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Magnesium isoglycyrrhizinate positively affects concanavalin A-induced liver damage by regulating macrophage polarization

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ABSTRACT

The deficient functional polarization of macrophages is implicated in the disease progression of autoimmune hepatitis (AIH). This study aims to evaluate the impact of Magnesium isoglycyrrhizinate (MgIG) on concanavalin A (Con A)-induced hepatitis in a mouse model, thereby clarifying the molecular mechanisms with which it is associated. MgIG was periodically administered to C57BL/6 mice before one intravenous injection of Con A (20 mg/kg). The MgIG treatment demonstrated a protective function in mice for Con A-induced AIH, the expression of proinflammatory cytokines, and the serum levels of alanine aminotransferase and aspartate aminotransferase. In addition, the MgIG pre-treatment had a significant effect on the number of F4/80⁺ cells entering the liver. MgIG efficiently facilitated macrophage polarization toward an M2 phenotype. The results indicate that a relationship may exist between the protective impacts of MgIG with respect to Con A-induced liver injury and the capability of the hepatoprotective agent to regulate macrophage polarization.

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1. Introduction

As a progressive inflammatory liver disease involving the diminishment of self-tolerance, ultimately resulting in the emergence of autoantibodies paired with lymphoplasmacytic inflammatory infiltrates in the portal tracts, autoimmune hepatitis (AIH) is also characterized by the appearance of the fragmentary necrosis of periportal hepatocytes (McFarlane, 1999). The underlying causal mechanisms that give rise to AIH are not yet understood. Although it is now known that autoreactivity to certain hepatocyte components is crucial in the course of the disease's development, the molecular mechanisms that contribute to the impairment of immune tolerance in AIH are not understood (Heneghan, Yeoman, Verma, Smith, & Longhi, 2013). The disease is characterized by the difficulties healthcare practitioners have when attempting to control its progression. This is especially

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the case among patients who are refractory to (or intolerant of) conventional therapy, which involves prednisone and azathioprine (Rubin & Te, 2016). Hence, the exploration of novel therapeutic agents and treatment modalities is a priority in the literature.

Concanavalin A (Con A)-induced liver injury refers to a frequently used mouse model that can identify immune cell-mediated acute hepatitis and is comparable to the pathology of human AIH in several ways (Wahl, Wegenka, Leithauser, Schirmbeck, & Reimann, 2009). One of the crucial contributors to the Con A-induced breakdown of liver immune homeostasis is the macrophage, which is implicated in the production of pro-inflammatory cytokines (Assis et al., 2014). Macrophages are defined as highly plastic cells that change in view of the surrounding tissue microenvironment (Peng & Tian, 2016; Tacke & Zimmermann, 2014; Wynn & Vannella, 2016). These cells engage in various functions in the body, such as immune responses, tissue repair, and homeostasis (Wynn, Chawla, & Pollard, 2013). Usually, resident macrophages in the liver, classified as Kupffer cells, were considered as being neither diverse nor dynamic. Lloyd et al. showed F4/80+, CD11b or CD68 could be used to identify Kupffer cells (Beyazit et al., 2015). Meanwhile, nitric oxide may be a potential mediator of hepatic inflammation and fibrogenesis in AIH. The serum nitric oxide metabolites like inducible nitric oxide synthase (iNOS) and nitric oxide activity can reflect oxidant stress in the liver and detect the inflammation and disease progression (Lloyd, Phillips, Cooper, & Dunbar, 2008; Yang, Zhang, Han, Jin, & Yang, 2017). In addition, various surface markers and cytokines are expressed and secreted, respectively, by the classical activation (M1) and the alternative activation phenotypes (M2), both of which have variable impacts in the context of immunoregulation (Ivashkiv, 2013; Ka, Daumas, Textoris, & Mege, 2014; Ying, Cheruku, Bazer, Safe, & Zhou, 2013). In recent years, the literature has made it clear that macrophage imbalances are a fundamental contributor to the pathogenesis of autoimmune diseases (Alleva, Pavlovich, Grant, Kaser, & Beller, 2000; Cosin-Roger et al., 2013; Espinoza-Jimenez, Peon, & Terrazas, 2012; Parsa et al., 2012; Zigmond et al., 2014).

