched female mice housed in laminar flow units and fed autoclaved chow. The UC Davis Institutional Animal Care and Use Committee approved all experiments.

### Experimental autoimmune encephalomyelitis (EAE)

Experimental autoimmune encephalomyelitis was actively induced in C57BL/6 mice by subcutaneous immunization with 150  $\mu$ l of a CFA emulsion containing 150  $\mu$ g of *Mycobacterium tuberculosis* H37Ra (Difco) and 150  $\mu$ g of MOG p:35-55. On days 0 and 2 mice were injected i.p. with 150 ng of purified pertussis toxin (PTX, List Biologicals) dissolved in 0.5 ml PBS. This protocol yields consistent results. EAE severity was scored as follows: 0, no clinical signs of disease or mild tail weakness; 1, complete tail limpness; 2, limp tail and moderate hind limb weakness or unsteady gait; 3, complete hind limb paralysis; 4, hind limb paralysis and some forelimb paralysis; 5, moribund [9]. Mice treated with  $\gamma$ C399tr or  $\gamma$ C390polyA received 100  $\mu$ g daily via intraperitoneal injections starting on day four after MOG p:35-55 immunization. Representative mice were sacrificed and CNS samples were sent for H&E analysis to determine the extent of the CNS pathology.

### Cell culture

The medium employed in all cell culture was RPMI 1640 (Invitrogen), supplemented with  $5 \times 10^{-5}$  M 2-mercaptoethanol (Sigma, St. Louis, MO), 4mM L-glutamine (Invitrogen), 100 U/ml benzylpenicillin (Invitrogen), 100  $\mu$ g/ml streptomycin sulfate (Invitrogen) and 10% heat-inactivated fetal bovine serum (Hyclone). All cultures were incubated at 37°C in a humidified atmosphere of 5% CO<sub>2</sub>. For T cell stimulation assays splenocytes isolated from OT II mice (Jackson Labs) were incubated at  $4 \times 10^5$  cells per well in a 96-well flat bottom plate with indicated amount of peptide antigen. ELISA antibodies were purchased from BD biosciences and the manufacturer's protocol was followed.

### Flow cytometry analysis

All samples were pretreated with Fc-block. PE conjugated antibodies directed against CD3, CD19, and CD11c (BD Biosciences) were used in conjunction with a PE-specific selection kit (Stem Cell Technologies) to isolate T cells, B cells, and dendritic cells from spleen cell suspensions. Alexa 488-labeled  $\gamma$ C399tr was then used to stain purified leukocyte populations for flow cytometry analysis. Peritoneal derived macrophages; a mast cell line, RBL (ATCC); and a human cutaneous T cell lymphoma line, CRL2105, were also stained with Alexa 488-labeled  $\gamma$ C399tr for flow cytometry analysis. Finally, freshly isolated splenocytes were depleted of red blood cells and stained with Alexa 488-labeled  $\gamma$ C399tr. These cells were also stained with antibodies to CD11c, NK1.1, and PDCA-1 conjugated to PE, APC and Alexa 647, respectively. All samples were run on a BD FACSCalibur flow cytometer and acquired data was analyzed using Flowjo 8.7 software.

## Mutant $\gamma$ C399tr

Individual amino acid residues in the NRLSIGE sequence (residues 390-396 of  $\gamma$ C) were simultaneously substituted to alanine as previously described [8].

## Results

### $\gamma$ C399tr binds to leukocytes

In order to determine which cells of the immune system have the ability to bind  $\gamma$ C399tr, PE antibodies directed against CD19 and CD11c were used in conjunction with PE-specific magnetic selection beads to isolate B cells and dendritic cells respectively from spleen cell suspensions. These isolated populations were then stained with Alexa 488-labeled  $\gamma$ C399tr. Figure 1 reveals that  $\gamma$ C399tr-Alexa 488 effectively bound to B cells (Figure 1A). As expected Alexa 488-labeled  $\gamma$ C399tr also bound to peritoneal derived macrophages, which express  $\alpha$ M $\beta$ 2 (Figure 1B). Binding of  $\gamma$ C399tr-Alexa 488 to CD11c-isolated cells revealed two distinct  $\gamma$ C399tr-binding populations, one with brighter staining than the other (Figure 1C). We therefore set out to better characterize the CD11c positive population. Staining with Alexa647-conjugated PDCA-1-specific antibody revealed that the CD11c positive population did not include significant numbers of plasmacytoid dendritic cells (Figure 1D). However, staining red blood cell-depleted splenocytes with APC-conjugated NK1.1-specific antibody and PE-conjugated CD11c specific antibody revealed a CD11c<sup>low</sup>, NK1.1 positive population. This population clearly bound to  $\gamma$ C399tr. Thus, CD11c positive splenocytes are comprised of both myeloid dendritic cells and NK1.1 positive cells, both of which bind to  $\gamma$ C399tr, albeit with different affinities.

