Glatiramer acetate treatment negatively regulates type I interferon signaling

Glatiramer acetate (GA; Copaxone), a random basic copolymer, is prescribed for treatment of relapsing-remitting multiple sclerosis (MS). Previously, it was demonstrated that GA therapy induces anti-inflammatory M2 (type II) myeloid cells.\(^1\)\(^-\)\(^3\) GA-induced M2 monocytes reversed ongoing paralysis in experimental autoimmune encephalomyelitis (EAE) and, as...
antigen-presenting cells (APC), promoted expansion of regulatory T cells (Treg) and T helper (Th) 2 cells. Thus, induction of M2 APC is a proximal event responsible for T-cell regulation associated with GA therapy.

While the molecular mechanisms required for GA-mediated M2 differentiation and therapeutic efficacy of GA have not been fully elucidated, it is recognized that GA inhibits monocyte reactivity in response to engagement of Toll-like receptors (TLRs), suggesting that modulation of innate signaling could represent a principal mechanism of action of GA. In general, TLR engagement triggers activation through interaction with adaptor molecule myeloid differentiation primary response gene 88 (MyD88) or Toll-IL-1 receptor domain-containing adaptor inducing interferon (IFN)–β (TRIF). Both pathways are important in CNS autoimmunity. MyD88-dependent signaling is associated with activation of the nuclear factor (NF)–κB pathway and is required for Th17 differentiation in EAE. In contrast, signaling via TRIF leads to activation of IFN regulatory factor 3 (IRF3) transcription factor and subsequent production of IFN-β. Signaling through TRIF, IRF3, or the type I IFN receptor (IFNAR) also influences development of Th17 cells and EAE. Thus, reduction of proinflammatory cytokines observed following stimulation of M2 monocytes could indicate that GA modulates MyD88, TRIF, or both pathways.

The goal of this investigation was to identify the molecular pathways participating in differentiation of M2 APC by GA. Our results, which demonstrate that GA downregulates type I IFN signaling, provide a key mechanism responsible for inhibition of proinflammatory APC polarization by GA treatment.

**METHODS**

**Mice.** Female 8- to 10-week-old C57BL/6 wild-type (WT), IFN-α/β receptor subunit-1 (IFNAR1)–deficient, TRIF-deficient, and CD11b-deficient mice were obtained from Jackson Laboratories (Bar Harbor, ME). C57BL/6 MyD88-deficient mice were provided by Dr. A. DeFranco (University of California, San Francisco). Mice were maintained and handled in accordance with UCSF Institutional Animal Care and Use Committee regulations for the humane use of animals in research.

**EAE induction.** C57BL/6J mice were injected SC with 50 μg MOG p35-55 (Auspep, Parkville, Australia) emulsified in complete Freund’s adjuvant containing 200 μg heat-killed Mycobacterium tuberculosis on day 0. Mice received 200 μg Bordetella pertussis toxin (List Biological Laboratories, Campbell, CA) IV in 0.2 mL phosphate-buffered saline (PBS) on days 0 and 2. Mice were observed daily. Mice received one SC injection of a 0.1-mL emulsion consisting of 250 μg GA (provided by Teva Pharmaceutical Industries, Petah Tikva, Israel) in an equal volume of PBS and incomplete Freund’s adjuvant in the upper flanks on the same day of immunization (day 0). Clinical scores were assessed as follows: 0 = no clinical disease, 1 = loss of tail tone only, 2 = mild monoparesis or paraparesis, 3 = severe paraparesis, 4 = paraplegia or quadriplegia, and 5 = moribund or death.

