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# **Modeling toxicodynamic effects of trichloroethylene on liver in mouse model of autoimmune hepatitis**

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# **Abstract**

Chronic exposure to industrial solvent and water pollutant trichloroethylene (TCE) in female MRL +/+ mice generates disease similar to human autoimmune hepatitis. The current study was initiated to investigate why TCE-induced autoimmunity targeted the liver. Compared to other tissues the liver has an unusually robust capacity for repair and regeneration. This investigation examined both time-dependent and dose-dependent effects of TCE on hepatoprotective and proinflammatory events in liver and macrophages from female MRL+/+ mice. After a 12-week exposure to TCE in drinking water a dose-dependent decrease in macrophage production of IL-6 at both the transcriptional and protein level was observed. A longitudinal study similarly showed that TCE inhibited macrophage IL-6 production. In terms of the liver, TCE had little effect on expression of pro-inflammatory genes (*Tnfa*, *Saa2* or *Cscl1*) until the end of the 40-week exposure. Instead, TCE suppressed hepatic expression of genes involved in IL-6 signaling (*Il6r*, *gp130*, and *Egr1*). Linear regression analysis confirmed liver histopathology in the TCE-treated mice correlated with decreased expression of *Il6r*. A toxicodynamic model was developed to estimate the effects of TCE on IL-6 signaling and liver pathology under different levels of exposure and rates of repair. This study underlined the importance of longitudinal studies in mechanistic evaluations of immuntoxicants. It showed that later-occurring liver pathology caused by TCE was associated with early suppression of hepatoprotection rather than an increase in conventional pro-inflammatory events. This information was used to create a novel toxicodynamic model of IL-6-mediated TCE-induced liver inflammation.

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#### **Keywords**

Trichloroethylene; liver; IL-6; immunotoxicity; modeling

# **Introduction**

Trichloroethylene (TCE) is a chlorinated hydrocarbon that has been used as a degreasing agent since the 1920s. Because of inappropriate disposal over the years TCE is now a common pollutant at Superfund toxic waste sites and at many industry and government facilities. It is found in soil and surface water as a result of direct discharges and in groundwater due to leaching from disposal operations. As noted by a recent National Research Council report evidence on human health hazards from TCE exposure, either occupational or environmental, has strengthened in recent years (Committee on Human Health Risks of Trichloroethylene, 2006). One of the predominant non-cancer outcome associated with TCE exposure in humans is immunotoxicity, most notably the development of hypersensitivity responses including autoimmune scleroderma and autoimmune liver diseases (Byers *et al.*, 1988; Yanez Diaz *et al.*, 1992; Hansen *et al.*, 1988; Saihan *et al.*, 1978; Flindt-Hansen *et al.*, 1987; Czirjak *et al.*, 1994; Lockey *et al.*, 1997; Dubrow *et al.*, 1987; Gist *et al.*, 1995). TCE is still widely used in Asia, where it has become a serious work-related health concern due to the induction of dermal and systemic hypersensitivity diseases often accompanied by non-viral, immune-mediated hepatitis (Kamijima *et al.*, 2008).

Using a mouse model we and others have found that long-term exposure to TCE in drinking water at concentrations lower than sanctioned occupational exposure levels generated a T cell-mediated liver disease commensurate with human idiopathic autoimmune hepatitis (AIH)(Griffin *et al.*, 2000; Gilbert *et al.*, 2008; Cai *et al.*, 2008). This TCE-induced liver inflammation was not usually accompanied by markers of acute liver injury such as increased blood levels of alanine transaminase or liver fibrosis, but was associated with the development of antibodies specific for liver microsomal proteins similar to those in patients with type 2 AIH.

The development of toxicant-induced immune pathology such as the autoimmune hepatitis caused by TCE exposure is almost certainly a complex multifactorial process. Developing conceptual models can be a way to delineate and quantify the contribution of different toxicant-induced alterations to the actual pathology. As a first step in this direction a model was developed here to describe a specific part of the process, namely IL-6-mediated liver events. IL-6 is one of the most important regulators of hepatic inflammation. The pathogenesis of AIH requires circumvention of the well-known propensity of the liver to induce T cell tolerance (Carambia *et al.*, 2010). Pre-existing inflammation in the liver may subvert its tolerogenicity and help sustain an immune response by entering T cells (Crispe, 2009). The ability of toxicant exposure to generate such inflammation depends on opposing forces of tissue injury and tissue repair. Distress signals triggered during initiation of toxicant-induced liver injury (e.g. lipid peroxidation, reactive intermediate formation) can promote inflammation. However, they also stimulate protective (anti-apoptotic) and

regenerative (cell division) mechanisms in the liver. One of the mechanisms that determine whether toxicant exposure ultimately leads to tissue repair or to injury-induced inflammation is regulated by IL-6.