Magnesium isoglycyrrhizinate (MgIG) is a magnesium salt of the 18- α -glycyrrhizic acid stereoisomer and is classified as a new molecular compound identified in the roots of the *Glycyrrhiza glabra* (licorice) tree (Wu et al., 2018). The compound is notable due to its anti-inflammatory and antioxidant effects and has been used to treat liver conditions (Ming & Yin, 2013; van Rossum, Vulto, de Man, Brouwer, & Schalm, 1998). Nevertheless, the impact of MgIG on Con A-induced AIH is not well understood. Therefore, this study investigates the preventive value of the compound with respect to liver injury using a mouse model of hepatitis (induced via intravenous Con A injection) and its underlying molecular mechanisms.

2. Materials and methods

2.1. Chemicals and reagents

The MgIG used in this study was procured from Jiangsu Chia-Tai Tianqing Pharmaceutical Co., Ltd. (Nanjing, Jiangsu, China). The Con A and lipopolysaccharide (LPS) used were procured from Sigma-Aldrich (St. Louis, MO, United States). Unless otherwise stated, the remaining chemicals used were also purchased from Sigma-Aldrich (St. Louis, MO, United States).

2.2. Animals and treatment

The Experimental Animal Centre at the Tianjin Medical University (TMU, Tianjin, China) was used to purchase male C57BL/6 mice aged 6–8 weeks. All of the mice were kept at $25 \pm 2^\circ\text{C}$ and $50 \pm 10\%$ relative humidity. This was paired with a 12-hour light/dark cycle. The animals were fed using a standard pellet diet with drinking water. The research proceedings were authorized by the ethical committee at the TMU and all of the procedures followed the Care and Use of Laboratory Animals regulations.

The mice were separated into the following categories: control mice ($n = 7$), mice that received Con A treatment ($n = 7$), mice that received both Con A treatment and 50 mg/kg MgIG ($n = 7$), and mice that received both Con A treatment and 100 mg/kg MgIG ($n = 7$). To prepare the Con A-induced liver injury mouse model, the mice received 20 mg/kg Con A via intravenous injection. The MgIG-treated mice were administered 50 or 100 mg/kg MGL (diluted using saline) 2 h before Con A challenging. Before sacrifice, the mice in the control group received an equal volume of saline. All of the mice were sacrificed 24 h after Con A injection for blood and liver tissue analysis.

2.3. Cell culture and treatment

Murine macrophage-like RAW264.7 cells were acquired from Sigma Chemical, Co., Ltd. (St. Louis, MO, United States) and were pre-cultured using the DMEM medium (Gibco BRL, Grand Island, NY, United States). The DMEM was combined with 10% fetal bovine serum. The cells were then randomly allocated into one of the following groups: a control group (comprised of cells that received phosphate-buffered saline [PBS] treatment), an LPS group (treated with 100 ng/mL LPS), an MgIG intervention + LPS group (treated with 50 μM MgIG + 100 ng/mL LPS), and an MgIG control group. For the MgIG intervention + LPS group, the cells were exposed to 100 ng/mL LPS and MgIG for 24 more hours. Following treatment, all of the cells underwent washing and harvesting (the latter by centrifugation), thereby facilitating RNA harvesting.

2.4. Evaluation of liver histopathology

Each liver sample was fixed in 4% paraformaldehyde, embedded in paraffin, and then cut into 5- μm sections. The sections were then stained using hematoxylin and eosin (H&E). Every sample underwent histological assessment by drawing on a pair of professional pathologists, which involved the use of the Knodell scoring system (KSS) to assess the extent of liver damage (Knodell et al., 1981). The KSS assesses liver damage based on the following criteria: (1) absence of inflammation or spotty necrosis, (2) fragmentary or confluent necrosis (less than 10%), (3) confluent necrosis (10–50%), and (4) significant confluent necrosis (more than 50%) accompanied by or not accompanied by bridging necrosis.

2.5. Biochemical analysis

After the mice from each group were sacrificed at the end of the experiment, blood was extracted from the vena cava. In turn, using the Automatic Chemical Analyzer 7600-100 (Hitachi, Ltd., Tokyo, Japan), the alanine aminotransferase (ALT) and aspartate aminotransferase (AST) serum levels were examined.

Table 1. Mouse primer sequences used for qRT-PCR.