Next we stained immortalized cell lines with Alexa 488-labeled  $\gamma$ C399tr. When compared to control, Alexa 488-labeled  $\gamma$ C399tr effectively bound to a macrophage cell line JM774A-1 and poorly to a cutaneous T cell lymphoma cell line (Figure 2). In addition, Alexa 488-labeled  $\gamma$ C399tr did not bind to the mast cell line, RBL (Figure 2).

### $\gamma$ C399tr an Inhibitor of Autoimmunity

After discovering  $\gamma$ C399tr's ability to bind to leukocytes, the next logical step was to determine whether or not this molecule possessed any therapeutic potential. We therefore set out to characterize its effect on the induction of autoimmunity. We hypothesized that  $\gamma$ C399tr would be able to bind to leukocytes via their surface integrin molecules and alter the immune response. C57BL/6 animals were immunized subcutaneously with MOG p:35-55 in a fashion as to induce EAE. Animals were then mixed randomly and divided into two groups, a control group to receive refolding buffer alone and a group to receive  $\gamma$ C399tr treatments. Treatments started on day four after immunization. Animals were treated daily until the end of the experimentation on day 25. Table 1 reveals that none of the  $\gamma$ C399tr-treated animals developed EAE (0/6). Four of the six control animals treated with refolding buffer developed severe and chronic EAE (maximum disease scores: 5,4,4,2,0,0). All animals were scored blindly.

Central nervous system tissue samples with coded labels were also sent to a pathologist for Hematoxylin and Eosin (H&E) staining and analysis (Figure 3). The pathologist was blinded, unaware of the treatment groups. All samples from the  $\gamma$ C399tr-treated group were read as having no CNS inflammation. CNS samples isolated from the control mice treated with refolding buffer were read as positive for CNS inflammation. The pathologist photographed the H&E slides and representative slides are shown in Figure 3A-3F.

To further characterize the anti-inflammatory properties of  $\gamma$ C399tr we then designed a crossover study in which MOG p:35-55 immunized animals were again divided randomly

into two groups, a control group receiving refolding buffer alone and a  $\gamma$ C399tr-treated group. Then, on day 16 after MOG p:35-55 immunization, treatments were switched; the control group receiving refolding buffer was switched to  $\gamma$ C399tr treatments and the  $\gamma$ C399tr-treated group began to receive refolding buffer alone. Figure 4 shows the results of our crossover study. Prior to day 16 none (0/8) of the  $\gamma$ C399tr-treated mice developed EAE, while seven out of eight animals in the control group developed severe disease (maximum disease scores: 4,4,4,4,4,3,0). In contrast, after switching therapies on day 16, four of the eight mice initially receiving  $\gamma$ C399tr developed EAE (maximum disease scores: 4,3,2,2,0,0,0,0); and all of the animals that had initially developed EAE recovered once they began receiving  $\gamma$ C399tr-treatment (Figure 4). Coded CNS samples were again sent to a pathologist for H&E analysis. The results confirmed our clinical scores. On day 25 no CNS inflammation was detected in any of the mice receiving  $\gamma$ C399tr since crossover day 16 (Figure 3). Thus,  $\gamma$ C399tr possesses strong anti-inflammatory properties and may be a potential therapeutic agent for the treatment of states of chronic inflammation.

