Intravenous Immunoglobulin with Enhanced Polyspecificity Improves Survival in Experimental Sepsis and Aseptic Systemic Inflammatory Response Syndromes

Running Head: Modified IVIg improve survival in sepsis

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Abstract

Sepsis is a major cause for death worldwide. Numerous interventional trials with agents neutralizing single pro-inflammatory mediators have failed to improve survival in sepsis and aseptic systemic inflammatory response syndromes. This failure could well be explained by the widespread gene expression dysregulation known as "genomic storm" in these patients. A multifunctional polyspecific therapeutic agent might be needed to thwart the effects of this "storm". Licensed pooled intravenous immunoglobulin preparations seemed to be a promising candidate but they have also failed in their present form to prevent sepsis-related death. We report here the protective effect of a single dose of intravenous immunoglobulin preparations with additionally enhanced polyspecificity in three models of sepsis and aseptic systemic inflammation. The modification of the pooled immunoglobulin G molecules by exposure to ferrous ions resulted in their newly acquired ability to bind some pro-inflammatory molecules, complement components and endogenous "danger" signals. The improved survival in endotoxemia was associated with serum levels of pro-inflammatory cytokines, diminished complement consumption and normalization of the coagulation time. We suggest that intravenous immunoglobulin preparations with additionally enhanced polyspecificity have a clinical potential in sepsis and related systemic inflammatory syndromes.

Introduction

Sepsis remains a leading cause of death in intensive care units. It is the result of a severe and uncontrolled activation of inflammatory and coagulation pathways in response to infection, accompanied by a variable degree of immune paralysis (1-3). Despite adequate antibiotic therapy and the use of sophisticated life-supporting measures the prognosis of patients with this syndrome has only marginally improved in recent years. This frustrating lack of progress, especially when novel experimental treatments aimed to target individual mediators of inflammation were used, has been hard to explain so far (1, 2, 4, 5).

A recent study has found significant changes of expression patterns of more than 80% of human genes (referred to as "genomic storm") in trauma patients and bacterial LPS-injected volunteers with severe inflammatory response syndrome (SIRS) (6). This "genomic storm" could well explain the discouraging results from the efforts to treat severe generalized inflammatory syndromes by neutralizing a single pro-inflammatory mediator. Targeting only one or very few components in a system-wide network disturbance, may not be successful in exerting control. A multifunctional therapeutic agent may be needed instead. Passive immunotherapy with pooled immunoglobulin preparations (administered as intravenous immunoglobulin, IVIg) is a logical choice as they contain a vast array of antibody specificities, some of which could well affect key products of the genomic storm. In addition to antibodies that neutralize pathogens and their virulence factors, IVIg have diverse immunomodulatory and anti-inflammatory activities (7). The latter are mediated through versatile interactions with receptors on immune cells, components of the complement system, cytokines, etc. The outcome is down-regulation of T-, B-lymphocyte activity and dendritic cell functions and modulation of the cytokine network (for

review see (7, 8)). The results from numerous clinical trials, using IVIg infusions as adjunctive therapy in sepsis patients have been, however, inconclusive (3, 9-11).

All commercially available IVIg preparations are generally believed to have identical biological and therapeutic properties. This may not be the case as the licensed therapeutic immunoglobulins, produced using a fractionation step at pH 4.0 have been shown to possess an increased potential to bind to self-antigens (12). Importantly, this increased reactivity to antigens correlates with different functional activity of immunoglobulin preparation (13, 14). Thus, the administration of the acid pH-treated, but not of the same unmodified preparation significantly decreased mortality in animals with endotoxemia (12, 13). Previous studies by our and other groups have proven that, in addition to low pH buffers, the exposure to a number of other substances (e.g. ferrous ions, heme, reactive oxygen species, etc.) also increases the antigenbinding polyspecificity of some IgG molecules (5, 15-17). IVIg modified by Fe(II) exposure acquired the ability to bind to the human pro-inflammatory cytokine IFN-γ and to improve survival in mice injected intravenously with 5.108 live E. coli or i.p with bacterial LPS (15, 18). Infusions of Fe-ions modified IVIg were also shown to have an anti-inflammatory activity in an experimental diabetes model (14). On the basis of these preliminary data we have hypothesized that the passive immunotherapy with pooled immunoglobulin preparations with additionally enhanced polyspecificity could neutralize some of the products of the genomic storm and thus should be beneficial in systemic inflammatory syndromes regardless of their primary insult. Three models of systemic inflammation in the presence or absence of infection were used to check this hypothesis: induced by LPS, by zymosan and by cecal ligation and puncture (CLP). While the infusion of native IVIg had no effect on survival, the administration of the same single dose of the Fe(II)-exposed IVIg improved significantly the survival of mice in all three models.

The studies of the mechanisms of beneficial action of the latter preparation revealed its ability to bind to pro-inflammatory molecules, complement components and extracellular histones.

Materials and Methods

Mice

Outbred female ICR mice (8-12 weeks old, 18-22 g) were purchased from the Breeding Farm of the Bulgarian Academy of Sciences and kept in a conventional animal facility. C57Bl/6 mice (8-12 weeks old, 18-22 g) were from Charles River Laboratories, Sulzfeld, Germany. The animals were housed under specific pathogen-free conditions, 6 per cage, with *ad libitum* access to food and water and maintained in a temperature-controlled environment. The animals were allowed to adapt to laboratory conditions for three days. The experimental protocols were approved by the Animal Care Commission of the Institute of Microbiology and the Thuringian State Office for Consumer Protection and Food Safety in accordance with National and European Regulations.

Blood (0.05 ml / mouse) was collected from the retro-orbital sinus by a Pasteur pipette after a local anesthesia with 0.5% tetracaine hydrochloride and let to clot at +4 $^{\circ}$ C. Sera samples were aliquoted and stored at -80 $^{\circ}$ C.

Experimental sepsis models

All survival experiments were performed using ICR mice in groups of twelve animals each. Endotoxemia was induced by the i.p. injection of 10 mg/kg *E. coli* LPS (B 055:B5, #L2880, Sigma-Aldrich St. Luis, MO). Cecal ligation and puncture (CLP) was performed as described in the literature (19). Briefly, the mice were anesthetized i.p. with ketamine (80 mg/kg) and xylazine (10 mg/kg), a 1.5 cm long laparotomic incision was made, the cecum was ligated at 2/3

of its length and punctured once by a sterile 21*G* needle. The puncture was confirmed by a delicate pressing and the abdomen was closed by suture. No antibiotics or fluids were administered. Multiple organ dysfunction was induced by the intraperitoneal injection of 500 mg/kg zymosan (Sigma-Aldrich) (20). The dose of the LPS or zymosan was adjusted to cause 80 to 100% mortality in the control animals, treated with PBS only.

Modified pooled therapeutic human immunoglobulins

The intravenous immunoglobulin preparations Endobulin S/D (Baxter, Deerfield, IL) and a special maltose- and albumin-free batch of Immunovenin-intact (BulBio Ltd, Sofia, Bulgaria, kindly provided by Dr Julia Nacheva), were used in the experiments. Both preparations are referred to as "native", as they are produced without using a fractionation step at acid pH (12). The pooled immunoglobulins were exposed to freshly prepared ferrous sulfate solution (1 mM final concentration) for 2 h at 4°C and then dialyzed against phosphate-buffered saline (PBS; pH 7.4) for at least 48 h as previously described (15). In a separate experiment the ferrous ionsmodified IVIg was dialyzed against 4 mM EDTA in PBS and then against PBS.

Treatment schedules

Single dose of the native, the modified IVIg preparations or PBS alone was injected intravenously (i.v.) 10 min before the administration of LPS, of zymosan or at the beginning of CLP procedure. In all experiments survival was observed daily for seven days.

In a separate experiment groups of animals were treated i.v. with 50 mg/kg of the native or Fe(II)-exposed IVIg at different time points – minutes before, one, or four hours post-LPS injection.

Determination of serum levels of cytokines

The levels of TNF-α, IFN-γ, IL-6, IL-10 and IL-22 in the sera obtained 4, 24 and 48 h after LPS injection were measured using commercial ELISA kits (PeproTech EC Ltd., London, UK).

The Mouse inflammation antibody array kit (RayBiotech Inc., Norcross, GA). was used to evaluate semi-quantitatively the levels of 40-inflammation-related molecules. Briefly, membranes spotted with 40 capture antibodies were blocked with the manufacturer's BSA buffer and incubated with the studied serum samples for 2 h at room temperature. After washing, a cocktail of biotinylated anti-cytokine antibodies was incubated with membranes for 2 h at room temperature. Membranes were then washed, incubated with HRP-conjugated streptavidin for 2 h, washed and exposed to the ECL peroxidase substrate for 2 minutes. Next, the membranes were drained and exposed for 20 seconds to a X-ray film (Kodak X-Omat AR film). The gray color intensity of the spots was quantified by densitometry and analyzed using the Image-Tool v2.0 for Windows (UTHSCSA, San Antonio, TX, USA) software package.

Measurement of plasma C3 levels

The C3 levels in plasma were measured at 4, 24 and 48 hour after LPS administration using the Mouse C3 ELISA kit (GenWay Biotech Inc., San Diego, CA).