**Monocyte isolation and culture.** CD11b+/CD11c– macrophages (purity >98%) were obtained from C57BL/6 WT, MyD88-deficient, IFNAR1-deficient, TRIF-deficient, and CD11b-deficient mice. Bone marrow–derived monocytes (BMDM) were obtained following in vitro GA or vehicle (PBS) treatment, for 7 days in RPMI, 10% FCS, and 10% L929 cell-conditioned medium (L929-CM), which was used as a source of murine M-CSF. Mouse BMDM and peripheral macrophages stained positively for CD11b, F4/80, and major histocompatibility complex (MHC) class II (I-A) cell surface molecules. For cyclic adenosine 3′,5′-monophosphate (cAMP) quantification, DNA binding, and ELISA assays, monocytes were cultured in 96-well plates at 2.5 × 10⁶ cells/well in 200 μL culture media. For Western blot analyses, monocytes differentiated in the presence or absence of GA were resuspended at 10 × 10⁶ cells/mL in culture media, and 2 mL were plated in 6-well plates at 37°C for 1 hour, then activated with indicated stimuli. Human peripheral blood buffy coat monocytes were isolated from healthy volunteers as previously described. Human monocytes expressed cell surface CD14, MHC class II (HLA-DR), and CD68 molecules. Monocytes were cultured in 96-well plates at 5 × 10⁶ cells/well in 200 μL media, and activated with the indicated stimuli for 24 hours. Monocytes were stimulated with forskolin (an agonist of adenylyl cyclase and serves as a positive control for induction of intracellular cAMP), lipopolysaccharide (LPS), Poly(I:C), lipoteichoic acid (LTA), and Pam3CSK4, which were purchased from InvivoGen (San Diego, CA).

**ELISA.** Production of tumor necrosis factor (TNF), interleukin (IL)–6, and IFN-β was measured in serum or culture supernatants by ELISA (eBioscience, San Diego, CA; R&D Systems, Minneapolis, MN), per manufacturer protocol. Intracellular cAMP (cAMP) was detected using Biotrack kit (GE Healthcare, Cleveland, OH).

**DNA binding.** DNA binding of ATF-2 and IRF3 was quantitated from monocyte nuclear extracts using the ELISA-based Transfactor (Clontech, Mountain View, CA) and TransAM (Active Motif, Carlsbad, CA) kits, respectively.

**Western blot.** Total cell lysates were prepared from monocyte cultures and subjected to Western blot analysis as described previously. Nitrocellulose membranes were probed (4°C, overnight) with antibodies against phospho-Akt (Ser473), phospho-JNK (Thr183/Tyr185), phospho-p38 MAP kinase (Thr180/Tyr182), phospho-IκBα (Ser32/36), glyceraldehyde-3-phosphate dehydrogenase (GAPDH), total STAT1, and total Akt1/2/3 (Cell Signaling Technology, Danvers, MA). Horseradish peroxidase–conjugated goat anti-mouse (Thermo Scientific, Waltham, MA) or sheep anti-rabbit (Coslab, Villeurbanne, France) antibodies were used for secondary detection. Antibody-bound proteins were detected using horseradish peroxidase–conjugated secondary antibodies (anti-rabbit, Covalab, Villeurbanne, France) and reaction with a chemiluminescent substrate (Immobilon Western Chemiluminescent HRP Substrate, Millipore, Billerica, MA).

**Intracellular cAMP analysis.** cAMP production was measured using the cAMP ELISA kit with an intra-assay CV of less than 10% (Coat-ELISA, Nottingham, UK).

**Intracellular DNA binding assay.** Nuclear extracts were prepared as previously described. DNA binding to the NFκB consensus sequence was assessed using the ELISA-based Transfactor kit (Clontech, Mountain View, CA).

**Western blot analysis.** Total cell lysates were prepared from monocyte cultures and subjected to Western blot analysis as described previously. Nitrocellulose membranes were probed (4°C, overnight) with antibodies against phospho-Akt (Ser473), phospho-JNK (Thr183/Tyr185), phospho-p38 MAP kinase (Thr180/Tyr182), phospho-IκBα (Ser32/36), glyceraldehyde-3-phosphate dehydrogenase (GAPDH), total STAT1, and total Akt1/2/3 (Cell Signaling Technology, Danvers, MA). Horseradish peroxidase–conjugated goat anti-mouse (Thermo Scientific, Waltham, MA) or sheep anti-rabbit (Coslab, Villeurbanne, France) antibodies were used for secondary detection. Antibody-bound proteins were detected using horseradish peroxidase–conjugated secondary antibodies (anti-rabbit, Covalab, Villeurbanne, France) and reaction with a chemiluminescent substrate (Immobilon Western Chemiluminescent HRP Substrate, Millipore, Billerica, MA).

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by the Uptight HRP Blot Chemiluminescent substrate (GE Healthcare). Western blots were digitized and quantified using NIH imageJ software (version 1.44).

**Mouse phospho-Akt (Ser473) cell-based ELISA.** PathScan Phospho-Akt1 (Ser473) Sandwich ELISA Kit (Cell Signaling Technology) was used to detect endogenous phospho-Akt1 (Ser473) protein. Equal concentrations of protein extract (100 μg/mL) were analyzed for each sample. The magnitude of absorbance (450 nm) was proportional to the quantity of phospho-Akt1 (Ser473) protein. Values were expressed as a fraction of the amount of phospho-Akt1 (Ser473) observed in BMDM extract, which was normalized to 1.