Target gene	Forward primers (5'-3')	Reverse primers (5'-3')
IFN- γ	ATGGCTGTTCTGGCTTACT	AATGACGCTTATGTTGTGCTG
TNF- α	GACGTGGAAGTGGCAGAAGA	ACTGATGAGAGGGAGGCCAT
IL-1 β	GTGCAAGTGTCTGAAGCAGC	CAAAGGTTTGAAGCAGCCC
IL-6	GGAGTCACAGAAGGAGTGGC	CGCACTAGGTTTGCCGAGTA
iNOS	GAGCCACAGTCTCTTTGCTA	TGTCACCACCAGCAGTAGTTG
Arg-1	CGTTGTATGATGCACAGCCG	CCCCACCAGTGATCTTGAC
GAPDH	TCTCTGCGACTTCAACA	TGTAGCCGTATTCATTGTCA

2.6. RNA preparation and analysis

Frozen liver tissues were cut to small sections and washed with PBS buffers. 5–10 mg of tissue was added to 700 μ l lysis buffer and homogenised with a TissueLyser II (Qiagen). Then Trizol reagent (Invitrogen, Carlsbad, CA, United States) was used to extract total RNA from frozen liver tissue and cultured cells according to the manufacturer's guidelines. The RNA was reverse transcribed with random hexamers and avian myeloblastosis virus reverse transcriptase using a commercial kit (Perfect Real Time, SYBR^R PrimeScriPTM TaKaRa, Japan). In addition, the gene expression analysis involved the use of qRT-PCR with SYBR Premix EX TaqTM II, facilitated by the ABI PRISM 7900 sequence detector (Applied Biosystems, Foster City, CA, United States). Sangon Biotech Co., Ltd. (Shanghai, China) was the source of the primer sequences for the target genes (Table 1). The standard curve, paired with $2^{-\Delta\Delta C_t}$ methods, was used to identify quantification.

2.7. Immunofluorescence analysis

Deparaffinization of the paraffin-embedded liver sections was conducted using xylene. Furthermore, the sections were rehydrated and incubated using a blocking solution (5% bovine serum albumin in PBS). In turn, fluorescein isothiocyanate conjugated F4/80 antibodies were incubated (BD Biosciences, San Diego, CA, United States) overnight at 4°C. After washing the PBS three times, coverslips were applied to the sections and observations were made using a confocal microscope (Olympus Inc., Center Valley, PA, United States).

2.8. Statistical analysis

The results are expressed as mean \pm SD. The data were examined using Student's *t*-test and one-way analysis of variance. To facilitate the statistical analysis, GraphPad Prism 5.0 was used (GraphPad Software, San Diego, CA, United States). Statistical significance was determined at the 5% level.

3. Results

3.1. MgIG preconditioning positively affects con A-induced liver damage

MgIG was administered intraperitoneally (depending on the mouse group, 50 or 100 mg/kg), as outlined in the previous section. This facilitated the study's attempt to identify the degree of MgIG's protective role in Con A-induced hepatitis (Figure 1(A)). When comparatively examined against the saline control groups, the serum ALT and AST levels evidently increased as a consequence of the Con A treatment. Furthermore, in a dose-