Binding of modified IVIg to the C1q complement component and biological activity assays

Increasing concentrations of native and of ferrous ions exposure-modified IVIg were added to
the wells of a plate coated with human C1q (Calbiochem, Merck KGaA, Darmstadt, Germany).

Bound IgG was detected using anti-human IgG-HRP, (Fc-specific, from Southern Biotech) and OPD substrate.

The ability of modified IVIg to interfere with the binding of human C1q to its ligands was analyzed by ELISA. C-reactive protein (from Calbiochem) or human IgG_1 were coated to the plates at a concentration of 10 μ g/ml and the wells were blocked with 2% BSA. IVIg (at 10 mg/ml) was pre-exposed to ferrous ions (0; 0,5 and 1mM) and was mixed at 1,5 mg/ml in the wells of a second plate with increasing concentrations of C1q (7 serial dilutions starting from 10μ g/ml). The mixtures were transferred to the CRP- and human IgG_1 -coated plates. After incubation for 1 h at 37 °C and extensive washing, anti-C1q biotin-conjugated antibody (purified from goat anti-C1q antiserum from Quidel, San Diego, CA and biotinilated in house) was added in a 1:500 dilution. Finally streptavidin-HRP was added (from DAKO, diluted 1:1000) for 30 minutes. After washing TMB substrate was used, the reaction was stopped with 2M sulfuric acid and the optical density was read at 450 nm.

Coagulation and organ dysfunction tests

Coagulation time was measured as previously described (21). Animals were bled 4 hours after the injection of LPS and the plasma concentrations of bilirubin and creatinine were measured by commercially available kits (from DIALAB GmbH, Wr. Neudorf, Austria).

Detection of antibodies to cytokines and endogenous «danger» molecules in the immunoglobulin preparations

The Bio-Plex Cytokine Assay (BioRad, Richmond, CA) was used to detect antibodies to human IL-1Ra, IL-6, IL-8, IL-10, IL-15, TNF, G-CSF, IFN-γ, IP-10, RANTES, MCP-1, MIP-1 alpha

and MIP-1 beta in a) the native IVIg, b) Fe(II) –exposed IVIg and c) pooled human IgM (kindly provided by Biotest AG, Dreieich, Germany). A human IgG1 monoclonal antibody was used as a control (22). The immunoglobulin preparations were added (at a concentration 200 µg/ml) to serial dilutions of the respective cytokines and incubated for 1 h at 37 °C following the manufacturer's protocol. The presence of antibodies to mouse IFN-γ, human MIG, HMGB1 (from HMGBiotech, Milan, Italy), recombinant human heat shock proteins (HSP 60 and HSP 70 obtained from Enzo Life Sciences, Lausen, Switzerland) and to human histones H3 and H4 (Roche Diagnostics, Mannheim, Germany) were tested for by ELISA. Plates (Nunc MaxiSorp) were coated with 2 µg/ml of the respective antigens in 0.1 M carbonate buffer and incubated overnight at 4 °C. After blocking and washing, the plates were incubated for 2 h at room temperature with increasing concentrations of the immunoglobulin preparations under study. The plates were then extensively washed and goat anti-human IgG antibody coupled to alkaline phosphatase was added and incubated for 1 h at room temperature. Immunoreactivities were revealed with p-nitrophenyl phosphate (Sigma-Aldrich) and the OD values were read at OD=405 nm.

Statistical analysis

All statistical analyses were performed using the GraphPad Prism 4 software (La Jolla, CA, USA) and the R software (v.3.0.2). Data were expressed as mean \pm SD. Differences in survival curves were compared using the Mantel-Haenszel test. For other data, the differences in the mean values between groups were analyzed with the two-tailed Student t test. Differences were considered significant when p < 0.05. Pearson correlation coefficient was used to compare gene expression profiles in different organs.

Results

Protective effect of ferrous ions-exposed IVIg improved survival in sepsis and aseptic SIRS

The intravenous administration of a single dose of 50 mg/kg of the Fe(II)-ions-modified IVIg significantly decreased the mortality of animals with endotoxemia (Fig. 1A - left panel). The same treatment (at dose of 250 mg/kg) was protective in zymosan-induced SIRS and in polymicrobial CLP sepsis (Fig. 1A, middle and right panels). The survival curves of all animals in the experimental series are shown on Fig. S1. The native immunoglobulin preparation, regardless of the dose used, had no effect thus confirming results from numerous clinical trials. The mechanisms of protective activity of IVIg with enhanced polyspecificity in endotoxemia were studied in more details. The infusion of the modified IVIg one hour post-LPS injection still had a protective effect while the treatment delayed for 4 hours failed to improve the survival (Fig. 1B). The iron content of IVIg increased after the Fe(II) treatment from 0.028 mg/g IgG to 7.2 mg/g IgG (Table S1). Nevertheless the beneficial effect of the preparation was still present even upon dialysis against EDTA which decreased the iron concentration (Fig. 1C).

Gene expression analysis

Microarray data of organ samples from Fe(II)-exposed IVIg treated endotoxemic mice were clustured by an unfiltered unsupervised approach. Unique responses to Fe(II)-exposed IVIg were observed in all organs, but were most pronounced in the liver, supported by a weaker correlation (Pearson correlation coefficient r (IVIg vs Fe(II)-exposed IVIg) = 0.83 in kidney and r = 0.45 in liver). Although, several differentially expressed genes (DEG) could be identified, they did not constitute an easily interpretable pattern of signaling pathways (Figure S2).

Treatment with IVIg with enhanced polyspecificity affects plasma levels of pro- and antiinflammatory cytokines

IVIg exposure to Fe(II) ions did not result in its enhanced binding to bacterial LPS (Fig. S3), indicating that the beneficial effect of modified IVIg was due to its interactions with host defense mechanisms rather than to a direct neutralization of LPS. The observed beneficial effect on survival suggests that the levels of pro-inflammatory molecules might be influenced by the treatment. Indeed, plasma levels of TNF-α, IL-6 and IFN-γ were significantly reduced in animals injected with modified IVIg (Fig. 2A). In contrast, the levels of IL-10 and IL-22 were significantly elevated. A semi-quantitative dot-blot protein array technique allowed the analysis of a larger panel of inflammation-related molecules. Significantly lower levels of IL-6, LIX (CXCL5), IL-12p70 and MIP-1gamma (CCL9) were detected after treatment with the IVIg with increased polyspecificity (Fig. 2B, 2C).

Effect of ferrous ions-exposed IVIg on the complement system, coagulation and organ dysfunction of mice with LPS-induced sepsis

Complement activation and the release of anaphylatoxins play an important role in the pathogenesis of sepsis (23, 24). The levels of C3 complement component in the circulation mirror the severity of on-going inflammation. The LPS injection resulted in a sharp and prolonged drop in C3 levels and the administration of the native IVIg did not significantly affect this level. Interestingly, the infusion of the same single dose of the IVIg with enhanced polyspecificity led to a significantly diminished consumption of C3 (Fig. 3A). We analyzed in more detail the interaction of the immunoglobulin preparations with human complement components. The binding of the pooled human IgG to the C1q component was enhanced after its

exposure to ferrous ions (Fig. 3B). While the pre-incubation of C1q with Fe(II)-exposed IVIg did not affect the interaction of the latter with immobilized IgG1 (not shown), it significantly inhibited C1q binding to CRP (Fig. 3C). Interestingly, the modified immunoglobulin preparation bound to C3a, but not to C5a anaphylatoxins (Fig. 3D).

Systemic inflammation in sepsis is accompanied by an over-activation of the coagulation cascade and results in disseminated clots in small blood vessels, contributing to organ failure. A single infusion of ferrous ions-exposed IVIg to endotoxemic mice normalized the blood clotting time (Fig. 3E). This could well explain also the observation that liver injury was significantly ameliorated in the group treated with the modified IVIg as assessed by the capacity of ferrous ions-exposed IVIg to decrease the LPS-induced elevation of serum bilirubin levels (Fig. 3F). Creatinine levels at the same early time point were not affected.

The Fe(II)-modified IVIg acquires additional polyspecificity that includes ability to bind pro-inflammatory mediators and endogenous "danger" molecules

The exposure of therapeutic IVIg to increasing concentrations of ferrous ions results in its ability to bind to a broader number of antigens in a HUVEC lysate (Fig. S4). In addition, anti-human IP-10, IFN-gamma, anti-MIG and anti-HMGBox1 reactivities were observed in Fe(II)-exposed IVIg using a BioPlex or ELISA assays, whereas native IVIg and pooled IgM preparations had no detectable antibodies to the studied cytokines. Antibodies to endogenous "danger" molecules - free Histone 3 and Histone 4, reported to possess a toxic effect on endothelial cells in sepsis (25), as well as to heat shock proteins were also present in the modified preparation (Table 1 and Fig. S5).

Discussion

Passive immunotherapy with commercially available therapeutic IVIg exposed *in vitro* to a low concentration of pro-oxidative ferrous ions, significantly improved survival in animals with experimental sepsis and aseptic SIRS. The same immunoglobulin preparations were not protective in their native form. The improved survival was accompanied by lower serum levels of pro-inflammatory cytokines, normalization of the coagulation time, diminished consumption of C3 complement component and attenuated organ injury or dysfunction. These multimodal effects strongly suggest that the ferrous-exposed IVIg has acquired novel biological properties. Pooled IVIg are highly polyspecific therapeutic preparations. Previous studies point that their exposure to Fe(II)-ions or to ROS, released in inflammation sites, amplify further this polyspecificity (15).