### $\gamma$ C399tr inhibits activated T cell

Since EAE is known to be a CD4<sup>+</sup> T cell mediated autoimmune disease we sought to further characterize the specific effects that  $\gamma$ C399tr has on activated T cells. The first step was to determine if  $\gamma$ C399tr had a differential binding capacity for naïve and activated T cells. To do this spleen cells obtained from C57BL/6 mice were incubated with anti-CD28 and anti-CD3 antibodies for 48 hours to activate the resident T cells. Following activation PE conjugated anti-TCR antibodies were used in conjunction with a PE selection kit (StemCell Technologies) to isolate activated T cells. Naïve T cells were also isolated from fresh spleen using the same PE-selection kit. The cells were then stained with  $\gamma$ C399tr-Alexa 488 and analyzed by flow cytometry. Figure 5 shows that  $\gamma$ C399tr binding is increased on activated T cells. With this in mind we then sought to identify the physiologic effect that  $\gamma$ C399tr binding had on T cells, using the OT II TCR transgenic mouse model [11]. These mice have T cells specific to the OVA peptide, p:323-339. Splenocytes isolated from OT II mice were incubated with increasing concentration of  $\gamma$ C399tr in the presence or absence of OVA peptide at a concentration of 5  $\mu$ g/ml. Results shown in Figure 6 reveal that  $\gamma$ C399tr effectively inhibits IL-4, IFN- $\gamma$ , and IL-17 produced by OT II T cells in response to antigen (Figures 6A, 6B and 6C, respectively). To determine if this inhibition was due to a nonspecific effect of  $\gamma$ C399tr, we conducted a similar experiment in which splenocytes were incubated with increasing concentrations of  $\gamma$ C399tr in the presence or absence of anti-CD3 and anti-CD28 antibodies. Figure 6D reveals that  $\gamma$ C399tr was unable to inhibit T cell cytokine secretion when cells were stimulated non-specifically in an antigen independent fashion. This rules out a non-specific toxic effect of  $\gamma$ C399tr on cultured cells. From this data we hypothesize that the ability of  $\gamma$ C399tr to inhibit autoimmunity is partly due to its ability to inhibit activated T cells.

### Mutating the $\alpha$ M $\beta$ 2 -binding site within $\gamma$ C399tr does not eliminate its therapeutic potential

Importantly, the Mac-1 binding site within  $\gamma$ C399tr has been well-characterized and appears to be distinct from the  $\alpha$ v $\beta$ 3-binding site [12]. Alanine substitutions within the Mac-1 binding site have been shown to abolish Mac-1 binding [8]. To determine if the protective effect of  $\gamma$ C399tr is dependent on this Mac-1-binding site we mutated this site within  $\gamma$ C399tr to alanine. The resulting  $\gamma$ C390PolyA was then administered to mice once daily beginning on day four after EAE induction with MOG p:35-55. Similar to the results obtained with wild-type  $\gamma$ C399tr, the  $\gamma$ C390PolyA molecule was effective at inhibiting EAE. As shown in Figure 7 only 2/6 mice developed EAE. Thus, even in the absence of an  $\alpha$ M $\beta$ 2-binding site  $\gamma$ C399tr possessed some inhibitory properties.

## Discussion

Fibrinogen is deposited at sites of tissue injury, including sites of inflammation. The ability of cells to migrate through the deposited fibrin scaffold is key to a variety of processes. Integrins bind to extracellular matrix proteins, including fibrin(ogen). Given the importance of this interaction one might predict that the integrin-binding motifs within fibrinogen would be conserved evolutionarily. However, the classic RGD-integrin-binding motif is not conserved within fibrinogen across different species. Thus, it was hypothesized that fibrin(ogen) likely contains other integrin-binding sequences. The  $\gamma$ C region of fibrinogen is conserved evolutionarily and has been found to bind a variety of integrins. We have sought to determine the leukocyte populations capable of binding to  $\gamma$ C399tr and to characterize the effect that binding of soluble  $\gamma$ C399tr has on the immune response. Although we were unable to prove that the  $\gamma$ C399tr-leukocyte interactions were mediated through cell surface integrins, a variety of circumstantial evidence was found in support of this possibility. For one, the fibrinogen fragment  $\gamma$ C399tr has been already been shown to bind to a variety of integrins, some of which are present on immune cells. For example, an  $\alpha$ M $\beta$ 2-binding site within  $\gamma$ C399tr has been well-characterized [8]. Not surprisingly,  $\gamma$ C399tr bound well to macrophages, which express  $\alpha$ M $\beta$ 2 (Figure 1B). However, interactions between fibrin fragments and integrins on the surface of T cells, B cells, and dendritic cells have not been reported. Given that fibrin and its degradation products are deposited perivascularly at sites of inflammation [10], one would expect that such an interaction would enhance T cell and B cell migration and navigation through inflammatory tissue. It would also facilitate cellular interactions between these cells and growth factors bound to the deposited fibrin scaffold. Activated integrins extend out from the surface of the cell, and are more accessible for ligand binding. When T cells are activated or are exposed to chemokines their integrins also assume a more active state;  $\gamma$ C399tr bound with a higher affinity to activated T cells. Integrins also participate in antigen recognition by helping the T cell bind to the antigen presenting cell (APC). They form part of the immunologic synapse, which is required for antigen-dependent T cell activation. The formation of this synapse is not required when T cells are stimulated with anti-CD3 and anti-CD28;  $\gamma$ C399tr was only able to inhibit antigen dependent T cell activation. Lastly, it appeared that the integrin expression level correlated with the binding capacity of  $\gamma$ C399tr. Cells that expressed higher concentrations of integrins bound more strongly to  $\gamma$ C399tr. Together this data identifies  $\gamma$ C399tr-integrin interactions as one possible mechanism for  $\gamma$ C399tr binding to leukocytes.