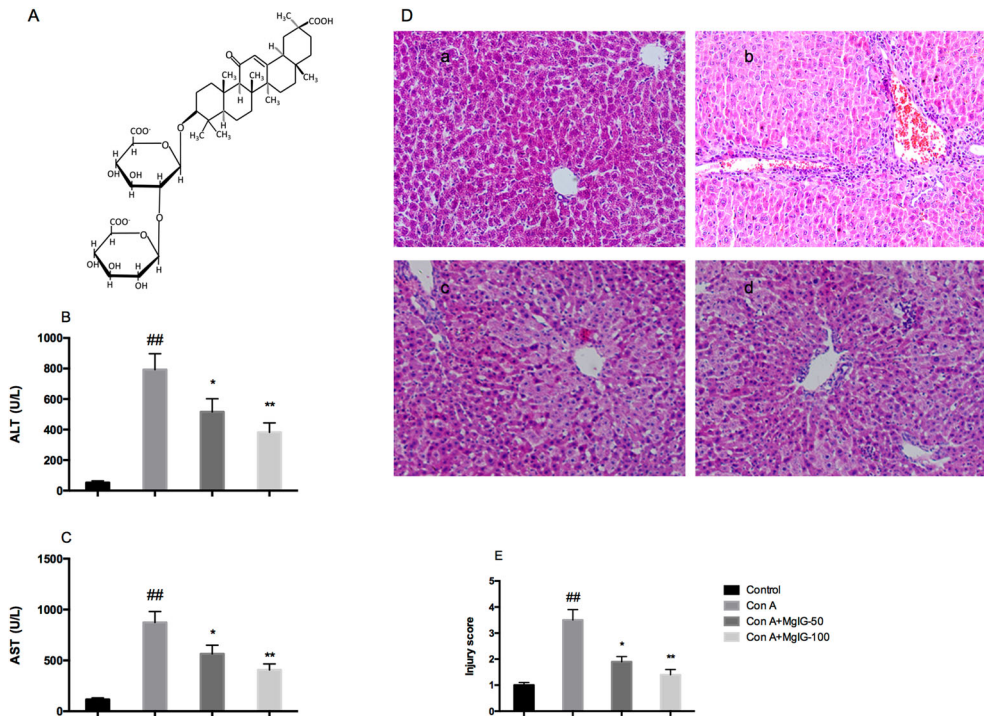


Figure 1. Impacts of pre-treatment using MgIG (intraperitoneal administration of 50 or 100 mg/kg) for Con A-induced liver damage. (A) MgIG's chemical structure; (B) Serum ALT levels; (C) Serum AST levels; (D) H&E staining of liver tissue; (E) Liver injury score. The data are expressed as mean \pm SD ($n = 7$, $##P < 0.01$ versus control; $*P < 0.05$, $**P < 0.01$ versus Con A). a: Control; b: Con A; c: Con A + MgIG-50; d: Con A + MgIG-100.

dependent way, MgIG pre-treatment reduced ALT and AST activity ($P < 0.05$), thereby indicating the protective role of MgIG in the context of Con A-induced hepatitis (Figures 1(B and C)). As indicated in Figure 1(D), morphological examination provided additional evidence suggesting the protective impact of MgIG. Administering Con A induced significant liver damage (in particular, hepatocyte necrosis accompanied by sinusoidal hyperemia associated with hemorrhage). However, after MgIG was administered, hepatocyte necrotic injury improved considerably and the number of infiltrating inflammatory cells in the necrotic area decreased. In addition, when comparatively examining the MgIG pre-treated mice to the Con A-treated mice (Figure 1(E)), the liver injury score was favorable for the former.

3.2. MgIG preconditioning inhibits the secretion of proinflammatory cytokines for con A-treated mice

Given that the secretion of large quantities of inflammatory cytokines performs an important function in the disease progression of Con A-induced AIH, one of the aims of this study is to determine the extent to which MgIG pre-treatment affects the expression of inflammatory cytokines in mice. As outlined in Figure 2, the mRNA levels of pro-inflammatory cytokines (IL-1 β , IL-6, IFN- γ , TNF- α , and inducible nitric oxide synthase [iNOS])