IVIg are known to suppress-the harmful effects of activated complement cascade components by preventing their binding to the respective receptors (26). In the fluid phase the IVIg with enhanced polyspecificity did not perturb the capacity of C1q to recognize IgG immune complexes. The latter is in agreement with previous studies, showing that even at high doses IVIg do not alter C1q binding and complement activation on sensitized sheep erythrocytes used as a model of cell-bound immune complexes (27). Therefore, the administration of Fe(II) ions-modified IVIg will not affect C1q functions during the early phase of defense against pathogens. Moreover, C1q binds better to the surface immobilized Fe(II) ions-modified IVIg (as a model of immune complex) than to the native IVIg, suggesting that the modified preparation could be more effective in opsonizing invading bacteria. When allowed to interact in a fluid phase with C1q Fe(II) ions-exposed IVIg inhibited the binding of C1q to surface-bound CRP, suggesting that attenuation of the complement activation on damaged host cells or on bacteria could result.

Since CRP levels are increased up to 50 fold in sepsis (28) the inhibition of its effector functions may be beneficial by reducing the ongoing inflammation. In our experiments the Fe(II) ions-modified IVIg also showed enhanced binding to C3a anaphylatoxin expected to result *in vivo* in its rapid neutralization and elimination.

The net beneficial effect of the blunted host response upon administration of IVIg with enhaced polyspecificity were reflected in less organ damage primarily in kidneys (Fig. S6). The transcriptome analysis showed a high variability between the individual animals suggesting that detection of the differential expression of particular genes might be a transitory phenomenon in a highly disordered (chaotic) gene network dynamics, characteristic of the septic state (29). For instance, some of the untreated septic animals clustered with the IVIG treated and some - with the modified IVIg. Nevertheless, the experiment showed detectable changes in the gene expression patterns, specific for the modified IVIg. A different set up with larger experimental groups will be necessary to trace these changes of the cascade ultimately dampening the genomic storm (Fig. S2).

The idea that polyspecific natural antibodies may act as a buffering system that prevents brisk changes in the levels of circulating cytokines, hormones, etc., has been proposed twenty years ago (30). The immune-suppressive properties of plasma during sepsis have been discussed extensively before. Diverse factors are currently considered in this context: IL-10, extracellular ubiquitin, circulating CD14, epinephrine and other inducible signals (31). Such a convergence of different activities towards an anti-inflammatory plasma compartment indicates the importance of this principle. It is thus not surprising that the quasi-complete repertoire of circulating natural antibodies can be conditionally recruited to that function too.

It is counter-intuitive that naturally polyspecific IgG antibodies should generally limit or block cytokine activities in the tissues. On the other hand, in vivo induced polyspecificity would be limited in space (to the site of inflammation) and time (IgG polyspecificity would result in a short half-life (32) and molecules gaining systemic access would be quickly diluted). Thus, its effect would be tightly controlled. It is possible that the *in vitro* treatment of IVIg with proteindestabilizing agents serendipitously activated this function on a large scale. Intravenous infusion of immunoglobulin preparation with additionally enhanced polyspecificity would apply systemically a "buffer" system that evolution have meant to have a local effect. During the event of uncontrolled systemic inflammation (sepsis) this may be helpful as it would enhance the inherent anti-inflammatory activity of plasma. Our data provide evidence to support this hypothesis as only the modified, but not the native IVIg preparation binds to a) a panel of antigens in a HUVEC lysate, b) mediators of inflammation – IFNy, IP-10/CXCL10, MIG/CXCL9 and HGMB1, c) the complement components C1q and C3a, and d) the hydrophobic "danger signals" heat shock proteins and the free extracellular histones H3 and H4 (see Refs. (15, 25) and Figs. 3, S4 and S5). Importantly, our studies revealed that Fe(II) ions exposure is associated with considerable increase in the hydrophobicity of antigen binding sites of IgG (15). As endogenous danger signals have frequently hydrophobic nature, antibodymediated non-specific quenching of hydrophobicity may also well explain the systemic antiinflammatory effect of modified IVIg (33, 34).

Furthermore, the elimination of the formed immune complexes by Fc receptor-expressing phagocytes may contribute to the down-regulation of the ongoing severe generalized inflammatory reaction.

Although the dependence of the observed therapeutic effect of modified IVIg on lymphocytes and macrophages is compatible with the "immunosomatics" mechanism (30), an effect, mediated by a particular mononuclear cell population, cannot be ruled out. Innate response activator (IRA) CD19^{low}CD5^{low}CD93⁺ B cells have been implicated recently in the control of experimental sepsis in mice (35). Indeed, our recent studies show that, unlike native IVIg, ferrous ions exposure-modified IVIg induced in the LPS-injected mice higher levels of IRA B cells and the beneficial effect of the treatment on survival correlated with this increase (36).

Conclusion

This work contains evidence that pleiotropic therapeutic approaches offer a viable strategy for treatment of sepsis and aseptic SIRS. Apart from the mechanistic explanation, the pleiotropy principle also justifies the use of mouse models despite the controversy on differences in transcriptome profiles in humans and mice with severe inflammatory syndromes (37, 38).

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For More Information The microarray data generated in this publication have been deposited in NCBI's Gene Expression Omnibus (GEO) and are accessible through the accession number GSE55964.

Competing financial interests

The authors declare competing financial interests: TV is applicant and TV and JD are inventors of a patent.

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Table 1. Exposure of IVIg to ferrous ions results in the appearance of reactivity to some of the tested human cytokines, complement components and "danger" molecules. Antibody reactivity to mouse IFN- γ were measured in a control ELISA experiment. NS* - not shown.

	Inflammation-related and "danger" molecules	Binding appears after IVIg exposure to ferrous ions	Method used	Reference	
Cytokines					
1	MIG	Yes	ELISA	Fig. S5	
2	IP-10	Yes	BioPlex Array	Fig. S5	
3	HMGB1	Yes	ELISA	Fig. S5	
4	IFN-γ	Yes	BioPlex Array	Fig. S5 and Ref.15	
5	Mouse IFN-γ	No	ELISA	Fig. S5	
6	IL-8	No	BioPlex Array	NS	
7	IL-10	No	BioPlex Array	NS	
8	IL-15	No	BioPlex Array	NS	
9	G-CSF	No	BioPlex Array	NS	
10	MCP-1	No	BioPlex Array	NS	
11	MIP-α	No	BioPlex Array	NS	
12	МІР-β	No	BioPlex Array	NS	
13	TNF-α	No	BioPlex Array	Fig. S5	
14	IL-6	No	ELISA	NS	
15	IL1-RA	No	ELISA	NS	
16	RANTES	No	BioPlex Array	NS	
	Complement components				
17	C1q	Yes	ELISA	Fig. 3B	
18	C3a	Yes	ELISA	Fig. 3D	
19	C5a	No	ELISA	Fig. 3D	
	"Danger" molecules				
20	Histone 3	Yes	ELISA	Fig. S5	
21	Histone 4	Yes	ELISA	NS	
22	HSP 60	Yes	ELISA	Fig. S5	
23	HSP 70	Yes	ELISA	Fig. S5	
24	HSP 90	Yes	ELISA	NS	

Figure legends:

Figure 1. Passive immunotherapy with IVIg modified by Fe(II) exposure improves animal survival in two aseptic SIRS and one polymicrobial sepsis models. Outbred female ICR mice were used in the experiments. (A) Endotoxemia was induced by the i.p. injection of 10 mg/kg E. coli LPS (left panel). Zymosan-induced systemic inflammation (middle panel) was caused by i.p. injection of 500 mg/kg zymosan. Polymicrobial CLP-induced sepsis (right panel) was performed as described in the Materials and methods section. The survival was compared between the control group, injected i.v. with PBS (open squares, n = 12), the group with endotoxemia treated with 50 mg/kg or of 250 mg/kg (in zymosan-induced inflammation and CLP sepsis) native IVIg (triangles, n = 12), and a group treated with 50 mg/kg (endotoxemia) or 250 mg/kg (zymosaninduced inflammation and CLP sepsis) Fe(II)-exposed IVIg (circles, n = 12); *p<0.05, Mantel-Haenszel logrank test). (B) The beneficial effect of the Fe(II)-exposed preparation on survival was still observed if its administration was delayed. Endotoxemia was induced as described above and the mice were treated i.v. with 50 mg/kg of the native (open bars) or Fe(II)-exposed IVIg (black bars) at different time points – minutes before, one, or four hours after the i.p. injection of LPS. (C) Residual iron ions in Fe(II)-exposed IVIg did not contribute to its protective effect in LPS sepsis. ICR mice (n=12) were injected i.p with 10 mg/kg E. coli LPS and treated i.v. with 50 mg/kg Fe(II)-exposed IVIg with high (7.2 mg/g IgG, black circles) or low (0.66 mg/g IgG, white circles) iron concentration or with PBS only (open squares).