Treatments to prevent or reverse immunological liver injury in mouse models have been associated with an increase in liver expression of *Il6* (Liu *et al.*, 2006). Disruption of IL-6, or its receptors IL-6R or Gp130, has been shown to promote liver inflammation and/or mortality following partial hepatectomy (Wuestefeld *et al.*, 2003), ethanol-induced liver disease (Gao, 2012), carbon tetrachloride-induced liver necrosis (Bansal *et al.*, 2005), obesity-associated insulin resistance (Wunderlich *et al.*, 2010), autoimmune cholangitis (Zhang *et al.*, 2010), and Con A-induced hepatitis (Lutz *et al.*, 2012). Thus, IL-6 appears to prevent immunological liver injury. In addition to its documented ability to promote liver regeneration and/or protection in the face of damage or trauma IL-6 also appears to be required for normal liver maintenance. Liver weight and total DNA and protein contents were decreased 26–28% in older (5–10month-old) female IL-6-deficient mice as compared to age-matched wild-type controls (Wallenius *et al.*, 2001). This suggests that IL-6 is needed for normal hepatocyte turnover, and that over time a loss of this cytokine is detrimental to liver function.

In an attempt to define why TCE-induced autoimmunity targets the liver, mice exposed to a single dose of TCE for 4, 10, 16, 22, 28, 34 or 40 weeks were evaluated in the current study for time-dependent alterations in IL-6 as well as other pro-inflammatory mediators. This was complemented by a second study that examined the dose-dependent effects of TCE on these mediators at a single time point.

The development of autoimmune hepatitis in our mouse model of TCE exposure involves alterations in both the liver and the immune system. This multi-factorial process mimics the complicated etiologies of human autoimmune diseases. Developing conceptual models can be a way to delineate and quantify the contribution of different disease-induced alterations to actual pathology. As a first step in this direction the results obtained here were used to model the part of the TCE-induced disease process revealed in the current study, namely the effect of TCE on IL-6-mediated liver events. Taken together, the results suggest that lateroccurring TCE-induced liver damage was due to an early decrease in IL-6-mediated hepatoprotection rather than an increase in pro-inflammatory events.

# **Materials and Methods**

#### **Mouse treatment**

Eight week-old female MRL+/+ mice (Jackson Laboratories; Bar Harbor, ME) were housed in polycarbonate ventilated cages and provided with drinking water (ultrapure from Milli-Q Integral Water Purification System, Millipore) *ad libitum*. TCE (purity 99± %; Aldrich Chemical Co. Inc.; Milwaukee, WI) was suspended in drinking water with 1% emulsifier Alkamuls EL-620 from Rhone-Poulenc (Cranbury, NJ). Freshly made TCE-containing drinking water was provided every 2–3 days. In one experiment the mice (12 mice/group) received either 0, 0.02, 0.1 mg/ml or 0.5 mg/ml TCE in their drinking water for 12 weeks. In a second experiment the mice (8–9 mice/group) received 0 or 0.5 mg/ml TCE in their

drinking water for 4, 10, 16, 22, 28, 34 or 40 weeks. The mice were weighed weekly and water consumption was monitored. All studies were approved by the Animal Care and Use Committee at the University of Arkansas for Medical Sciences. When the mice were sacrificed at the different time points adherent macrophages isolated from pooled peritoneal exudates from 2–3 mice (n=3–4/treatment group) were incubated for 20 hours in the presence or absence of LPS (1 μg/ml). Approximately 80% of adherent peritoneal exudate cells (PEC), regardless of treatment group, expressed the transmembrane protein F4/80, a marker of mature macrophages (data not shown). Culture supernatants from the peritoneal macrophages were then collected for cytokine evaluation. RLT Lysis Buffer (Qiagen Sciences, Germantown, MD) was then added directly to the remaining adherent cells before freezing for subsequent qRT-PCR analysis. Liver tissue harvested at the time of sacrifice was stained with H&E and examined for liver pathology. Liver and sections were examined microscopically and scored in a blinded manner by a veterinary pathologist for the severity of inflammation and fibrosis based on a four point scale  $(0 - 3)$ , ranging from no change to severe, respectively) as described (Gilbert *et al*., 2008). Some liver tissue was instead frozen for subsequent qRT-PCR analysis.

#### **qRT-PCR**

Fluorescence-based quantitative reverse transcriptase polymerase chain reaction (qRT-PCR) was conducted using RNA isolated (using RNeasy, Qiagen, Germantown, MD) from peritoneal macrophages or liver. Details of the technique as well as quality controls and rationale for choice of reference gene *Eef2* (eukaryotic translation elongation factor 2) has been described (Gilbert *et al*., 2008).