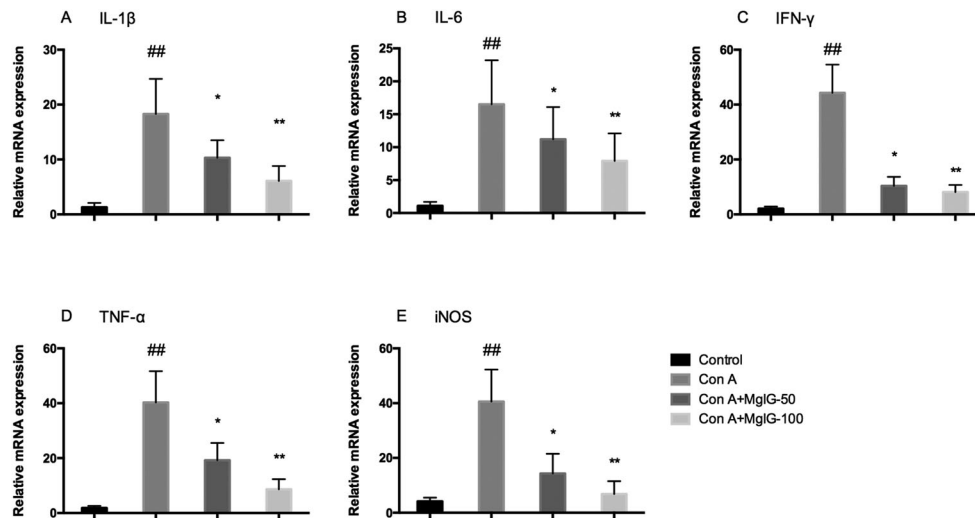


Figure 2. MglG pre-treatment of mice decreases the release of proinflammatory cytokines in the liver tissue following induction of Con A ($n = 7$, ## $P < 0.01$ versus control; * $P < 0.05$, ** $P < 0.01$ versus Con A).

in the liver were considerably greater in response to Con A exposure than those of the control mice ($P < 0.01$). Nevertheless, MgIG administration rehabilitated this increasing trend among the Con A-induced mice in a dose-dependent manner ($P < 0.05$).

3.3. Preconditioning with MgIG-Impaired macrophage infiltration of the liver for con A-treated mice

Immunofluorescence assays were used to illuminate the impact of MgIG on the infiltration of F4/80⁺ macrophages in the livers of the Con A-induced mice. As indicated in [Figure 3](#), the F4/80⁺ macrophage numbers increased considerably following Con A challenging ($P < 0.01$). However, it was possible to move them back to basal levels after MgIG treatment ($P < 0.05$ or $P < 0.01$).

3.4. MgIG regulates M1 and M2 inflammatory macrophage polarization in LPS-Stimulated RAW264.7 cells

Finally, to examine the impact of MgIG on the polarization of macrophages within the RAW264.7 cells, the degree to which MgIG could convert macrophages into an anti-inflammatory M2 phenotype was assessed. iNOS, the M1 phenotype marker, increased considerably due to LPS stimulation in the RAW264.7 cells and LPS-induced iNOS generation was inhibited following MgIG treatment. In contrast, the M2 marker arginase-1 (Arg-1) expression in the LPS-induced RAW264.7 cells was markedly upregulated by MgIG ([Figure 4](#)). Thus, MgIG facilitated M1 phenotype switching to the M2 macrophage.

4. Discussion

Licorice has a long history as a medicinal agent. Appearing in an early *materia medica* manuscript, “Shennong’s Classic Materia Medica,” it has since been applied across China in many contexts as a result of its characteristics (Yang et al., 2016). MgIG, a magnesium salt of the 18- α -glycyrrhizic acid stereoisomer and classified as a new molecular compound identified in the roots of the *Glycyrrhiza glabra* (licorice) tree (Lu, Xu, et al., 2017), is associated with anti-inflammatory and antioxidant properties and hepatic protection and liver function enhancement (He et al., 2010; Sun et al., 2007). For the inhibition of ethanol-induced lipid peroxidation, MgIG has also been used as a pharmaceutical for liver protection (Xie et al., 2015). Other liver-protective properties stem from how the drug prevents ischemia/reperfusion-induced liver injury (Huang, Qin, & Lu, 2014) and free fatty acid-induced hepatic lipotoxicity (Cheng, Zhang, Shang, & Zhang, 2009). Nevertheless, with respect to AIH, the literature on MgIG’s anti-inflammatory and immunoregulatory impacts is not extensive. Con A-induced liver damage is a common mouse model that has been used to identify immune cell-mediated acute hepatitis and is similar to the pathology of human AIH in various respects (Wahl et al., 2009). In this study, the mice suffering from Con A-induced hepatitis had elevated ALT and AST serum levels accompanied by hepatic damage. Nevertheless, following MgIG pre-treatment, the inflammatory pathway was inhibited, thus reducing the pathological damage of liver cells and enhancing liver cell function.