Figure 2. Serum levels of inflammation-related molecules in treated endotoxemic mice. (A) Endotoxemia was induced by the i.p. injection of 10 mg/kg *E. coli* LPS. A control group was

injected i.v. with PBS (n = 10, light grey bars), a second group was injected i.v. with 50 mg/kg of native IVIg (n = 10, dark grey bars), and a third group - with 50 mg/kg of Fe(II)-exposed IVIg (n = 10, black bars). Cytokine levels in untreated mice are represented with white bars. Sera were obtained 4, 24 and 48 h after LPS injection and cytokine levels were measured by ELISA. Data represent mean pg/ml values \pm SD of quadruplicate wells, *p<0.05 (paired Student t-test). (B) Scan of dot blots evaluating semi-quantitatively of serum levels of a larger panel of inflammatory molecules two hours after LPS injection. Each blot is representative of three individual experiments. (C) The individual dots (see panel B) were subjected to densitometry and the data were analyzed using the Image-Tool v2.0 for Windows (UTHSCSA, San Antonio, TX, USA) software package and were represented in relative units of grey intensity (n = 3, mean \pm SD, *p<0.05, paired Student t-test).

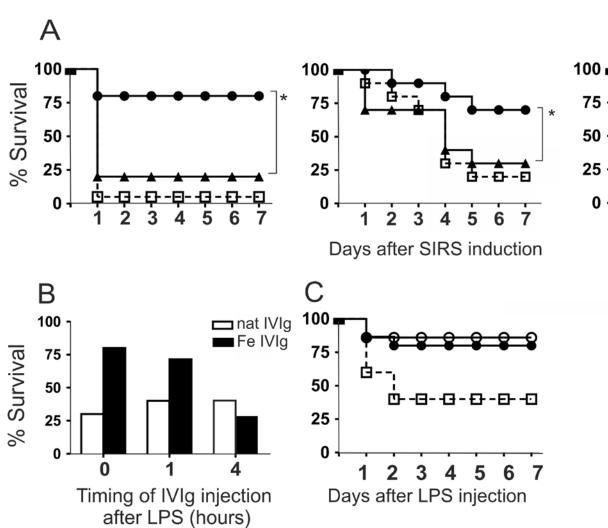
Figure 3. A single administration of Fe(II) exposure-modified IVIg overcomes the coagulation and complement abnormalities in LPS-induced sepsis. (A) Endotoxemia was induced to ICR mice by the i.p. injection of 10 mg/kg *E. coli* LPS. A control group (n=10, open squares) was injected i.v. with PBS, a second group was treated with 50 mg/kg native IVIg (n = 10, black triangles), and a third was treated with 50 mg/kg Fe(II)-exposed IVIg (n=10, black circles). Serum samples were collected at 4, 24 and 48 hour after LPS administration and the C3 levels in plasma were measured using a commercial Mouse C3 ELISA kit, *p<0.05, paired t-test. (B) The exposure of IVIg to Fe(II)-ions results in its increased binding to C1q molecules (black circles) compared to that of native IVIg (black triangles), mean±SD. (C) C1q pretreatment with Fe(II)-exposure modified IVIg inhibits the binding of the former to CRP (black circles); in comparison to C1q pretreated with native IVIg (black triangles). The binding of untreated C1q is

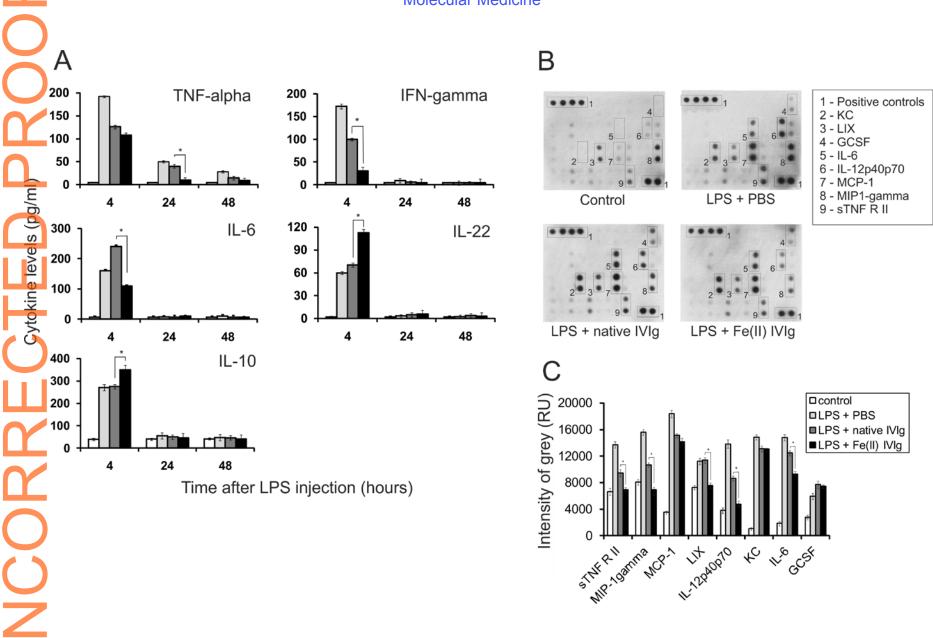
presented with open squares. (D) The exposure of IVIg to increasing concentrations of ferrous ions results in an enhanced binding to the C3a, but not the C5a component Anti-C3a and anti-C5a reactivity was compared by ELISA in IVIg, exposed to increasing concentrations of ferrous ions. (E) Endotoxemia was induced to ICR mice by the i.p. injection of 10 mg/kg *E. coli* LPS. A control group was treated i.v. with PBS (n=10, light grey bars), a group was treated with 50 mg/kg native IVIg (n = 10, dark grey bars), and another was treated with 50 mg/kg Fe(II)-exposed IVIg (n=10, black bars). Animals were bled 4 hours after the injection of LPS and coagulation time was measured as previously described (21), *p<0.05; ***p<0.001, t-test. (F) Bilirubin serum levels measured four hours after LPS administration in groups of mice treated with native (n=10, dark grey bars), Fe(II) -exposed IVIg (n=10, black bars) or PBS alone (n=10, light grey bars). *p<0.05, t-test.

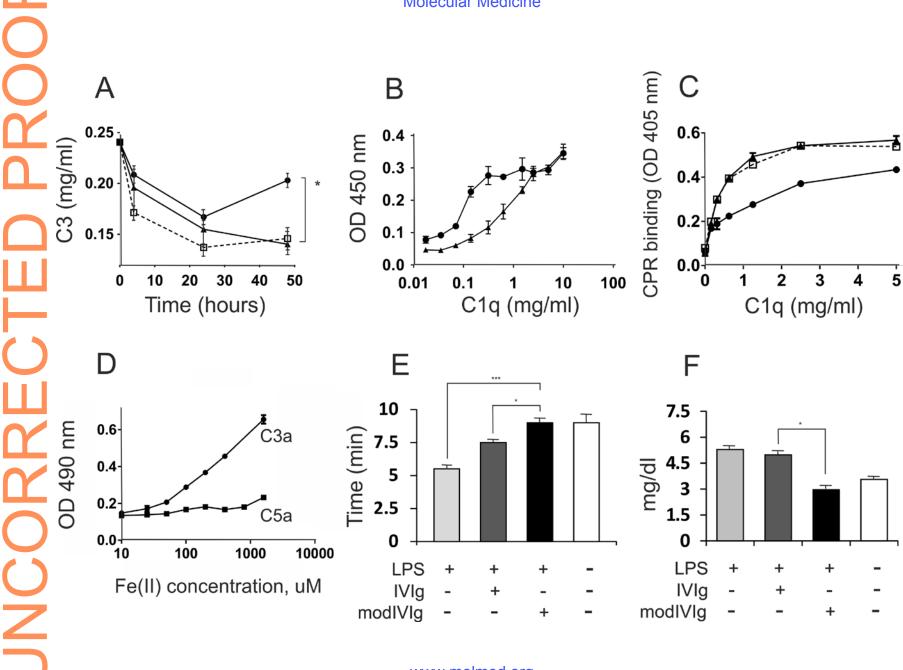
→ control

- nat IVIg
- Fe IVIg









Supporting information:

Materials and Methods

Flame atomic absorption spectroscopy

The concentration of iron ions in the IVIg preparations under study was measured by flame atomic absorption spectrometry. The samples were diluted 1:2 with a 20% (W/V) trichloro acetic acid solution, and heated in a heating block for 15 minutes at 90°C. They were next cooled, centrifugated and the iron concentration was determined using a Perkin Elmer M 5000 atomic absorption spectrometer.

Gene expression analysis

C57Bl/6 mice were used in this analysis. Blood samples, liver and kidney homogenates were obtained and processed by standard protocols. RNA was extracted using the RNeasy minikit (Qiagen, Hilden, Germany) in a balanced design comprising three biological replicates from blood and four from liver and kidney of each treatment group (vehicle, native IVIg, Fe(II)-exposed IVIg) respectively. Biotinylated probes were prepared using the TargetAmpTM-Nano Labeling Kit for Illumina® Expression BeadChip® (Illumina, San Diego, CA). The arrays were scanned using the Illumina BeadArray Reader 500 X.