This manuscript demonstrates that  $\gamma$ C399tr is a potent inhibitor of EAE, the animal model of MS. It is possible that some of this protection is mediated through the  $\alpha$ M $\beta$ 2 binding site within  $\gamma$ C399tr. However, this site is not solely responsible for all of the protective effects of  $\gamma$ C399tr. When mutated to polyalanine the resulting  $\gamma$ C390polyA was still able to inhibit EAE induction (Figure 7). Thus, other properties of  $\gamma$ C399tr are likely contributing to its role in EAE inhibition. One possibility is that  $\gamma$ C399tr-has an inhibitory effect on T cells. The secretion of T cell cytokines IFN- $\gamma$  and IL-17, known to be important for EAE induction [13-15], were inhibited by soluble  $\gamma$ C399tr (Figure 6).

Since fibrinogen degradation products are deposited at sites of inflammation, the ability to bind to fibrin(ogen) would allow leukocytes to better migrate through inflammatory sites. Why then was soluble  $\gamma$ C399tr able to inhibit autoimmunity? In contrast to a fibrin scaffold, binding of soluble  $\gamma$ C399tr would not help leukocytes orient themselves for migration. In addition,  $\gamma$ C399tr was able to inhibit antigen-dependent T cell activation, something a fibrin scaffold would not be expected to do.

Importantly, our results reveal a novel approach to identify integrin-binding mediators of autoimmunity. We have focused on identifying “self” molecules rather than identifying

monoclonal integrin-blocking antibodies. Integrins have relatively low affinities for their ligands ( $K_d=10^{-6}$  to  $10^{-7}$  M). Thus, any interaction between  $\gamma$ C399tr and its integrin ligands is likely quite different than that of a high affinity specific monoclonal antibody. Although the broad binding specificity of this ligand might limit its potential for drug development, we are hopeful that this study will aid in the design of future biologic medications.