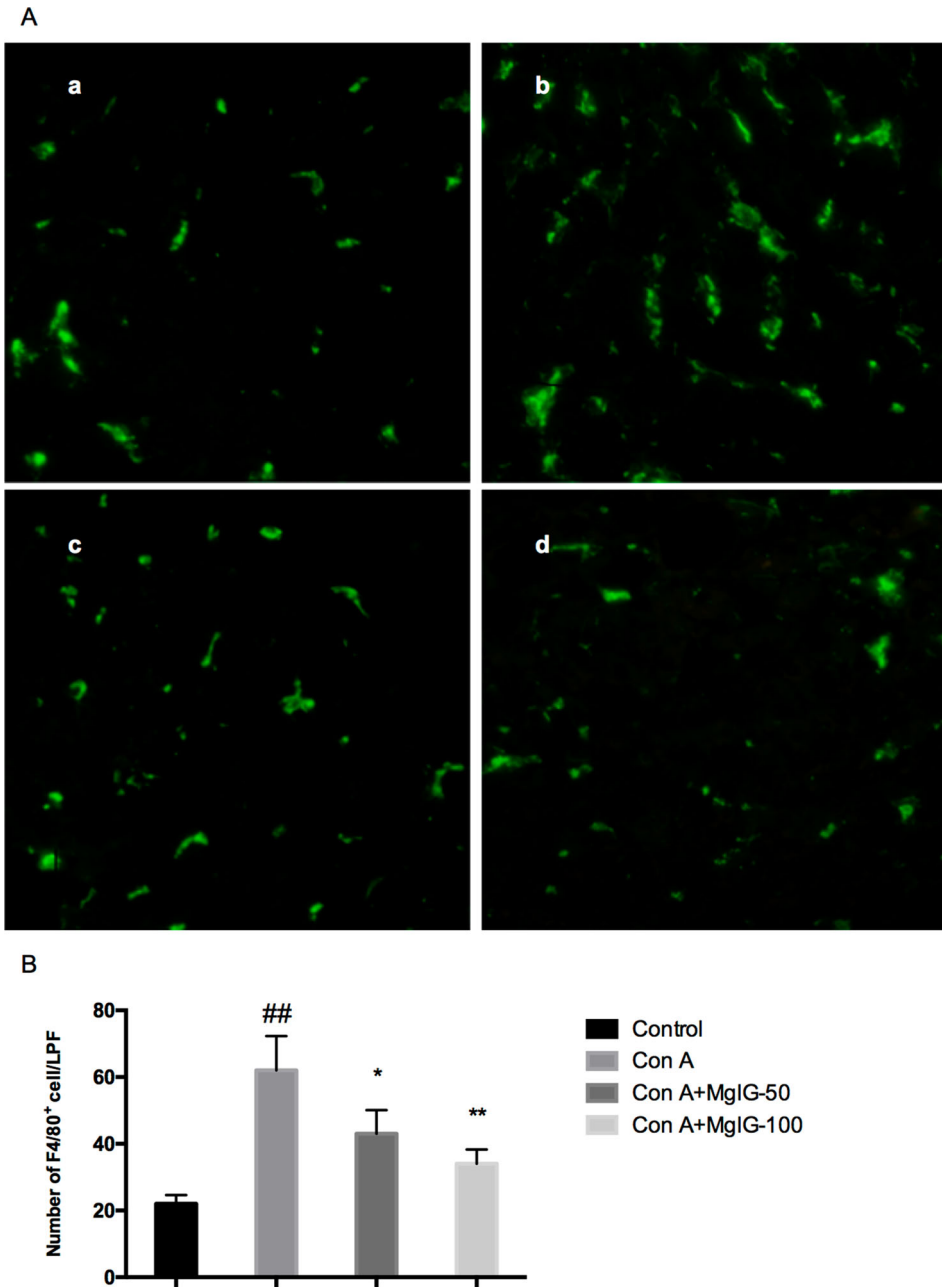


Figure 3. Impact of MglG on F4/80⁺ macrophage infiltration in the liver samples of Con A-induced mice. (A) Immunofluorescence assays were used to identify F4/80⁺ macrophage infiltration; (B) Quantification of F4/80⁺ macrophages in the mice livers took place for each group. The data are expressed as mean \pm SD ($n = 7$, $##P < 0.01$ versus control; $*P < 0.05$, $**P < 0.01$ versus Con A). a: Control; b: Con A; c: Con A + MglG-50; d: Con A + MglG-100.