#### **Cytokine Analysis**

The macrophage culture supernatants were examined using the Mouse IL-1β, IL-6 and TNF alpha Ready-Set-Go! ELISA kits from eBioscience, Inc. (San Diego, CA). The amounts of IL-6 produced by macrophages in the 12-week study were generally lower than those generated in the 40-week study. This is likely due to difference in the number of macrophages/well. In the 12-week study pooled peritoneal cells from 3 mice were used to generate four 1ml wells (two unstimulated and two LPS-treated) for an n=4 per treatment group. With 8–9 mice/group in the 40-week study pooled peritoneal cells from 2–3 mice were used to generate two 1ml wells (one unstimulated and one LPS-treated) for an n=3 group.

Levels of IL-6R in the livers of individual mice were examined by Western blotting using goat IgG anti-mouse IL-6Rα (R&D Systems, Minneapolis, MN). The results were represented as mean  $\pm$  standard deviation of the densitometric analysis of the IL-6R compared to reference protein GAPDH (glyceraldehyde 3-phosphate dehydrogenase) run in the same lane, and detected by rabbit IgG anti-GAPDH (Trevigen, Gaithersberg, MD).

**Antibody Production—**Using previously described methodology (Gilbert *et al*., 2008), microsomal liver protein (30 μg) obtained from an untreated MRL+/+ mouse was separated on 12% SDS-PAGE, electrotransferred onto nitrocellulose, and subsequently probed with pooled sera (1:500) obtained from control or TCE-treated MRL+/+ mice followed by HRP-

conjugated polyclonal goat anti-mouse IgG (1:4,000). Densitometric analysis of mouse myeloma IgG run in adjoining lanes was used to normalize exposure times for the individual Western blots.

#### **Statistics**

The data are presented as means  $\pm$  standard deviations. Assays were conducted using samples from 8–12 individual mice per treatment group or samples from equal numbers of pooled cells for n=3 or 4 per treatment group. The threshold for statistical significance was set at  $a = 0.05$ . Differences between experimental groups were tested first with analysis of variance (ANOVA), and where the *F* test was significant, subsequent pairwise contrasts were tested using a two-sample *t*-test. Homogeneity of variance between groups was tested using studentized Breusch-Pagan, and normality of residuals using Shapiro-Wilk. Where significant deviations from homoscedasticity or normality were observed, the nonparametric Kruskal-Wallis and Wilcoxon rank sum tests were applied instead of ANOVA/ttest. For experiments involving multiple timepoints, ordinal logistic regression models were fitted and the significance of TCE exposure was computed using a likelihood-ratio test; this is a two-factor generalization of Kruskal-Wallis. Linear regressions were fitted to evaluate the relationship between gene expression and histopathology, with statistical significance judged using an *F* test. Macrophage concentration and gene expression values were rightskewed, and therefore these data were log-transformed for statistical analyses.

#### **Mechanistic Toxicodynamic Modeling**

The effects of IL-6 signaling on liver events in TCE-treated mice were modeled to link changes in TCE mediated IL-6 signaling outcomes to the observed pathology following lowdose chronic exposure to TCE. The objective in developing a mathematical model for chronic, low dose exposure to TCE was to provide a means to quantitatively describe the role of IL-6 as a maintenance mechanism and predict downstream effects, such as changes in pathology, due to modifications of this IL-6 repair pathway. To this end, a time-dependent mathematical description of the health state of discrete volumes of liver ("liver units") and the IL-6 and TCE-dependent transition between these health states was developed. Using *in* vivo results reported in this study, this toxicodynamic model will create a link between TCE exposure and the resulting histopathology. While not measurable *in vivo*, the state and number density of individual liver units serve as an intermediate measure to quantify the relationship between impaired cytokine signaling and the resulting autoimmune hepatitis.

**Liver unit health state model—**For the purposes of mathematical modeling, the characteristics of the liver units (LUs) were as follows: the entire liver comprises LUs, each of which is of equal volume; an LU is relatively small in volume compared to that of the entire liver, but consists of a large enough number of cells to be represented as a continuum; each LU exists in one of three health states: healthy (H), compromised (C), and inflamed (I), and is homogeneous with respect to its properties and health state; and the health state of the entire liver may be estimated through a number-weighted average of the health states of the constituent LUs. Table 1 lists the characteristics and assumptions for each mathematical state.

The resulting transition between health states is described by

$$
H \overset{k_{DAM,H-C}}{\underset{k_{REP,H-C}}{\rightleftarrows}} C \overset{k_{DAM,C-I}}{\underset{k_{REP,C-I}}{\rightleftarrows}} I, \quad (1)
$$

where the k's represent transition rates, the subscripts *DAM* and *REP* refer to damage- and repair-associated phenomena, respectively, and the subscripts H-C and C-I refer to their respective transition pathways.