**Statistics.** Data are presented as the mean ± SD. For ELISA data, n = 3 separate samples for each condition were examined. When required, statistical significance of differences between groups was evaluated using Student paired t test. Differences were considered significant at a value of p < 0.05. For clinical scores, significance between groups was examined using the Mann-Whitney U test. A value of p < 0.05 was considered significant.

**RESULTS**

**GA induces activation of PI3K, but not cAMP.** Whether GA selectively engages specific receptors is unclear. cAMP and phosphoinositide 3-kinase (PI3K) are second messenger systems that can be activated after receptor engagement, and have been associated with early signaling events in anti-inflammatory responses. As shown in figure 1A, incubation of monocytes with GA did not induce intracellular cAMP levels, as compared to stimulation by the positive control, adenylylcyclase activator forskolin, suggesting that GA-induced M2 differentiation does not involve cAMP. Phosphorylation of Akt on serine 473 is a surrogate marker of PI3K activation. Within 15 minutes, GA induced phosphorylation of Akt as measured both by Western blot (figure 1B) and ELISA (figure 1C). These results suggest that GA activates at least one intracellular second messenger signaling pathway associated with reduction of inflammation in innate immune cells.

**GA inhibits TRIF-dependent, but not MyD88-dependent, signaling.** MyD88 and TRIF represent the 2 essential adapters of innate immune signaling. Previous data demonstrated that GA treatment was associated with reduction of proinflammatory cytokine production and STAT1 phosphorylation in monocytes following stimulation with LPS, a TLR4 agonist. All TLRs utilize MyD88 except TLR3, which depends exclusively on TRIF. TLR4 signals via both MyD88 and TRIF. Therefore, it was possible that GA could interfere with MyD88 or TRIF. We employed distinct TLR agonists in order to distinguish between these 2 possibilities. First, we observed that GA itself did not alter baseline or ligand-induced expression of TLR2, TLR3, or TLR4 (figure e-1 at Neurology.org/nn), indicating that GA did not influence the capacity to sense TLR agonists. As shown in figure 2A and figure e-2, activation of GA-treated monocytes by the

![Figure 1](image-url)
TLR2 ligands, Pam3CSK4 or LTA, did not alter TNF and IL-6 secretion. In contrast, secretion of these proinflammatory cytokines was reduced when GA-treated monocytes were activated by the TLR3 ligand, Poly(I:C), or the TLR4 ligand, LPS. These results therefore suggested that GA inhibited TRIF-dependent signaling and was independent of MyD88. Similar to monocytes from WT mice, monocytes from MyD88-deficient mice exhibited reduced IL-6 and TNF levels following GA treatment (figure 2A). Engagement of MyD88 activates NF-κB. As shown in figure 2B, GA exposure did not impair phosphorylation and degradation of the inhibitor protein IκBα following LPS treatment. GA exposure also did not significantly alter DNA binding of p50 or p65 NF-κB subunits.

Figure 2  Glatiramer acetate treatment induced M2 differentiation through a MyD88-independent pathway

(A) As described previously,3 M2 monocytes were treated in the presence or absence of glatiramer acetate (GA) for 6 days. They were then stimulated with lipopolysaccharide (LPS), Poly(I:C), or Pam3CSK4 for 24 hours. (B) Wild-type (WT) monocytes cultured in the presence or absence of GA were stimulated with LPS (100 ng/mL) for the indicated duration. Cell lysate proteins were separated by sodium dodecyl sulfate polyacrylamide gel electrophoresis and membranes were probed for phosphorylated IκBα (Ser32/36). Data are representative of 2 separate experiments. (C) Human peripheral blood monocytes were preincubated for 1 hour with or without 50 μg/mL GA and then cultured for 24 hours in the presence or absence of Poly(I:C) (10 μg/mL) or Pam3CSK4 (100 ng/mL). Tumor necrosis factor (TNF) (left panels) and interleukin (IL)–6 (right panels) secretion was quantitated in cell supernatants by ELISA. Results are presented as mean ± SD (n = 3); **p < 0.01, ***p < 0.001 by Student t test. Data presented are representative of 3 independent experiments. MyD88 = myeloid differentiation primary response gene 88; TRIF = Toll-IL-1 receptor domain-containing adaptor inducing interferon-β.
(data not shown). These results suggest that the NF-κB pathway is dispensable for the immunologic effects of GA on monocytes. Further, GA did not reduce proinflammatory cytokine secretion in TRIF-deficient monocytes (figure 2A), providing further support that inhibition of TRIF-dependent signaling is required for GA-mediated reduction of proinflammatory cytokine secretion. As GA treatment of murine monocytes suppressed proinflammatory cytokine production by Poly(I:C), but not Pam3CSK4, we tested how GA-treated human monocytes responded to these 2 TLR ligands. GA treatment of human peripheral blood monocytes decreased TNF and IL-6 secretion induced by Poly(I:C), but not Pam3CSK4 (figure 2C). Thus, as with murine monocytes, these results indicate that GA inhibits proinflammatory cytokine production by human monocytes by blocking TRIF-mediated signaling.