Illumina mouseRef8 V2 chips were read by Genome Studio (v.1.9, and annotation MouseRef-8_V2_0_R3_11278551_A) yielding 25,697 bead probes for each sample. Analysis was performed consistently using R software version 3.02 (http://www.r-project.org/) and Bioconductor (1) packages. Raw data from each organ was separately subjected to robust spline normalization, quality control and background correction with the R package Lumi (2). Two samples (one from blood cells, one from kidney) did not pass quality control and were discarded from analysis. Bead types having detection values p < 0.05 of the RNA samples

were called "present" and taken for further analysis. Differentially expressed genes (DEG) were filtered according to microarray quality control criteria (MAQC, (3)) by: 1) at least two-fold changes and 2) p-values from moderated t-statistics <0.05 using "Limma" (24). Further, these statistics were computed for all the contrasts (IVIg, Fe(II) exposure-modified IVIg (modIVIg), vehicle) using linear model fits, moderated t-and F-statistics, and log-odds of differential expression by empirical Bayes moderation. Gene set enrichment analysis was performed by hypergeometric tests with the WebGestalt tool (4). Annotation, log2 fold changes, top enriched categories and p-values are provided as Supplemental gene expression data.

Binding of and modified IVIg to bacterial lipopolysaccharide

ELISA plates were coated with *E. coli* LPS (B 055:B5, Sigma #L2880, at 5 ug/ml in PBS) and blocked. Serial dilutions of the IVIg preparations were added in triplicate wells and the binding of the pooled human IgG was measured by ELISA.

In a separate experiment the native- and the Fe(II)-exposed IVIg were mixed in the wells of a blocked ELISA plate with increasing concentrations of *E.coli* LPS and incubated for two hours at room temperature. These mixtures were transferred to a second LPS-coated plate and the binding of human IgG was measured by ELISA as described in the Methods section. Its inhibition in the presence of free LPS was determined and presented in arbitrary units.

Immunoblot analysis of binding to HUVEC antigens

HUVEC cells lysates were subjected to 10% gel SDS-PAGE and transferred to nitrocellulose membranes (Scheicher & Schuell, Dassel, Germany) by a Mini Transfer Blot system (BioRad, Richmond, CA). Next, the membranes were incubated for 1 hour at room temperature in TBS containing 0,4% Tween 20, cut into strips and incubated for 1 h at room

temperature with native or Fe(II)-exposed IVIg preparations at concentrations of 0,2 mg/ml. After extensive washing the strips were incubated with an anti-human IgG antibody (Fcspecific), conjugated to alkaline phosphatase (Sigma Chemical Co, St. Louis, MO) and revealed using the nitro blue tetrazolium/bromo-chloro-indolyl-phosphate substrates (Sigma).

Antibodies to pro-inflammatory and "danger" molecules in the modified IVIg

The presence of antibodies to cytokines, complement components and "danger" molecules was studied using ELISA and BioPlex Array (see the Materials and Methods section of the main text).

Histology

ICR mice (8-12 weeks old, 18-22 g) were used in this experiment. Endotoxemia was induced by the i.p. injection of 10 mg/kg *E. coli* LPS (B 055:B5, #L2880, Sigma-Aldrich St. Luis, MO). Single dose of native, modified IVIg preparations or PBS alone was injected intravenously (i.v.) 10 min before the administration of LPS. Kidneys and livers from each experimental group were collected 24 h after the injection of LPS. The organs were fixed in 4% buffered formalin, embedded in paraffin, sectioned into 3.5 μm thick slices and stained with Hematoxylin and Eosin (H&E). Pathologists blinded for the treatments estimated kidney damage in mice challenged with LPS + vehicle (n=8), LPS + native IVIg (n=10), LPS + Fe(II)-exposed IVIg (n=10) using the following criteria: the presence of brush-border defects, flattening of epithelia, loss of nuclei and presence of debris all indicative of an acute tubular injury (ATI) (5). Presence of liver damage was graded based on sinusoidal congestion, nuclear and cytoplasmic irregularities, signs of cellular ischemia, sinusoidal infiltration and single cell necrosis (6).

Table S1. The concentration of iron in the studied immunoglobulin preparations was measured by atomic absorption spectrometric analysis.

Preparation	Iron concentration	
Native (untreated) IVIg	0.028 mg/g IgG	
Fe(II)-exposed IVIg	7.2 mg/g IgG	
Fe(II)-exposed IVIg dialyzed additionally	0.66 mg/g IgG	
against 4mM EDTA		

Figure legends:

Figure S1. Reduced mortality in three different experimental models of sepsis after passive immunotherapy with an IVIg preparation, modified by Fe(II) exposure. (A) Survival curves of ICR mice with bacterial LPS-induced endotoxemia. Animals were injected i.p. with 10 mg/kg E. coli LPS. A survival study was performed comparing a control group, treated i.v. with PBS (open squares, n = 12), a group treated with 2 mg/kg (left panel), 10 mg/kg (middle panel) or 50 mg/kg (right panel) native IVIg (triangles, n = 12), and a group treated with 2 mg/kg (left panel), 10 mg/kg (middle panel) or 50 mg/kg (right panel) Fe(II)exposed IVIg (circles, n = 12); *p<0.05, Mantel-Haenszel logrank test). (B) Survival curves of ICR mice with zymosan-induced systemic inflammation. Animals were injected i.p. with 500 mg/kg zymosan. A survival study was performed comparing a control group, treated i.v. with PBS (open squares, n = 12), a group treated with 10 mg/kg (left panel), 50 mg/kg (middle panel) or 250 mg/kg (right panel) native IVIg (triangles, n = 12), and a group treated with 10 mg/kg (left panel), 50 mg/kg (middle panel) or 250 mg/kg (right panel) Fe(II)exposed IVIg (circles, n = 12); *p<0.05, Mantel- Haenszel logrank test). (C) Survival curves of ICR mice with polymicrobial CLP-induced sepsis. Survival was compared between a control group, injected i.v. with PBS (open squares, n = 12), a group treated with 10 mg/kg (left panel), 50 mg/kg (middle panel) or 250 mg/kg (right panel) of native IVIg (triangles, n =12), and a group treated with 10 mg/kg (left panel), 50 mg/kg (middle panel) or 250 mg/kg (right panel) of Fe(II)-exposed IVIg (circles, n = 12). *p<0.05, Mantel- Haenszel logrank test).

Figure S2. Blood cell, liver and kidney gene expression in LPS-injected mice treated with native- or Fe(II)-exposed IVIg (modIVIg). Heatmaps depicting the differentially expressed genes (DEG, average two-fold change and p<0.05) for Fe(II)-exposed IVIg versus

native IVIg in blood (A), liver (B) and kidney (C), respectively. Resolved log2 signals of biological replicates were row-wise scaled (z-score) accompanied by vertical and horizontal hierarchical correlation clustering (technical probe replicates are indicated by ".1". Definition of gene symbols is provided in the supplement files. The Venn diagrams depict the number of DEG for all contrasts (D).

Figure S3. The exposure of IVIg to ferrous ions does not affect its binding to LPS. (A) Increasing concentrations of native- and Fe(II)-exposed IVIg preparations (marked with squares and circles respectively) were left to interact with immobilized LPS and human IgG binding was evaluated by ELISA. (B) Both preparations were pre-incubated with increasing concentrations of LPS and the mixtures were transferred to the LPS-coated wells. Next, the binding of human IgG was measured by ELISA and its inhibition (presented in arbitrary units - AU) was plotted against the concentration of free LPS used.

Figure S4. Exposure of IVIg to increasing concentrations of Fe(II) ions results in its enhanced binding to antigens in a HUVEC lysate. The membranes were incubated with $200 \,\mu\text{g/ml}$ of the native IVIg or of the same preparation exposed to ferrous ions.

Figure S5. Ferrous ions-exposed IVIg acquires the ability to bind to various human pro-inflammatory molecules. The figure presents several examples whereas all results are showed in Table 1 of the main text. Reactivity to IFN- γ , IP-10 and TNF α were measured using BioPlex Array and in all other cases ELISA was used (see the Materials and Methods section of the main text). In a control experiment the reactivities of both studied preparations to mouse IFN- γ (mIFN- γ) were compared. The pie-chart summarizes all data and points out that modified IVIg acquires the ability bind to almost half of the unselected tested inflammation-related molecules.