The corresponding system of differential equations governing the time-dependent fraction of LUs in each state may be written as

$$
\frac{d[H]}{dt} = -k_{DAM,H-C} [H] + k_{REP,H-C} [C]
$$
\n
$$
\frac{d[C]}{dt} = k_{DAM,H-C} [H] + k_{REP,C-I} [I] - (k_{REP,H-C} + k_{DAm,C-I}) [C]. \quad (2)
$$
\n
$$
\frac{d[I]}{dt} = -k_{REP,C-I} [I] + k_{DAM,C-I} [C]
$$

Here, *t* is time and [*H*], [*C*], and [*I*] are the fractions of LUs in the healthy, compromised, and inflamed states, respectively. It is assumed that initially  $(t=0)$ ,  $[H]=1$  and  $[C]=[I]=0$ .

To complete the mathematical description of this system, four major assumptions were made:

- **1.** LUs normally exist in a state of IL-6-mediated hepatocyte turnover and protection.
- **2.** Events such as TCE exposure can initiate inflammatory processes and move the LUs into the "C" state. However, protective mechanisms mediated by IL-6 normally restore the LUs from the "C" to the "H" state. The rates of repair, *kREP,H-C* and *kREP,C-I*, are dependent on the fraction of IL-6 produced from the macrophage and the IL-6r expressed by the hepatocyte. If homeostasis levels of IL-6 and IL-6r are present, these pathways operate at the optimal repair rates.
- **3.** TCE initiates inflammatory processes (e.g. redox disequilibrium) that move the LUs from the "H" to "C" state. It also decreases the protective effects of IL-6 signaling that would normally restore the LUs to the "H" state. These dual effects of TCE allow the inflammatory processes to progress and to move the LUs from the "C" to the "I" state.
- **4.** Autoimmune hepatitis pathology does not occur without TCE, even if the IL-6 pathway is impaired in some other way.

Consistent with these assumptions, the rate terms above can be further specified as follows:

$$
k_{REP.H-C} = k_1 \cdot f_{IL6} \quad ; \quad k_{REP.C-I} = k_2 \cdot f_{IL6} \quad ; \quad k_{DAM.H-C} = k_3 \quad ; \quad k_{DAM.C-I} = k_4 \cdot f_{ICE}, \quad (3)
$$

where  $k_1$ ,  $k_2$ ,  $k_3$ , and  $k_4$  are constants to be estimated using experimental data,  $f_{\text{TCE}}$ represents the administered TCE dose normalized by the upper dose used in this study (0.5 mg/ml or a time-weighted average of approximately 54 mg/kg/day), and  $f_{IL6}$  is the fraction

of IL-6 expressed by the macrophage compared to control levels. To express the dependence of *f*IL6 on TCE dose, a sub-model based on a saturation mechanism was used:

$$
f_{IL6}=1-\frac{\alpha \cdot f_{TCE}}{\beta + f_{TCE}},
$$
 (4)

where  $\alpha$  and  $\beta$  are constants to be derived from experimental data.

**Predicting liver pathology scores—**To compute overall liver pathology scores, the [H], [C], and [I] calculated from equations (2), (3), and (4) at the desired time point were used as weighting factors for the individual PS values corresponding to each of the model states. Mathematically, this can be expressed as

$$
PS = \sum_{s=H,C,I} PS_s \cdot [s], \quad (5)
$$

where  $PS_s$  is the pathology score of a LU in state s (see Table 1).

**Software and modeling tools—**The system of differential equations were solved using a fourth-order Runge-Kutta method implemented in the Python programming language  $(v2.7.6)$  [\[https://www.python.org/\]](https://www.python.org/). Parameter estimation was conducted using lsqfit  $(v4.6.1)$ ] [\[https://github.com/gplepage/lsqfit\]](https://github.com/gplepage/lsqfit), a software package for non-linear least-squares fitting of noisy data.

#### **Results**

#### **Dose-dependent effects of TCE on peritoneal macrophage activity**

Since autoimmune diseases and hypersensitivity disorders in humans involve an ill-defined genetic component, we use young "autoimmune-prone" female MRL+/+ mice to study the immunotoxicity of TCE. As observed previously, TCE exposure did not alter weight gain or water consumption (data not shown). Peritoneal macrophages from the mice exposed to different concentrations of TCE for 12 weeks were examined for the production of macrophage-derived cytokines IL-6 and IL-1β. Macrophage secretion of IL-1β was unchanged by exposure to TCE (Figure 1). The peritoneal macrophages collected from control mice secreted low but measurable levels of IL-6 even in the absence of LPS. Stimulation with LPS increased IL-6 production in all groups. However, both LPSdependent and LPS-independent IL-6 production was suppressed in a dose-dependent manner in peritoneal macrophages from mice treated for 12 weeks with TCE. For example, LPS-induced IL-6 production in mice exposed to 0.5 mg/ml TCE was 70% lower than that of controls.