The in vivo contribution of TRIF-dependent signaling in modulating proinflammatory cytokine secretion was also examined during GA treatment. GA-treated WT mice that were stimulated with LPS (figure 3A), but not with Pam3CSK4 (figure 3B), exhibited a significant reduction of serum TNF and IL-6 levels. However, this modulation of TNF and IL-6 was not observed in GA-treated TRIF-deficient mice following LPS injection (figure 3B). As MyD88-deficient mice are unresponsive to Pam3CSK4 or LPS,12 these mice were evaluated by stimulation with Poly(I:C). GA treatment reduced proinflammatory cytokine serum levels in MyD88-deficient mice in response to Poly(I:C) (figure 3C). Thus, these in vivo results further establish that GA inhibits TRIF-dependent, but not MyD88-dependent, signaling.

We then addressed whether TRIF deficiency might alter the clinical effect of GA treatment in EAE. TRIF-deficient, but not MyD88-deficient, mice are susceptible to EAE.5-7 Clinical severity of EAE in untreated WT and TRIF-deficient mice was similar (figure 3D). However, GA suppressed EAE development in WT mice but not in TRIF-deficient mice. Specifically, the incidence, day of onset, and EAE severity were not altered by GA treatment of TRIF-deficient mice. These in vivo results further confirmed the importance of modulating TRIF-dependent signaling in the clinical activity of GA.

GA treatment inhibits IFN-β production by targeting regulatory elements of the IFN-β promoter. IFN-β is a key cytokine induced via the TRIF-dependent pathway that contributes to both innate and adaptive immune responses.4,6 Therefore, we analyzed expression of IFN-β in monocytes treated with GA. As anticipated, we observed a reduction of IFN-β secretion by GA-treated WT monocytes (figure 4A). Serum IFN-β was also decreased in GA-treated mice that were subsequently challenged with LPS (figure 4B).

Through binding to IFNAR and activating the signal transducers and activators of transcription (STAT) signaling pathway, IFN-β acts in a feedback manner to amplify monocyte responses to innate stimulation.13 Previously, we reported that STAT1 phosphorylation was impaired in M2 monocytes.3 Thus, we addressed the importance of IFNAR1 in GA treatment. As shown in figure 4C, IFNAR1 deficiency prevented GA-mediated modulation of IL-6 and TNF secretion. Together, these results suggest that reduction of type I IFN signaling may be a primary effect of GA action on monocytes.

Type I IFN expression requires the coordinated activation of the transcription factors NF-κB, IRF3, and IFNAR1.14 We evaluated the influence of GA treatment on activation of these signaling molecules and their regulators. As described above, GA treatment did not influence NF-κB activation. In contrast, we observed that GA suppressed nuclear translocation of IRF3 in monocytes following stimulation with either LPS or Poly(I:C) (figure 4D). Activation and nuclear translocation of IRF3 are dependent upon phosphorylation by c-Jun N-terminal kinase 1 (JNK1).15 In this regard, phosphorylation of JNK1 was reduced in GA-treated monocytes in response to LPS stimulation (figure 4E). Similarly, decreased DNA binding of ATF-2 was observed following incubation with GA (figure 4F and figure e-3), which corresponded to inhibition of p38 MAPK phosphorylation (figure 4G), the major upstream regulator of ATF-2.16 Taken together, these results demonstrate that GA treatment inhibited IFN-β production by interfering with activation of transcription factors participating in formation of the IFN-β enhanceosome.