## Acknowledgments

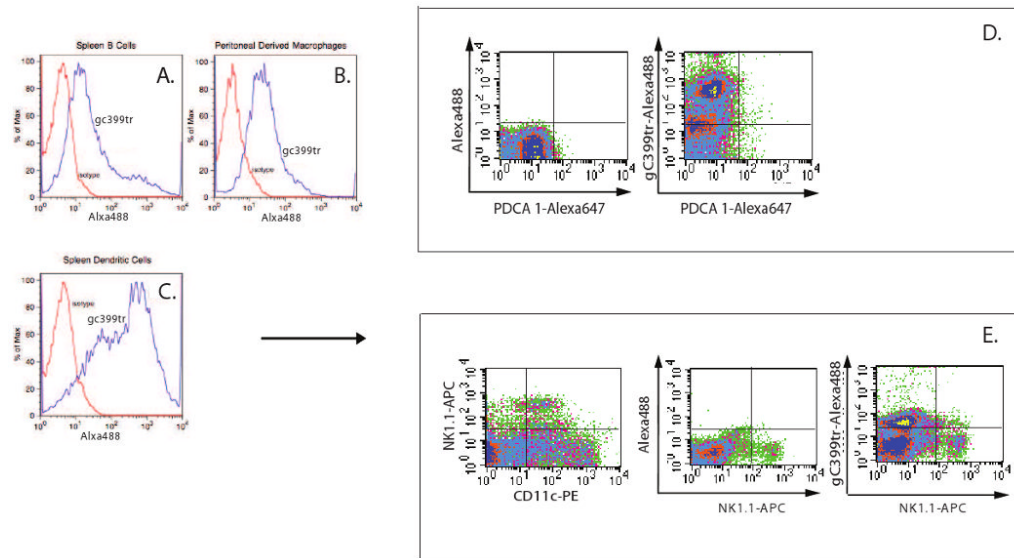
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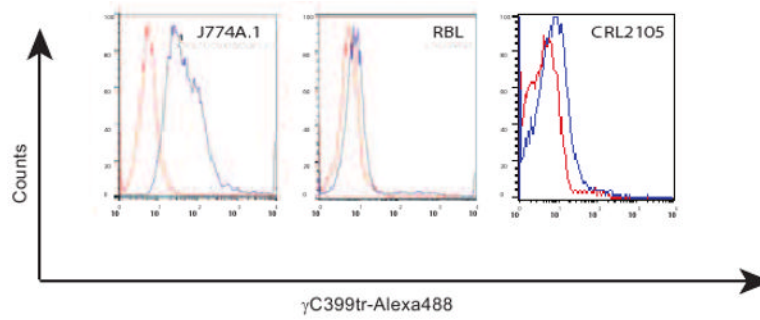
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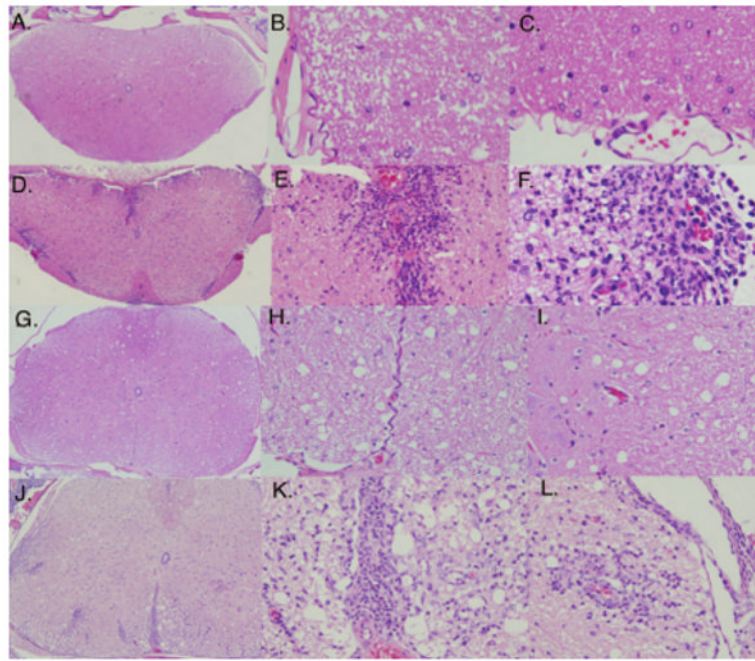
**Figure 1.  $\gamma$ C399tr binds to B cells, macrophages, and dendritic cells**

A) B cells were purified from splenocytes using a PE-labeled anti-CD19 antibody and a PE selection kit. Staining with Alexa 488-labeled  $\gamma$ C399tr revealed that this population effectively bound to  $\gamma$ C399tr. B) Peritoneal derived macrophages were further purified with a PE-labeled F4/80 antibody and a PE selection kit. Staining with Alexa 488-labeled  $\gamma$ C399tr revealed that this population also effectively bound to  $\gamma$ C399tr. C) CD11c<sup>+</sup> cells were purified from splenocytes using a PE-labeled anti-CD11c antibody and a PE selection kit. Staining with Alexa 488-labeled  $\gamma$ C399tr revealed that this population effectively bound to  $\gamma$ C399tr with a bimodal distribution. D) To further characterize the two CD11c<sup>+</sup> populations, splenocytes were depleted of red blood cells and then stained with PE-conjugated anti-CD11c, Alexa 488-conjugated  $\gamma$ C399tr and Alexa 647-conjugated anti-PDCA-1. This revealed that there were almost no contaminating PDCA-1<sup>+</sup> cells in the CD11c<sup>+</sup> population. Again, CD11c<sup>+</sup> cells bound well to Alexa 488-conjugated  $\gamma$ C399tr. E) Splenocytes were then double stained with PE-conjugated anti-CD11c and APC-conjugated anti-NK1.1. Results show that a population of CD11c<sup>low</sup> NK1.1<sup>+</sup> cells exists in the spleen. These cells bound poorly to  $\gamma$ C399tr.