IL-6 was also inhibited at the transcriptional level in macrophages from TCE-treated mice (Figure 2). Although LPS stimulation increased *Il6* expression, this effect was significantly suppressed in macrophages from mice treated with 0.1 or 0.5 mg/ml TCE as compared to control mice. Once again the suppressive effects of TCE were confined to IL-6, and did not encompass expression of genes for other macrophage-derived cytokines, including Lt-α,

IL-12, or IL-10. Taken together, a 12-week exposure to TCE selectively suppressed IL-6 gene expression and protein production by peritoneal macrophages in a dose-dependent manner. The ability of TCE to alter expression of genes for other macrophage-derived cytokines was intermittent and not dose-dependent.

#### **Time-dependent effects of TCE on peritoneal macrophage gene expression**

In a second study designed to examine time-dependency of TCE-induced effects mice were given drinking water alone or with 0.5 mg/ml TCE for 4, 10, 16, 22, 28, 34 or 40 weeks. TCE exposure did not alter the number of PEC recovered at any of the time points (data not shown). Once again TCE suppressed production of IL-6 (Figure 3). Also evident, but as yet unexplained, was the general time-dependent decrease in IL-6 production in both treatment and control groups. Production of TNF-α was not affected by TCE exposure. A longitudinal evaluation of cytokine gene expression showed that the TCE-induced decrease in *Il6* expression by peritoneal macrophages was evident by 16 weeks of exposure (Figure 4). The time-dependent expression of several other genes for macrophage-derived cytokines, *IL1b, Il12,* and *Mmp12* was for the most part unaltered by exposure to TCE (Figure 4 and data not shown). Thus, the primary effects of exposure to TCE on peritoneal macrophages was a decrease in *Il6* that was maintained for the duration of the study.

#### **Time-dependent effects of TCE on liver events**

Most of the protective and/or regenerative events in T cell-mediated liver injury are triggered by IL-6 signaling that is initiated when IL-6 binds to a complex comprised of the transmembrane protein gp130 and the IL-6R on hepatocytes (Klein *et al.*, 2005). As shown in Figure 5 hepatic expression of *Il6r* was suppressed by TCE at several time points, and only approached control values at the last time point. Protein levels of IL-6R were also lower in the livers of the TCE-treated mice. The gene that encoded for the other subunit in the IL-6R family, *Gp130,* was suppressed by TCE at early time points. Expression of IL-6 itself in the liver was undetectable (data not shown). Another molecule important in hepatoprotection is the transcription factor EGR-1. EGR-1 binds to the promoter region of *Il6* (Hoffmann *et al.*, 2008), and reciprocally, is important in mediating signaling from the IL-6R/STAT3 pathway (Pritchard *et al.*, 2011). Expression of *egr1* in the liver was suppressed midway through the TCE exposure, but then rebounded at the final 40-week time point. Increased levels of pro-inflammatory cytokines/chemokines such as TNF-α, osteopontin, serum amyloid A (SAA) and CXCL1 have been implicated in the induction or progression of chronic liver inflammation (Iwamoto *et al.*, 2013; Nagoshi, 2014; Gollaher *et al.*, 1990; Zhang *et al.*, 2012). Hepatic expression of these *Saa2*, *Cxcl1* and *Spp1* (encodes for osteopontin) were for the most part unchanged or decreased during all but the last 40 week time point of TCE exposure. Thus, unlike IL-6R associated genes hepatic expression of several pro-inflammatory cytokines and chemokines was primarily unchanged or decreased by TCE exposure until the last time point when expression was dramatically reversed in select TCE-treated mice. These results showed that during most of the exposure TCE appeared to negatively impact liver repair rather than directly promote inflammation. Only at the last time point was this reversed; several pro-inflammatory cytokines/ chemokines increased expression while the negative effect on hepatoprotective genes was overturned.

Histopathology in the form of lymphoplasmacytic portal infiltrate and lobular inflammation in the liver was not noted until week 28 of TCE exposure, and became more robust during the course of the 40-week experiment (Figure 6A). This pathology was characteristic of the early stages of autoimmune hepatitis; hepatocellular necrosis was only noted in a couple of instances. The mice were also examined for the generation of anti-liver antibodies as another readout of immune-mediated liver disease (Figure 6B). MRL+/+ mice are noted for their age-dependent increase in the production of autoantibodies such as anti-nuclear antibodies, even in the absence of toxicant exposure (Yoshida *et al.*, 1989). In accord with this predisposition the baseline production of anti-liver antibodies became more abundant in

control mice at the 40 week time point. However, exposure to TCE further increased the levels and diversity of the anti-liver antibodies. Thus, the MRL+/+ mice treated with TCE for 40 weeks demonstrated liver inflammation and anti-liver autoantibody production consistent with AIH.