**DISCUSSION** Cellular immune responses that result from GA treatment of MS and EAE have been well-described.17,18 In 2007, it was shown that it is the influence of GA on APCs that is responsible for the amplification of Th2 and Treg cells, a study that provided the first demonstration that M2 monocytes could reverse CNS autoimmune disease and promote T-cell immune modulation in vivo.3 At that time the long-held belief that GA-specific T cells required cross-reactivity with myelin self-antigen for therapeutic benefit was disproved. It was also observed that GA treatment was associated with inhibition of STAT1 in monocytes, indicating that GA affected at least one proinflammatory signaling pathway within these cells.3 TRIF and MyD88 represent 2 major innate pathways involved in the induction of several proinflammatory responses, including cytokine secretion. Our results...
demonstrate that GA inhibits the TRIF-dependent pathway, resulting in a reduction in activity of downstream signaling molecules that lead to production of IFN-β (figure 5). This immune modulation occurred independently of MyD88 and did not influence NF-κB activation. These findings provide a key anti-inflammatory mechanism connecting innate and adaptive immune modulation in GA therapy.

Engagement of some cell surface receptors has been associated with anti-inflammatory polarization of myeloid cells. Vasoactive intestinal peptide, which, similar to GA, also exerts anti-inflammatory effects,\(^{19}\)

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Figure 3 TRIF deficiency abrogates the immunomodulatory effects of glatiramer acetate treatment on cytokines and experimental autoimmune encephalomyelitis

Wild-type (WT) and Toll-IL-1 receptor domain-containing adaptor inducing interferon-β (TRIF)-deficient mice treated with glatiramer acetate (GA) or vehicle (n = 5 mice/group) were injected IP with (A) lipopolysaccharide (LPS) (100 ng/kg) or (B) Pam3CSK4 (1.00 ng/kg). (C) Myeloid differentiation primary response gene 88 (MyD88)-deficient mice were injected IP with Poly(I:C) (1.0 μg/kg). Serum levels of tumor necrosis factor (TNF) and interleukin (IL)-6 were quantitated by ELISA 5 hours after injection. Results are presented as the mean ± SEM (n = 3) of 2 experiments that provided similar results; *p < 0.05, **p < 0.01, ***p < 0.001 as determined by Student t test. (D) On day 0, C57BL/6J WT (left) or TRIF-deficient mice (right) mice were immunized with MOG peptide (p35-55; 50 μg). GA (250 μg) was administered once (SC in incomplete Freund’s adjuvant [IFA]) on the same day as immunization (day 0). Control mice received a single SC injection of IFA. For all experimental autoimmune encephalomyelitis experiments, mean disease score ± SEM is shown. *p < 0.05 as determined by Mann-Whitney U test. Results shown are representative of 3 independent experiments.
**Figure 4** Glatiramer acetate treatment negatively regulates IFN-β production by targeting components of the IFN-β enhanceosome