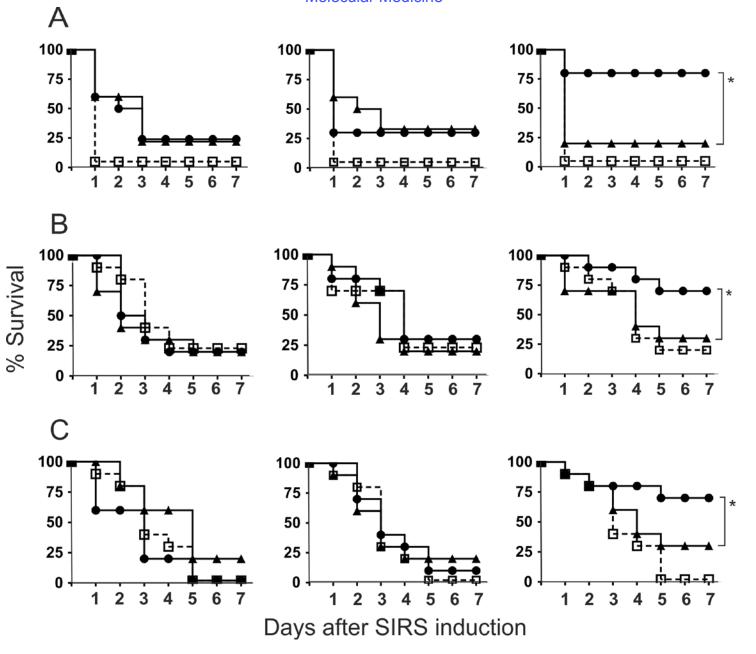
Figure S6. Tissue damage score from histological specimen (kidney - A and B, liver - C and D) from mice with endotoxemia treated with native or Fe(II)-exposed IVIg. Kidney injury was evaluated by the presence of brush-border defects, flattening of epithelia, loss of nuclei and presence of debris, summarized in the acute tubular injury (ATI) score (1). Liver damage scoring was based on sinusoidal congestion, nuclear and cytoplasmic irregularities, signs of cellular ischemia, sinusoidal infiltration and single cell necrosis. Summaries are provided in A and C for kidney and liver, respectively. *p<0.05, *t*-test (Welch-modified, accounting for differences in variances) Fe(II)-exposed IVIg vs the other groups given in parametric plots.

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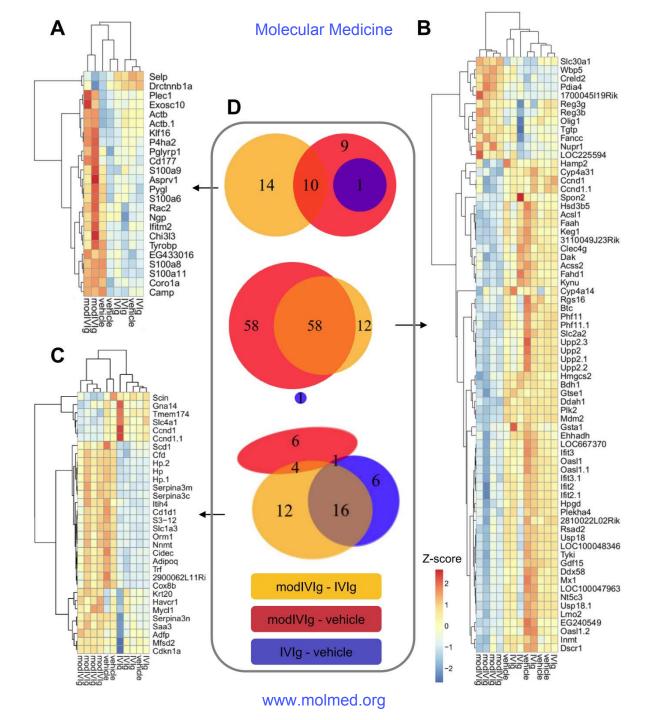
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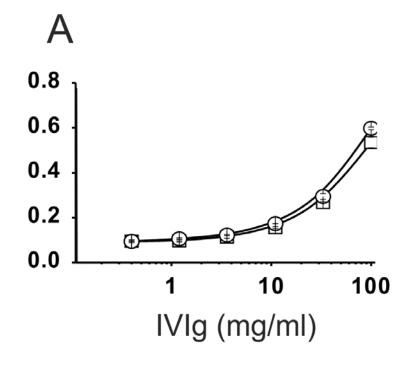
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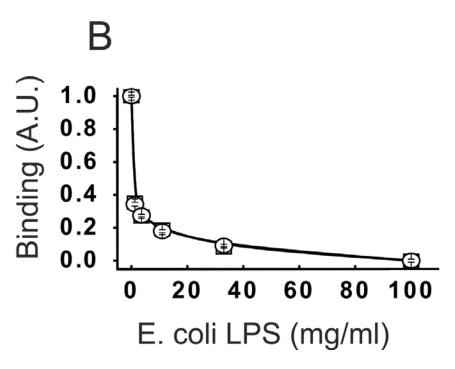
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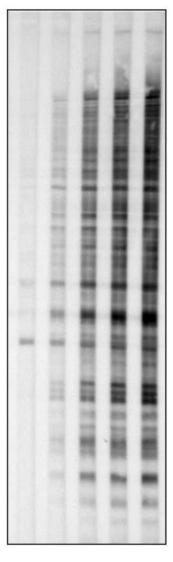




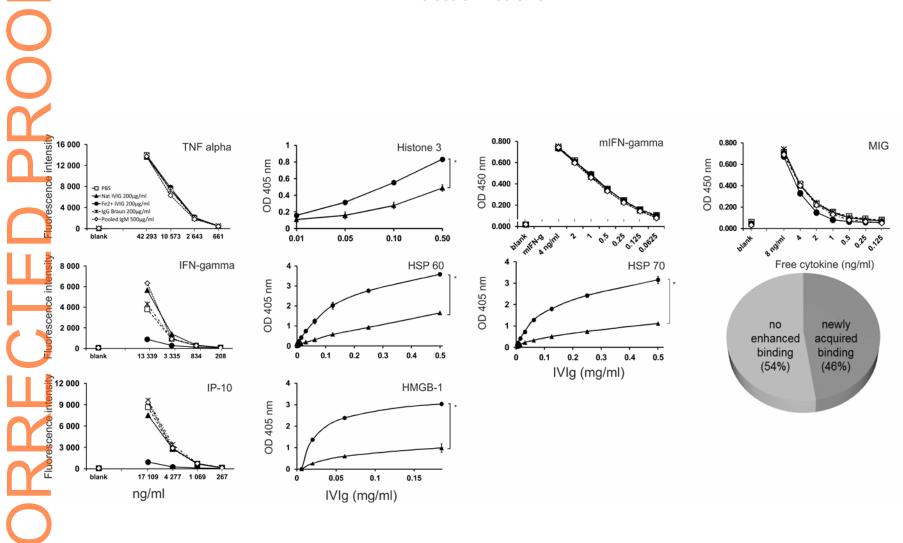


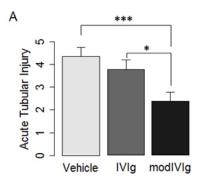
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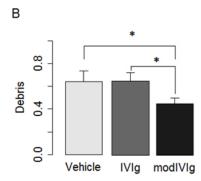
0 0.10 0.25 0.50 1.00