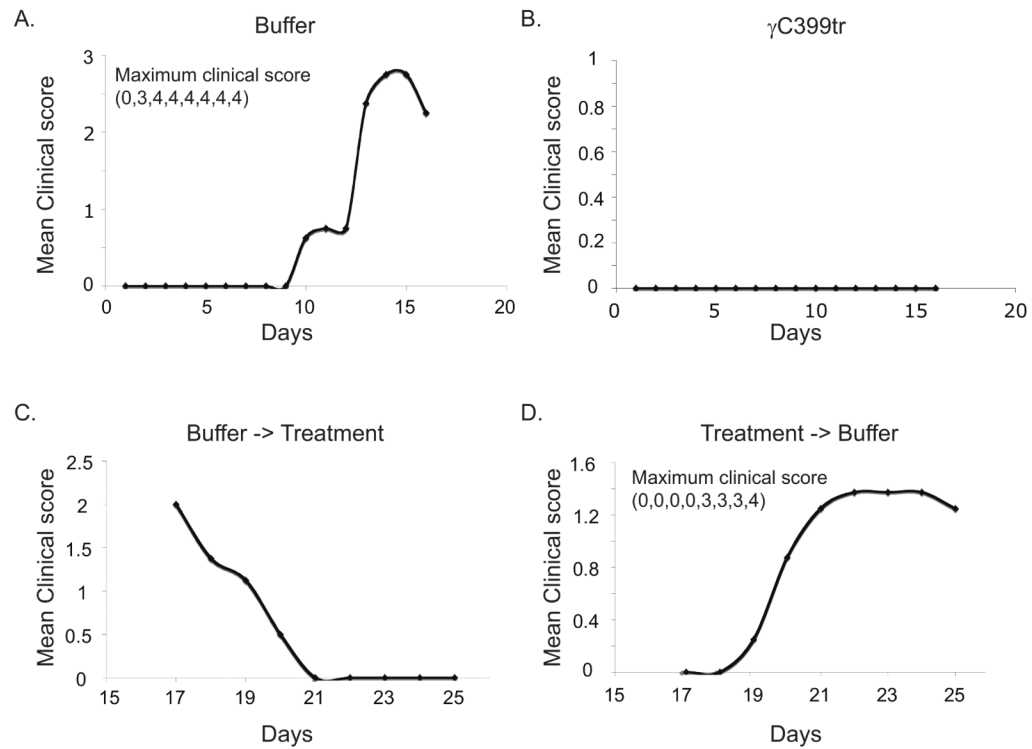
**Figure 2. A macrophage cell line binds to  $\gamma$ C399tr**

Alexa 488-labeled  $\gamma$ C399tr effectively bound to a macrophage cell line JM774A.1 and bound weakly to a cutaneous T cell lymphoma, CRL2105. However, Alexa 488-labeled  $\gamma$ C399tr did not bind to a mast cell line, RBL.



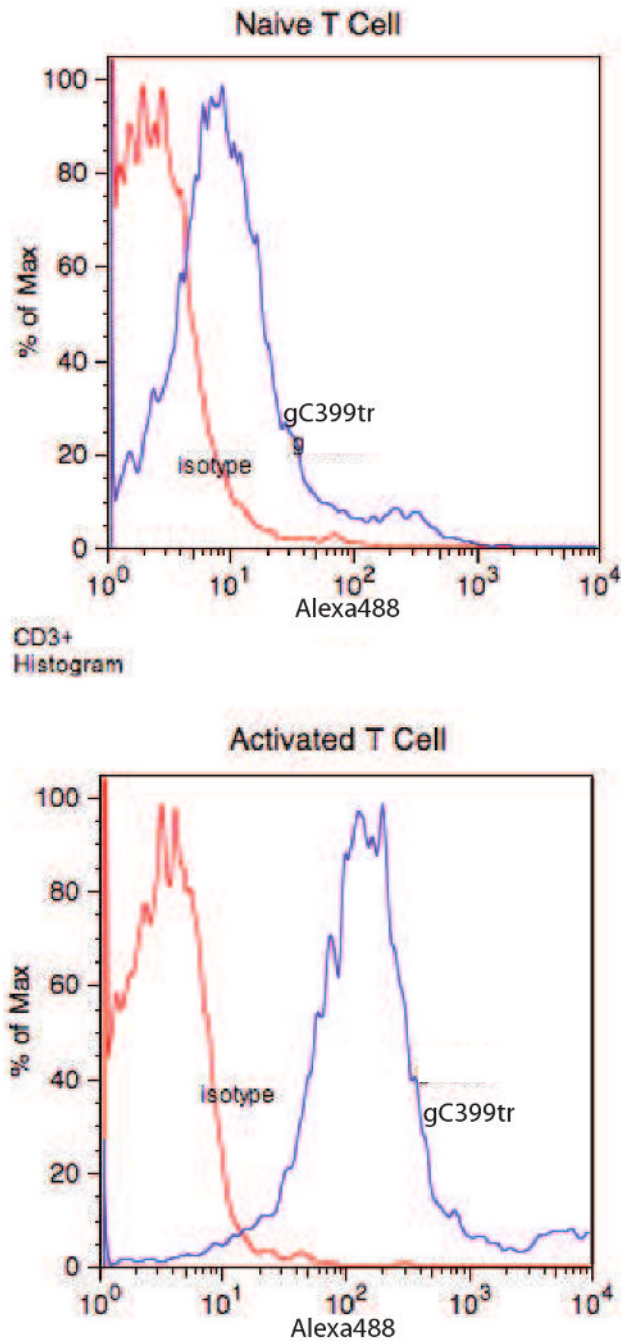
**Figure 3.  $\gamma$ C399tr protects animals from EAE induction**