(A) Wild-type (WT) monocytes differentiated in the presence or absence of glatiramer acetate (GA) were stimulated with lipopolysaccharide (LPS) (100 ng/mL) for 24 hours. Interferon (IFN)-β secretion was quantitated in cell culture supernatants by ELISA. Results are representative of 2 experiments. Data are presented as mean ± SEM; *p < 0.05, **p < 0.01 as determined by Student t test. (B) WT mice (n = 3/group) treated with GA or vehicle were injected IP with LPS (100 ng/kg). Serum levels of IFN-β were quantitated by ELISA, 5 hours following injection. Data are representative of 2 separate experiments. (C) Interferon-α/β receptor subunit-1 (IFNAR1)-deficient monocytes differentiated in the presence or absence of GA were stimulated with LPS at the indicated dose for 24 hours. Tumor necrosis factor (TNF) and interleukin-6 secretion was quantitated in cell supernatants by ELISA. Results are representative of 3 independent experiments (n = 5/group). (D) Monocytes generated in the presence or absence of GA were stimulated with LPS (100 ng/mL) or Poly(I:C) (10 μg/mL) for the indicated duration. IRF3 binding activity in nuclear extracts was measured with TransAM IRF3. (E) Monocytes generated in the presence or absence of GA were stimulated with LPS (100 ng/mL) for the indicated duration. Cell lysate proteins were separated by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and membranes were probed for phosphorylated SAPK/JNK (Thr183/Tyr185) and total GAPDH. Data are representative of 3 separate experiments. (F) Monocytes differentiated in the presence or absence of GA and stimulated with 100 ng/mL LPS for the indicated duration. DNA binding of ATF-2 was quantitated from nuclear extracts. (G) Monocytes generated in the presence or absence of GA were stimulated with LPS (100 ng/mL) for the indicated duration, and whole-cell lysates were subjected to SDS-PAGE and Western blot analysis for phosphorylated p38 MAPK (Thr180/Tyr182) and total STAT1. Data are representative of 3 separate experiments.
triggers the second messengers PI3K and cAMP following receptor engagement. We established that PI3K, but not cAMP, participates in M2 polarization by GA, and like a previous study of human monocytes, further supports a central role for PI3K/Akt in regulating inflammatory responses. These results also suggest that GA, or its peptide degradation products, might engage protein receptors. GA exhibits promiscuous binding to allelic surface MHC class II molecules. Although the cytoplasmic domain of MHC class II molecules can participate in intracellular signaling, MHC class II is neither required for GA-dependent anti-inflammatory monocyte functions nor necessary for targeting of GA to these cells in vivo. As data have indicated that GA can bind to CD11b (CD18), a cell surface integrin that participates in immune regulation, it was speculated that the interaction of GA with CD11b might contribute to GA-mediated modulation. However, we have observed that during in vivo GA treatment, GA or its byproducts accumulate equally well in CD11b-deficient monocytes as CD11b-bearing (WT) monocytes and that CD11b expression is not required for GA-mediated M2 differentiation (figure e-4). These findings highlight the challenges in identifying a specific receptor that binds GA and leads to activation of second messengers. Indeed, GA, a random linear arrangement of glutamate, lysine, alanine, and tyrosine (i.e., glat in glatiramer) that exhibits extensive sequence variation, may interact with many macromolecules.

Other random basic synthetic copolymers have also been considered for development as potential therapies in MS. In this regard, one random copolymer containing phenylalanine, lysine, alanine, and tyrosine and another composed of valine, lysine,
alanine, and tryptophan also prevent EAE and cause immune modulation.\textsuperscript{25,26} Further, lysine—the basic amino acid common to these copolymers—is essential for therapeutic benefit, as "gat"-ramer acetate, lacking lysine, was ineffective in preclinical EAE studies. Thus, the observation that other synthetic random basic copolymers that, like GA, contain alanine, lysine, and an aromatic amino acid (i.e., phenylalanine, tryptophan, or tyrosine) also induce immune modulation raises the possibility that the signaling pathways modulated by GA identified in this report may be common to this class of copolymers.

While the observation that GA exerts anti-inflammatory activity via inhibition of TRIF and its downstream signaling molecules is novel, it is known that the glucocorticoid dexamethasone also promotes downstream signaling molecules is novel, it is known that the glucocorticoid dexamethasone also promotes M2 polarization.\textsuperscript{27} Although some data indicate that steroids interfere with NF-kB-dependent signaling, other results have demonstrated that steroid-mediated M2 polarization is primarily associated with inhibition of both IRF3 activation and type I IFN signaling.\textsuperscript{28} Resveratrol, an antioxidant found in red wine, inhibits TRIF-dependent activation,\textsuperscript{29} and, like GA, reduces clinical EAE and macrophage-mediated inflammatory activity.\textsuperscript{30} Thus, while the mechanism of action of GA is unique among MS therapeutics, suppression of TRIF-dependent signal transduction by agents that exhibit anti-inflammatory activity is not unprecedented.

Our finding that GA prevents IFN-\(\beta\) synthesis in M2 monocyes is consistent with the earlier demonstration that STAT1 phosphorylation is reduced upon activation in these cells.\textsuperscript{3} One may find these observations surprising, considering that IFN-\(\beta\) is approved for treatment of MS and has been protective in some EAE models.\textsuperscript{3,6} However, GA and IFN-\(\beta\) promote immune modulation via distinct mechanisms. For example, GA and IFN-\(\beta\) not only exhibit opposing activities on STAT1 phosphorylation, an obligate initial step in IFN-\(\beta\) signaling,\textsuperscript{31} but also demonstrate distinct effects on proinflammatory T-cell differentiation. Whereas GA induces both Th2 and Treg cells that can modulate Th1 and Th17 responses,\textsuperscript{3,18} respectively, some data indicate that IFN-\(\beta\) downregulates pathogenic Th1, but not Th17 responses.\textsuperscript{32} Separately, it was observed that deficiency in either IRF3 or JNK1 reduced susceptibility to EAE and impaired expansion of Th17 cells.\textsuperscript{9,33} Thus, our findings that GA treatment inhibited activation of JNK1 and nuclear translocation of IRF3 in monocytes and reduced production of IFN-\(\beta\) are consistent both with the knowledge that inhibition of TRIF-dependent gene products reduces Th17 differentiation and that GA inhibits Th17 polarization.\textsuperscript{9} Together with the observation that GA prevented nuclear translocation of ATF-2, which contributes to the activation of the IFN-\(\beta\) promoter, our data indicate that GA inhibits formation of the IFN-\(\beta\) enhancosome, a complex that is required for maximal IFN-\(\beta\) gene transcription.