To help determine functional relevance hepatic gene expression in individual mice at the 40 week time point were plotted against liver pathology scores in the same mice. The linear regression showed that liver pathology was most closely correlated with a decrease in *Il6r* (p=0.003)(Figure 6C). Correlations between liver histopathology and expression of *Egr1* and *Spp1* were also moderately significant  $(p<0.07)$ . There was no significant correlation between liver histopathology and expression of *Tnfa* or *Cxcl1*. This evaluation suggested that liver pathology was most closely correlated with a decrease in IL-6 signaling as reflected in a decrease in the expression of *Il6r* and *Egr1* in the liver. There was also correlation with expression of *Spp1,* suggesting that the reversal of the TCE-induced decrease in OPN observed in the liver played a role in liver pathology.

#### **Toxicodynamic model for liver response to TCE exposure**

In order to develop a model to describe the effect of TCE on IL-6-mediated liver events certain required parameters were estimated based on the results described above.

**Parameter estimation—In** order to fit a curve that could be used to extrapolate IL-6 effects across a range of TCE doses values of  $a$  and  $\beta$  in the IL-6 submodel, Eq. (4), were estimated using a nonlinear least-squares approach with the non-LPS induced IL-6 results presented in Fig. 1. The resulting parameters values, mean (variance), were found to be  $\alpha$  = 1.01 (0.01) and  $\beta$  = 0.071 (0.003). Figure 7A illustrates the resulting fit of the experimental data to the IL-6 submodel.

It was similarly necessary to fit a curve to extrapolate liver pathology based on time of TCE exposure. The rate constants,  $k_i$ , defined in Eq. (3), were estimated based on experimental time-course pathology scores (Figure 6A) to be  $k_1 = 101.5$  (98.0),  $k_2 = 0.39$  (0.18),  $k_3 = 1.02$  $(0.08)$ , and  $k_4 = 0.21$  (0.16). The resulting fit of the data to the mathematical model is depicted in Figure 7B. The uncertainty shown in model simulations results from both the uncertainty in the parameters associated with the IL-6 submodel and that from *in vivo* pathology scores.

**Simulations of liver unit health states and the dose response—**Following parameter estimation, simulations of time-course LU health were conducted. Figure 8

illustrates results from several such studies, where the fraction of LUs in a particular health state are shown as a function of time at the two highest doses used in the experimental study. For the 0.1 mg/ml dose (Figure 8A), almost all of the LUs are in a healthy state. However, as the external TCE dose is increased to 0.5 mg/ml (Figure 8B), the abundance of healthy LUs decreases while those in the compromised/inflamed state increase in a non-linear manner. At doses less than 0.1 mg/ml, simulations indicated that virtually all of the LUs were in the healthy (H) state over time.

One of the benefits of the mathematical model is the prediction of system variables and endpoints not directly measured during the course of the studies. For instance, using 40 week pathology scores as an endpoint, the model was used to generate a dose-response curve (Figure 9). This curve can be used to relate this endpoint to any dose within the predicted range. As an example, for a benchmark response level (BML) corresponding to mild inflammation of 25% or less of the portal regions of the liver  $(PS=2)$ , the benchmark dose (BMD) was estimated to be *fTCE* = 0.55, corresponding to a 0.27 mg/ml dose, or approximately 37 mg/kg/day, of continuous TCE exposure.

**Examining the impact of varying relative rates of damage and repair—**Another important benefit to the mathematical modeling is the ability to vary system parameters and observe the effects on system states of interest. Here, the effects of varying the relative rates of damage and repair in the H-C and C-I state transitions were investigated through a parametric study. Since each damage and repair pathway is first order with respect to the abundance of LUs, ratios of rate constants were defined:

$$
\kappa_{H-C} = \frac{k_{REP,H-C}}{k_{DAM,H-C}} \quad ; \quad \kappa_{C-I} = \frac{k_{REP,C-I}}{k_{DAM,C-I}}. \quad (1)
$$

By conducting simulations with varying values for  $\kappa$  (Figure 10), the effects of relative rates of repair and damage in the system could be examined. If  $\kappa \gg 1$ , the repair mechanism dominates and the LUs tend toward a relatively low value of PS, even at higher TCE dose; conversely, if  $\kappa \ll 1$ , the damage mechanism dominates for the given pathway and LUs may acquire high values of PS values, even at relatively low doses. As can be seen, the shape and nonlinearity of the dose-response curve is highly dependent on the relative rates of repair and damage in both of the health state transitions. Overall, such predictions can help in the understanding of the interactions in this system and lend insight into the effects of non-TCE mediated events, such as additional stress from other hepatotoxicants or further impaired IL-6 repair mechanisms.