Coded CNS samples from mice depicted in Table 1 were paraffin-embedded and stained with H&E. A representative from each group is shown in images 3A-3F. All mice treated with  $\gamma$ C399tr (3A-3C) were free of CNS inflammatory cells. In contrast, marked inflammation was seen in mice treated with refolding buffer alone (3D-3F). In 3D scanning magnification reveals multiple sites of perivascular inflammation. At closer magnification (3E and 3F) the CNS inflammation can be identified as mixed cellular (macrophages and lymphocytes). G-L) A crossover experiment was then performed in which control mice receiving refolding buffer were switched to  $\gamma$ C399tr and  $\gamma$ C399tr-treated animals were switched to refolding buffer. CNS samples were obtained from mice in both groups at the end of the experiment, day 25. CNS samples obtained from mice in the refolding buffer to  $\gamma$ C399tr treatment group completely lacked CNS inflammation [3G (scanning), 3H and 3I (closer magnification)]. CNS samples from the  $\gamma$ C399tr to refolding buffer group 3J-3L had marked CNS perivascular inflammatory cells [3J (scanning), 3K and 3L (closer magnification)]. A pathologist blinded to treatment groups read all H&E samples.



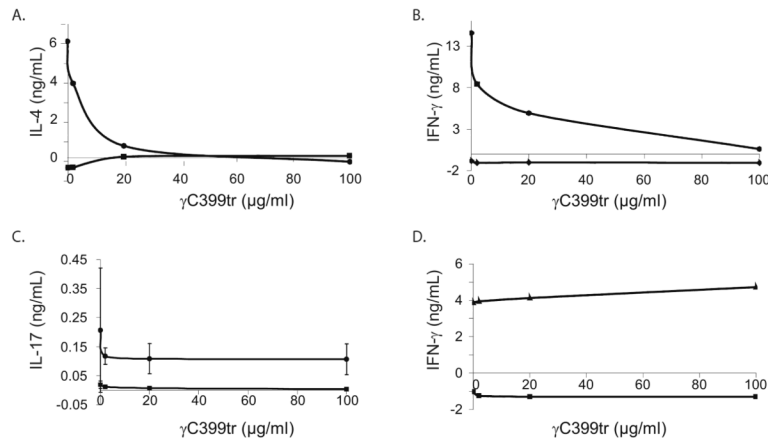
**Figure 4.  $\gamma$ C399tr protects mice from actively induced EAE**

Sixteen C57BL/6 mice were immunized subcutaneously with MOG p:35-55 in an encephalitogenic fashion. On day 4, mice were randomly divided into two groups of eight mice each. One group received daily injections of 100  $\mu$ g of  $\gamma$ C399tr while the other group received an equal volume of refolding buffer alone. The degree of paralysis was then scored daily as described in the materials and methods. On day 16, treatments were crossed; animals initially receiving refolding buffer started to receive  $\gamma$ C399tr and vice versa. Treatment injections were given in the morning and animals scored blindly in the evening. Averages for each group are graphed as days post injection versus EAE disease score. *4A*) Upper left- seven of eight mice receiving control treatments with buffer alone developed EAE. *4B*) Upper right- mice receiving  $\gamma$ C399tr treatment were protected from EAE induction (0/8) ( $P < 0.00001$ , student T test). *4C*) Lower left- once animals developed EAE i.p. treatments with  $\gamma$ C399tr could reverse their active disease. *4D*) Lower right- four of the eight mice that were initially protected by  $\gamma$ C399tr treatments developed EAE once this therapy was switched to refolding buffer.