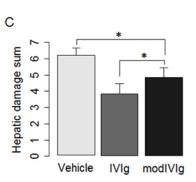


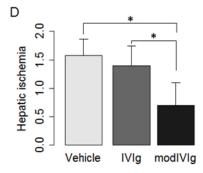
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ILMN_Gene	average log2FC	(Vig-Vehicle	modIVla-Vehicle	modNlg-IVIg	P (moderated t-statistics) IVIo-Vehick modIVIo	o-Vehicle m	glVI-glVlbor	Probe_ld	Probe_Sequence
2900062L11RIK		-1.416697695	-0.173654787	1.243042908	0.037431 0.775	5410146	0.0463929	ILMN_2609182	CCTTGTAGGTCCACATTGAATGCCGTTTTCTTATATCTTCTAGGGTGTGT Mus musiculus RIKEN
ADFP		-0.594814157	0.717284553	1.31209871		1508139 0			AGAAAATTCAGGGTGCTCAGGATAAGCTCTATGTCTCGTGGGTGG
ADIPOQ		-2.400097295	0.232938434	2.633035729		1813749 0			GACTGCAACTACCCATAGCCCATACACCAGGAGAATCATGGAACAGTCG Mus musculus adipone
AOC3		-1.531980725	-0.312202827	1.219777897		3477318 0			GAGAGGGAATCAGGTTCTGCCTGCCCTTCTGAACATCATGGATTGGAAC Mzs musculus amine o
APOC1 AQP4		-1.299479883 -1.01024592	-0.347932302 -1.330365875	0.951547381 -0.320119956		0706361 0 6900839 0	0.10537082 0.51716872	ILMN_2599794 ILMN 2757232	AAAGTGAAGGAGAAGTTGAAGACCACGTTCTCCTGAGCACCTGGCGGGI Mas musiculus apolipoj GTGCAGAGCATATGGACACCTCTGTGAGGAAGCTGGCATTGTCCATCGT Mas musiculus apolipoj
CAR3		-1.362158574	-0.203605447	1.158553127	0.0381835 0.729	9427688 0	0.05319186	II MN 2630047	CCCTGGCTCTGCTAAGACCATCCTGAACAATGGGAAGACCTGCAGAGTI Mus musiculus carboni
CCND1		0.381308916	-0.86206292	-1.243371836	0.3771588 0.062	2804125 0	0.00844535	II MN 2601471	GCCATAGACTGGTGGTGGGTCCACGGAATCTGCCCTGTGACATGAAAGGC
CCND1		0.398618927		-1.169531129		0707034 0			GGCCATAGACTGGTGGTGGGTCCACGGAATCTGCCCTGTGACATGAAAGG
CD1D1		-1.46195839	-0.084498452	1.377459938	0.028787 0.886	6047215 0	0.02683035	II MN 2734212	GAACTAGAAAGCATACTTCCTGCCCAAACAGACGCTCTGAGGTTAGTTG Mus musiculus CD1d1
CDKN1A		-0.598236148		1.248232406			0.02766547		CATCCTGGTCTGGACTGTCTACCCTTAGCCCGCACCCCAAGAACATGTA Mus musiculus circlin-d
CED		-2.965653662	0.530785528	3.496419189			0.0010017		CAGTOGAAGGTGTGGTTACGTGGGGGCTCTCGCGTCTGTGGCAATGGCA Mus musculus comple
CIDEA		-1.801244902	-0.559559857	1.241685045	0.0150816 0.385	5987296 0	0.05452063		GACGGGACAGTTCCTGGTCTATGCGGGCACATACATGCTCCGAGTACT(Mizi miziculus cell des
CIDEC		-1.456853318		1.317934099			0.00924771		GAGGTGTGCGAGAGAGTGTTAGAGGTGGAGATTAACTAAAGCCGGCATI Mus musiculus cell des
CLDN11		-1.645730543		0.224570363			89063747		GTGTCTAAGACTCTGGATACTGCAAGCTCCGTCCGGTGCATTTGTTCAG Mus musculus clisudin
COX88		-2.449745171		2.199432937			0.03034856	II MN 2727520	CGTTGGCTTCATGGTTCCAGCAGGATGGGTCTTAGCCCACTTGGAGAG(Mus musiculus cytochri
GNA14		0.487424158		-1.016789488		2600443 0			GCGCTGTGCCTTTCTGGATACTAAGAATACACTGTTCTACCCTGAACAC(Mus musiculus quantine
HAVCR1		-0.172270347					0.01981639		AAGCTTTGCAGAACGCAGCGGTTGTGCATTCCCGAGCTGAAGACAACAT Mga muscukus haqafti
HP		-2.096151341	0.392517347	2.488668689			0.00317183		CTGTTGTCACTCTCCTGCTCTGGGGTCAGCTTTTTGCTGTGGAGTTGGG Mus musiculus histoph
HP		-2.204083308		2.652672936		3855579 0			CTCTCCTGCTCTGGGGTCAGCTTTTTGCTGTGGGAGTTGGGCAATGATGC Mas musiculus historis
HP		-2 332394507	0.611622015	2 944016523			0.00181362		ACGAGTCCCGTTGGGGTCCAGCCTATCTTGAACGAGCACACCTTCTGTG Mas musiculus historis
IGFBP5		-0.499217044	-1.072298018			1436343 0			GGATAGTACAGTTCAGACAAGACTCCTTCAGATTCCGAGACGCCTACCG Mus musiculus insulin-li
IL4I1		-0.391947595				9367298 0			CCCAGCTCCTAAGCCAGCCCTCTTCAGGGCAGACAGACCACCTACACT/ Mus musiculus interfeul
ITIH4		-0.746467905	0.417565978				0.02270197		CACCTTATGCCAACAGGGACGCCTGTGAGGCCGAGACCTTGATGGGAA Mas musiculus inter sito
KRT20		0.083357367					0.01327319	II MN 2907285	GTGAACGGGTTCTCCAGGGAGGCAACTTCCCAGTTAATCTCATTGTCTC Mus musiculus kerstin 2
LOC100043671		-1.456136503				1242413 0			TCTGTCCACTTGTTCACAAAGCAGCCTTCACACGTGTCAGTGAGAGTTCI PREDICTED: Mus mu
MFSD2		-1.051508402	0.381609069	1.433117471		2914016 0			AGGGCTGCTACTGTGAATATGCCAAGGACTGACTGGGCCTAGCTCGGA: Mas musiculus major fa
MYCL1		-0.022587085		1.087032205			0.01412011		GGGACTCCTGGAACTATCTTGGGAGGACAAGTGGTGAACAGGCTAAAG* Mus musiculus v-mvc n
MYL1		-1.077625097		0.734201609			0.10004509		GATAATCTCAAGTCCACACACCACCGTTATCATCTACTTGGGTCCACAC Mus musiculus mirosin.
NMT		-1.199571192	0.25428225	1.453853442			0.0229501		CTCCCTAGTGGGGCGAAAGCCAGGCAGATCTGAATGACATCCTGTGAT(Mas musiculus nicotina
ORM1		-2.030115548	-0.101961618	1.92815393			0.03164228		CCTTTGCCCACCTTATAGTGCTGAGGAAACATGGGGCCTTCATGCTTGC Mgs musiculus grosom
83-12		-1.472158014	0.190004481	1.662162495			0.00826673		CCTTAGTGTCTGCTTCACTCAAGCTGGTAGAACAGGCTACTCGCTGCAC Mus musiculus plasma
SAA3		-0.78514295	0.678302447				0.00786362		GAGGACTCAAGAGCTGACCAGTTTGCCAATGAGTGGGGCCGGAGTGGC Mas musiculus serum s
SCD1		-1.647443514	-0.537645069	1.109798445			0.02437171		CCAGTCAAAGTGCAAGACTACCTACCCACTGCATGAAAACCATTGCAG Mga muscukus seesrovi
SCIN		-0.002361753	-1.293221152			3086022 0		E MM 1250174	GCTTCTAAGCATTTCCCCGTCTGCTACTCTGTTTGCAGTGAGCTTTACTT Mus musiculus scinderi
SERRINASC		-1.001675207	0.217593835	1 219269043			0.00373395		CAAGGGACTCATCTCAGACCTGGATACGGATACACTGATGGTGCTGGTC Mas musiculus serine (
SERPINASM		-0.885792445	0.417494031	1 303286476		0410176 0			CCTCTTTATGGCCAAAGTCACTAACCCCAAGTGAACCTAAAGCTCCCCA. Mga muscukus serine (
SERPINAIN		-1.378347276	0.440050877	1.818398152			0.00575328		CCCCTGGCACTCCTACTTAGAACAAAGTAGCCTTTCTTTTAGTTCCCAGC Mus musiculus serine (
SGK2		0.533528881		0.541113519		0434508 0			GGCTGCCCAGAAACCGTCTGGTGGCTTCCTTTAGTGTGTCGTAAATCTC Mas musiculus serum/s
SGK2		0.639027332	1.000316869	0.361289537		1458472 0			GAGAGAAAGGGGAGGATGTAGGTGACCAGCACTCTGGGAGAGAGCCTC Mas musiculus serum/c
SLC1A3		-1.259241472	0.321836021	1.581077493			0.00954689		CCCGGGCTCTGCCTATGTGTGTACCCACTGTGGACTGGGAAACTTCAC1 Mai musculus solute o
SLC4A1		0.52824231	-0.744134243			8739204 0			CGCTCTTCCCCCACCGCCGTGCTTGGGAATAGGTTTTGCCAATAAACGT Mas musiculus solute o
SLC5A3		0.750679233	1.037239005	0.286559772			0.38412317		CCCAGTGGCTTCCATGGGTCATTCAGAGGCAGAACACCAGTAGATGCT Mas musiculus solute o
TMEM174		0.249557844	-0.848911432				0.00224195		GGCTGCCCCCAAGAGAGATACAGGAATTGTCAGGTCTTGAAGGTTTTGC Mai musiculus trinnimis
TRE		-1.477109112	0.245021332			8774584 0			CTGGGGTAGACAGACCGCTGGTTGGAACATCCCTATGGGCATGCTGT/ Musi musiculus transfer
1194		-1.411109112	U.A 1021332	1.744130944	0.0175221 0.646	U117-09 U		£MN_2405323	CTOSTOST POR PROPERTY OF THE P