This study also seeks to identify the underlying molecular mechanism through which MglG gains its protective properties for mice suffering from Con A-induced hepatitis. Evidently, the breakdown of immune tolerance is important in AIH pathogenesis and

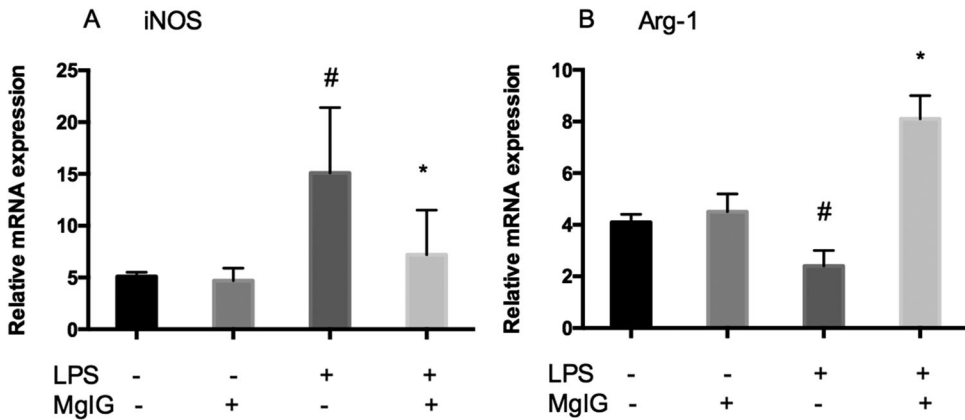


Figure 4. MglG underpins the regulation of LPS-induced M1 and M2 inflammatory macrophage polarization. (A) iNOS mRNA expression; (B) Arg-1 mRNA expression. The data are expressed as mean \pm SD ($n = 6$, $\#P < 0.05$ versus control alone; $*P < 0.05$ versus LPS alone).

macrophages perform critical functions in innate immune defense and homeostasis (Lavin et al., 2014). The literature is replete with publications showing how macrophages are similarly implicated in the Con A-induced hepatitis model (Higashimoto et al., 2013; Orsolich, Jazvinscak Jembrek, & Terzic, 2017). Hence, the inhibition of macrophage infiltration constitutes the therapeutic target for Con A-induced hepatitis. In this study, the number of infiltrating F4/80⁺ cells in the livers of the Con A-induced hepatitis mice increased considerably. However, this could be inhibited by MglG pre-treatment. Macrophages, which are key factors in promoting innate immunity, are not a cell population characterized by homogeneity and when exposed to certain signals, these cells can undergo classical M1 or alternative M2 activation (Lu, Chen, Ren, Yang, & Zhao, 2017; Udalova, Mantovani, & Feldmann, 2016). The cells are defined by how they express cell surface markers, how they secrete cytokines and chemokines, and their transcription and epigenetic pathways (Zhou et al., 2014). The findings cumulatively suggest that M1 macrophage activation plays an important role in liver inflammation and fibrosis. In particular, M1 macrophages are responsible for secreting a range of pro-inflammatory mediators, such as TGF- β 1, TNF- α , IL-1 β , and IL-6, which result in the inflammatory cascade (Beljaars et al., 2014; Sica, Invernizzi, & Mantovani, 2014). Contrastingly, several studies have demonstrated that M2 macrophages, typified by conventional M2 markers, such as Arg-1, YM1, FIZZ1, and MGL2 (Sica et al., 2008), can inhibit hepatic inflammation and are also important in solving fibrosis (Labonte, Sung, Jennelle, Dandekar, & Hahn, 2017; Tosello-Trampont et al., 2016). The phenotype alteration informs the part they play in liver immunoregulation, the implication of which is that M1/M2 macrophage polarization manipulation may yield effective therapeutic interventions (Tacke, 2017; Xu et al., 2015). This study indicates that MglG treatment constitutes an alternative to M2 macrophage polarization, as it induces Arg-1 expression and lowers the production of M1 inflammatory markers. In view of this, this study proposes that the anti-inflammatory impact of MglG stems from altering macrophage phenotype.

Taken together, this study's findings show that MglG constitutes a protective agent against Con A-induced hepatitis. Furthermore, it is clear that its positive action in this

respect can be accounted, as it facilitates the regulation of macrophage polarization. These findings may allow for the development of a new pharmacological treatment for AIH in future research.

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Disclosure statement

No potential conflict of interest was reported by the authors.

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