Considerable effort has been devoted to the identification of therapeutics that may provide additive or synergistic benefit when combined for treatment of MS.\textsuperscript{34–37} Even though mixed results were obtained when GA was tested in combination with type I IFNs in preclinical studies,\textsuperscript{38,39} GA and intramuscular IFN-\(\beta\)-1a were tested separately and together in a phase III clinical trial in relapsing-remitting MS.\textsuperscript{35} GA alone reduced relapse rate more than IFN-\(\beta\)-1a, and was equivalent to the combination of GA and IFN-\(\beta\). While tempting to translate our mechanistic findings in relation to those clinical results, one should exercise caution; our study has focused primarily on identification of the mechanisms employed by GA in M2 monocyte polarization. Further, while results of that phase III MS trial did not identify clinical benefit from the combination of GA and intramuscular IFN-\(\beta\)-1a, it did not provide clear evidence of antagonism either. Evidence of synergy or antagonism may be dependent upon dosing. For example, when cholesterol-lowering statins were administered in combination with low-dose (intramuscular) or high-dose (subcutaneous) IFN-\(\beta\) in MS clinical trials, antagonism was observed and appeared more evident with high-dose IFN-\(\beta\); this effect may relate to the opposing actions of these medications on STAT1 activation.\textsuperscript{36} In retrospect, our observation here that GA treatment inhibited expression of IFN-\(\beta\) in monocytes would not have provided mechanistic support for testing the combination of GA and IFN-\(\beta\) in MS. Recognizing that GA regulates TRIF-dependent IFN-\(\beta\) production in monocytes may provide insight regarding its therapeutic effects in MS and for identification of surrogate markers of its efficacy.

**AUTHOR CONTRIBUTIONS**

N.M., T.P., and S.S.Z. designed research, analyzed data, and wrote the article. J.C.P., C.M.S., M.S.W., and P.H.L. analyzed data, gave conceptual advice, and discussed the results. N.M. and T.P. performed the experiments. U.S.-T. conducted the EAE experiments. S.S.Z. supervised the study. All authors read, commented, and approved the final manuscript.

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**DISCLOSURE**

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CORRECTION

Glatiramer acetate treatment negatively regulates type I interferon signaling

In the article “Glatiramer acetate treatment negatively regulates type I interferon signaling” by N. Molnarfi et al. (Neuroimmunol Neuroinflamm 2015;2:e179. DOI: 10.1212/NXI.0000000000000179), originally published on November 9, 2015, there is an error in the description of figure 2B. This figure shows \( \text{I} \kappa \beta \text{a} \) degradation, not \( \text{I} \kappa \beta \text{a} \) phosphorylation as originally described. In Methods (“Western blot”), the second sentence should read “Nitrocellulose membranes were probed (4°C, overnight) with antibodies against phospho-Akt1 (Ser473), phospho-JNK (Thr183/Tyr185), phospho-p38 MAP kinase (Thr180/Tyr182), \( \text{I} \kappa \beta \text{a} \), glyceraldehyde-3-phosphate dehydrogenase (GAPDH), total STAT1, and total Akt1/2/3 (Cell Signaling Technology, Danvers, MA).” In figure 2, the label above panel B should read just \( \text{I} \kappa \beta \text{a} \) and the corresponding sentence in the legend should read “…and membranes were probed for \( \text{I} \kappa \beta \text{a} \).” In the second paragraph under Results (“GA inhibits TRIF-dependent, but not MyD88-dependent, signaling”), the 13th sentence should read “As shown in figure 2B, GA exposure did not impair degradation of the inhibitor protein \( \text{I} \kappa \beta \text{a} \) following LPS treatment.” The authors regret this error.