#### **DISCUSSION**

MRL+/+ mice can spontaneously develop autoimmune diseases such as lupus nephritis, pancreatitis, and Sjogren's syndrome late in life (after 1-year of age)(Kanno *et al.*, 1992; Toda *et al.*, 1999). However, before they reach one year of age most female MRL+/+ mice do not exhibit autoimmune tissue pathology, and are often used to examine the autoimmunepromoting capacity of a toxicant such as TCE. Based on water consumption and TCE

degradation in the water bottles, the mice given water containing TCE at 0.02, 0.1 or 0.5 mg/ml for 12 weeks were exposed to TCE at time-weighted levels of approximately 3, 14 or 64 mg/kg/day, respectively. Even the highest exposure is lower than the current 8-hour Permissible Exposure Limit [established by the Occupational Safety and Health Administration (OSHA)] for TCE of 100 ppm or approximately 76 mg/kg/day.

The effects of TCE on macrophage activity have been primarily studied in inhalation models. Inhalation exposure to TCE was shown to increase susceptibility to respiratory bacterial infection in mice, and to suppress phagocytosis in lung macrophages (Selgrade *et al.*, 2010). Similarly, multiple inhalation exposures to TCE lowered resistance to respiratory streptococcus infection (Aranyi *et al.*, 1986). Although the mechanism for this suppressive effect of inhaled TCE on macrophages was not defined, others have shown that an IL-6 deficiency increases susceptibility to viral and bacterial respiratory infections (Murphy *et al.*, 2008; Jones *et al.*, 2006). The results of the current study showed that oral exposure to TCE suppressed IL-6 at the level of protein production and gene expression in macrophages.

IL-6 is a pleiotropic cytokine, which can make it difficult to predict the cumulative impact of its altered production. Elevated levels of IL-6 in the blood have been observed in a number of pathological conditions associated with chronic inflammation including rheumatoid arthritis (Gottenberg *et al.*, 2012), systemic lupus erythematosus (Chun *et al.*, 2007), and active disease in Guillain-Barre syndrome (Weller *et al.*, 1991). IL-6 did not reach detectable levels in the blood of control or TCE-treated mice in the current study. Circulating levels of IL-6 are increased in children with AIH type 1, but not with AIH type 2 (Maggiore *et al.*, 1995), the type of AIH that most closely resembles TCE-induced disease in MRL+/+ mice. Some studies of idiopathic autoimmune liver disease in humans have found increased levels of IL-6 in liver biopsies (Zhao *et al.*, 2011), while other studies of autoimmune hepatitis have demonstrated decreased expression of hepatic *Il6* in the liver (Tovey *et al.*, 1991). On the other hand, treatments to prevent or reverse immunological liver injury in mouse models have been associated with an increase in liver expression of *Il6* (Liu *et al*., 2006). Thus, the majority of studies suggest that in the liver IL-6 is primarily protective. Increases in hepatic levels of IL-6 in some humans with AIH may represent a compensatory rather than pathological mechanism. Alternatively, changes in IL-6 may be specific for a certain stage of disease development, type of autoimmune hepatitis (e.g. type 1 vs type 2) (Maggiore *et al*., 1995), or cell type (e.g. peritoneal exudate macrophages vs Kupffer cells). Deletion of IL-6 in mice deficient in TGF-β receptor II improved colitis but exacerbated autoimmune cholangitis in association with increased numbers of activated T cells (Zhang *et al*., 2010).

Cytokine production by macrophages from MRL+/+ mice is reportedly aberrant even in the absence of TCE exposure. LPS-induced production of IL-6, IL-1β, TNF-α, and IL-12 by macrophages from untreated MRL+/+ mice were all significantly decreased in comparison to macrophages from C57BL/6, BALB/c or A/J mice(Hartwell *et al.*, 1995; Alleva *et al.*, 2000). Of these macrophage-derived cytokines only IL-6 was found in the present study to be further decreased by TCE exposure.