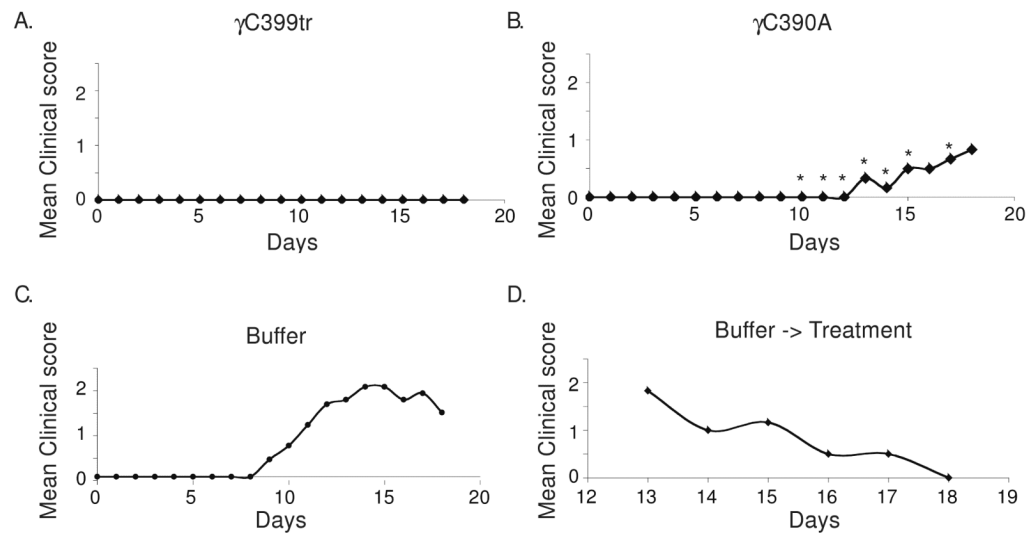
**Figure 5.  $\gamma$ C399tr binds with increased affinity to activated T cells**

Naïve T cells and T cells activated with a mixture of CD3 and CD28 antibodies were stained with Alexa 488-labeled  $\gamma$ C399tr. Upper histogram reveals that naïve T cells bound poorly to  $\gamma$ C399tr. In contrast, activated T cells bound very strongly  $\gamma$ C399tr (lower histogram).



**Figure 6.  $\gamma$ C399tr can inhibit cytokine production by activated T cells**

Splenocytes isolated from OTII Tg mice were *in vitro* cultured with either OVA peptide 5  $\mu$ g/ml (6A-6C) or anti-CD3/anti-CD28 antibodies (6D) in the presence of increasing concentrations of  $\gamma$ C399tr. 6A) Upper left- As the concentration of  $\gamma$ C399tr is increased OVA-specific T cells secrete less IL-4 (circles). OTII cells incubated with control peptide did not secrete IL-4 (squares). 6B) Upper right- As the concentration of  $\gamma$ C399tr is increased OVA-specific T cells secrete less IFN $\gamma$  (circles). OTII cells incubated with control peptide did not secrete IFN $\gamma$  (diamonds). 6C) Bottom left- As the concentration of  $\gamma$ C399tr is increased OVA-specific T cells secrete less IL-17 (circles). OTII cells incubated with control peptide did not secrete IL-17 (squares). 6D) Bottom right-  $\gamma$ C399tr DID NOT inhibit IFN $\gamma$  production by OTII T cells stimulated nonspecifically with anti-CD3 and anti-CD28 antibodies (circles). As the concentration of  $\gamma$ C399tr is increased secretion of IFN $\gamma$  is unchanged (triangles). OTII cells incubated with control peptide did not secrete IFN $\gamma$  (squares).



**Figure 7. The  $\alpha$ M $\beta$ 2-binding defective mutant  $\gamma$ C390polyA effectively inhibits EAE**  
 EAE was induced using the same protocol as in Figure 4.  $\gamma$ C399tr and  $\gamma$ C390polyA were able to effectively inhibit EAE (7A upper left and 7B upper right, respectively. Asterix represents  $P < 0.05$ , student T test). 7C) Lower left- mice treated with buffer alone as a control developed severe EAE. 7D) Lower right- animals that had initially developed EAE were able to recover after administration of  $\gamma$ C399tr. This figure better highlights the chronic nature of the untreated animals.

**Table 1**

Treatment	Maximum Disease score	Mean Day of Onset
$\gamma$ C399tr 100 $\mu$ g	(0,0,0,0,0,0)	NA
Refolding Buffer	(5,4,4,2,0,0)	12.25

(\*Mice were scored blindly.)