ILMN_122

ILMN_276

NA

NA

Ulk1

Camp

8475

6604

			Database	:molecular	function Name:c	alcium ion binding	ID:GO:00055	509	
	In	dex	UserID	C=479; (Value	O=5; E=0.94; R=5.32; Gene Symbol	rawP=0.0022; adjP=0.0 Gene Name	336 EntrezGer		Ensembl
	1		ILMN_288 3392	NA	S100a11	S100 calcium binding protein A11 (calgizzarin)	20	-	ENSMUS G0000002 7907
)	2		ILMN_280 3674	NA	S100a9	S100 calcium binding protein A9 (calgranulin B)	20	_	ENSMUS G0000005
			ILMN_271			S100 calcium binding		_	ENSMUS G0000000
	3		2120 ILMN_269	NA	S100a6	protein A6 (calcyclin) myosin, light	<u>20</u>	_	1025 ENSMUS G0000005
	4		1780	NA	Myl3	polypeptide 3 S100 calcium binding	<u>17</u>	_	ENSMUS
	5		ILMN_271 0905	NA	S100a8	protein A8 (calgranulin A)	<u>20</u>	<u>)201</u>	G0000005 6054
)	Da	tabase	e:cellular co	omponent	Name:cytoplasmi	c membrane-bounded v	vesicle ID):GO:	:0016023
			e:cellular co	C=506; (O=6; E=1.00; R=6.01;	c membrane-bounded v rawP=0.0004; adjP=0.0 Gene Name	088		
	In	tabase idex	UserID ILMN_271	C=506; (Value	O=6; E=1.00; R=6.01; Gene Symbol	rawP=0.0004; adjP=0.0 Gene Name	088 EntrezGer	ne	Ensembl ENSMUS G0000004
]] -	In		UserID	C=506; (O=6; E=1.00; R=6.01;	rawP=0.0004; adjP=0.0	088 EntrezGer	n e 	Ensembl ENSMUS G0000004
]	In 1		UserID ILMN_271 2986 ILMN_123 6889	C=506; (Value	O=6; E=1.00; R=6.01; Gene Symbol	rawP=0.0004; adjP=0.0 Gene Name	088 EntrezGer <u>12</u>	ne <u> </u>	Ensembl ENSMUS G0000004 0809 ENSMUS G0000002 6580 ENSMUS
	1 2 3		UserID ILMN_271 2986 ILMN_123	C=506; (Value	O=6; E=1.00; R=6.01; Gene Symbol Chi3l3	rawP=0.0004; adjP=0.0 Gene Name chitinase 3-like 3	088 EntrezGer <u>12</u>	ne	Ensembl ENSMUS G0000004 0809 ENSMUS G0000002 6580 ENSMUS G0000002
	1 1 2		UserID ILMN_271 2986 ILMN_123 6889 ILMN_266	C=506; C Value	O=6; E=1.00; R=6.01; Gene Symbol Chi3l3 Selp	rawP=0.0004; adjP=0.0 Gene Name chitinase 3-like 3 selectin, platelet	088 EntrezGer 12 20	ne	Ensembl ENSMUS G0000004 0809 ENSMUS G0000002 6580 ENSMUS G0000002 7301 ENSMUS G0000003

unc-51 like kinase 1

antimicrobial peptide

cathelicidin

G0000002

ENSMUS

G0000003

22241 9512

12796 8357

Liver

Index	UserID		:molecular function 5105; O=27; E=11.04; Gene Symbol	Name:catalytic activity R=2.45; rawP=5.83e-07; adjP Gene Name	ID:GO:0003824 =4.55e-05 EntrezGene	Ensembl
1	230073	NA	Ddx58	DEAD (Asp-Glu-Ala-Asp) box polypeptide 58	<u>230073</u>	ENSMUSG000000402 96
2	107569		Nt5c3	5'-nucleotidase, cytosolic	<u>107569</u>	ENSMUSG000000297 80
3	76654	NA	Upp2	uridine phosphorylase 2	76654	ENSMUSG000000268 39
						ENSMUSG000000716
4	240549		Gm4952	predicted gene 4952 dihydroxyacetone kinase 2		ENSMUSG000000343
5	225913	INA	Dak	homolog (yeast) radical S-adenosyl	<u>225913</u>	<u>/1</u>
6	58185	NA	Rsad2	methionine domain containing 2	<u>58185</u>	ENSMUSG000000206 41
7	13119	NA	Cyp4a14	cytochrome P450, family 4, subfamily a, polypeptide 14	<u>13119</u>	ENSMUSG000000287
				indolethylamine N-		ENSMUSG000000034
8	21743		Inmt	methyltransferase myxovirus (influenza virus)	<u>21743</u>	
9	17857	NA	Mx1	resistance 1 transformed mouse 3T3	<u>17857</u>	NULL ENSMUSG000000201
10	17246	NA	Mdm2	cell double minute 2	<u>17246</u>	<u>84</u>
11	14081	NA	Acsl1	acyl-CoA synthetase long- chain family member 1	14081	ENSMUSG000000187
12	71911		Bdh1	3-hydroxybutyrate		ENSMUSG000000465
				dehydrogenase, type 1 glutathione S-transferase,		ENSMUSG000000741
13	14857	NA	Gsta1	alpha 1 (Ya) phenazine biosynthesis-	<u>14857</u>	<u>83</u>
14	67307	NA	Pbld2	like protein domain containing 2	<u>67307</u>	ENSMUSG000000200 68
15	64697	NA	Keg1	kidney expressed gene 1	<u>64697</u>	ENSMUSG000000246 94
16	12304	NA	Pdia4	protein disulfide isomerase associated 4	<u>12304</u>	ENSMUSG000000258 23
17	24110	NA	Usp18	ubiquitin specific peptidase 18	<u>24110</u>	ENSMUSG000000301
18	21822			T cell specific GTPase 1	21822	ENSMUSG000000789
			Tgtp1	·		ENSMUSG000000217
19	20620	NA	Plk2	polo-like kinase 2	<u>20620</u>	
20	15360	NA	Hmgcs2	3-hydroxy-3-methylglutaryl- Coenzyme A synthase 2	<u>15360</u>	
21	14073	NA	Faah	fatty acid amide hydrolase	<u>14073</u>	ENSMUSG000000341 71
				hydroxy-delta-5-steroid dehydrogenase, 3 beta-		
22	15496	NA	Hsd3b5	and steroid delta- isomerase 5	<u>15496</u>	ENSMUSG000000380 92
23	70789	NA	Kynu	kynureninase (L- kynurenine hydrolase)	<u>70789</u>	ENSMUSG000000268 66
24	231655	NA	Oasl1	2'-5' oligoadenylate synthetase-like 1	<u>231655</u>	ENSMUSG000000418 27
25	60525	NA	Acss2	acyl-CoA synthetase short- chain family member 2	60525	ENSMUSG000000276 05
26	68636	NA	Fahd1	fumarylacetoacetate hydrolase domain containing 1	68636	ENSMUSG000000453
27	74147		Ehhadh	enoyl-Coenzyme A, hydratase/3-hydroxyacyl Coenzyme A dehydrogenase		ENSMUSG000000228

	Data	base:biological proces	ss Name:defense response	ID:GO:000695	52
		C=803; O=13; E=1.7	72; R=7.55; rawP=8.70e-09; adjP	=2.38e-06	
Index	UserID Va	alue Gene Symbo	Gene Name	EntrezGene	Ensembl
					ENSMUSG000000789
1	21822 NA	Tgtp1	T cell specific GTPase 1	21822	<u>22</u>
			DEAD (Asp-Glu-Ala-Asp)		ENSMUSG000000402
2	230073 NA	Ddx58	box polypeptide 58	230073	<u>96</u>
					ENSMUSG000000307
3	56312 NA	Nupr1	nuclear protein 1	<u>56312</u>	17
		·	regenerating islet-derived		ENSMUSG000000713
4	18489 NA	Reg3b	3 beta	18489	56
		· ·	interferon-induced protein		
			with tetratricopeptide		ENSMUSG000000459
5	15958 NA	lfit2	repeats 2	15958	32
			interferon-induced protein		
			with tetratricopeptide		ENSMUSG000000748
6	15959 NA	lfit3	repeats 3	15959	
			radical S-adenosyl		_
			methionine domain		ENSMUSG000000206
7	58185 NA	Rsad2	containing 2	58185	
			regenerating islet-derived		ENSMUSG000000300
8	19695 NA	Reg3g	3 gamma	19695	
		- 3 - 3	kynureninase (L-		ENSMUSG000000268
9	70789 NA	Kynu	kynurenine hydrolase)	70789	
		,	hepcidin antimicrobial		ENSMUSG000000569
10	66438 NA	Hamp2	peptide 2	66438	
		7 19111 _P =	myxovirus (influenza virus)		
11	17857 NA	Mx1	resistance 1	17857	NULL
			spondin 2, extracellular	<u></u>	ENSMUSG000000373
12	100689 NA	Spon2	matrix protein	100689	
_			2'-5' oligoadenylate		ENSMUSG000000418
13	231655 NA	Oasl1	synthetase-like 1	231655	
-			-,		

Kidney

Database:cellular component	Name:lipid particle	ID:GO:0005811

Index	UserID	Value	Gene Symbol	Gene Name	EntrezGene	Ensembl
	ILMN_258					ENSMUS G0000000
1	8249	NA	Plin4	perilipin 4	<u>57435</u>	<u>2831</u>
	ILMN_121			cell death- inducing DNA fragmenta tion factor, alpha subunit- like		ENSMUS G0000002
2	5446	NA	Cidea	effector A	<u>12683</u>	<u>4526</u>
	ILMN_122			cell death- inducing DFFA-like		ENSMUS G0000003
3	2679	NA	Cidec	effector c	<u>14311</u>	0278
	ILMN_122					ENSMUS G0000002
4	2246	NA	Plin2	perilipin 2	<u>11520</u>	8494

Database:molecular function Name:carboxylic acid binding ID:GO:0031406

C=178; O=4; E=0.38; R=10.48; rawP=0.0005; adjP=0.0145

Index	UserID	Value	Gene Symbol	Gene Name	EntrezGene	Ensembl
1						ENSMUS
*	ILMN_259			apolipopro		G0000004
1	9794	NA	Apoc1	tein C-I	11812	0564

2	ILMN_263 4317 NA	Slc1a3	solute carrier family 1 (glial high affinity glutamate transporte r), member 3	ENSMUS G0000000 20512 5360
3	ILMN_125 8600 NA	Acaca	acetyl- Coenzym e A carboxyla se alpha	ENSMUS G0000002 107476 0532
4	ILMN_273 8082 NA	Adipoq	adiponecti n, C1Q and collagen domain containing	ENSMUS G0000002 11450 2878