In addition to a decrease in macrophage-derived IL-6, TCE suppressed liver expression of Il6r and gp130, the dual components of the IL-6R. This TCE-induced decrease would seem to further ensure the lack of IL-6 signaling in the liver. The IL-6-induced liver protection to T cell-mediated liver injury has been attributed to a downstream increase in acute phase protein serum amyloid A2, (SAA2)(Klein *et al*., 2005). TCE suppressed hepatic expression of *Saa2* at two time points late in the exposure period, thus seeming to prevent the upregulation of this molecules needed for liver regeneration. Egr1 is a transcription factor required for wound healing, and which has been identified as a negative regulator of carbon tetrachloride-induced hepatotoxicity (Pritchard *et al.*, 2010). Egr1 has been described as both a trigger and a target for IL-6 (Zhang *et al.*, 2013; Maekawa *et al.*, 2010). Only at the final time point did TCE increase expression of *Egr1* and *Saa2*. It is not known why the earlier TCE-induced suppression was reversed, but presumably the late recovery of these genes was not sufficient to protect against liver damage.

The contribution of TCE to AIH in the present model is multidimensional; the healthy-toinflamed state model described here can be amended to include more immune parameters such as the contribution of  $CD4^+$  T cells as they are characterized. However, even in its present state, the model facilitated point-of-departure predictions based on dose-dependent changes in liver pathology. The model stemmed from the linear regression analyses showing that liver pathology in TCE-treated mice was best correlated with the decreased liver expression of macrophage *Il-6*r. We now have the tools to predict liver pathology based on relative rates of liver repair and damage. In addition to its predicted effect on IL-6 signaling the model also infers that TCE initiates inflammatory processes that transition LUs from "H" to "C". These processes were not investigated in this study, but probably include, but are not restricted to, alterations in redox equilibrium. In a previous study, a metabolomics analysis following chronic 32 week exposure to 0.5 mg/ml in MRL+/+ mice revealed significant alterations in several metabolites (e.g., cystathionine) involved in the generation of glutathione, which functions as the major intracellular antioxidant against oxidative stress and plays an important role in the detoxification of reactive oxygen species and subsequent oxidative damage from pro-oxidant environmental exposures. Others have shown the functional significance of oxidative stress in TCE-induced liver pathology (Wang *et al.*, 2007; Wang *et al.*, 2013). IL-6 has been shown to inhibit oxidative stress and steatosis in the liver (El-Assal *et al.*, 2004). Consequently, a TCE-induced loss of IL-6 signaling in the liver would be expected to exacerbate associated oxidative-stress and resulting inflammation. The first stage model development described here (i.e. generation of equations and description of parameters) was based on data from two different experiments, albeit with some differences in experimental design. Obtaining new data to validate and extend this model will be included in the design of future chronic TCE exposure studies.

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# **Abbreviations**

**TCE** trichloroethylene

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# **Highlights**

- **•** We developed a toxicodynamic model to study effects of trichloroethylene on liver.
- **•** We examined protective as well as pro-inflammatory events in the liver.
- **•** Trichloroethylene inhibits IL-6 production by macrophages.
- **•** Trichloroethylene inhibits components of the IL-6R in the liver.
- **•** Trichloroethylene inhibits events associated with IL-6-mediated hepatoprotection.



#### **Figure 1. TCE inhibits macrophage IL-6 production in dose-dependent manner**

Peritoneal macrophages were incubated with (open bars) or without LPS (shaded bars) following isolation from untreated control mice or from mice exposed to TCE at different concentrations for 12 weeks. Culture supernatants were examined for cytokines (mean ± SD). \*Significantly different ( $\alpha$ <0.05) compared to control values.



#### **Figure 2. TCE inhibits macrophage** *Il6* **expression in dose-dependent manner**

Cytokine gene expression was examined in peritoneal macrophages incubated with (open bars) or without LPS (shaded bars) after isolation from untreated control mice or from mice exposed to TCE for 12 weeks. The data represents the mean  $\pm$  SD. \*Significantly different  $(\alpha<0.05)$  compared to control values.





Peritoneal macrophages were incubated with LPS following isolation from untreated control mice or from mice exposed to TCE (0.5 mg/ml) for up to 40 weeks. Culture supernatants were examined for cytokines (mean  $\pm$  SD). \*Significantly different ( $\alpha$ <0.05) compared to control values.



#### **Figure 4. TCE inhibition of** *Il6* **expression is maintained over time**

Cytokine gene expression was examined in peritoneal macrophages incubated with or without LPS after isolation from untreated control mice or from mice exposed to TCE (0.5 mg/ml) for up to 40 weeks. The data represents the mean  $\pm$  SD. \*Significantly different (α<0.05) compared to control values.



## **Figure 5. TCE alters expression of hepatic genes over time**

A. Gene expression in individual liver tissue isolated from untreated control mice or from mice exposed to TCE (0.5 mg/ml) for up to 40 weeks. The data represents the mean  $\pm$  SD from 6–8 individual mice/treatment/time point. \*Significantly different (α<0.05) compared to control values. B. Relative protein levels (percentage reference protein GAPDH) of IL-6R in individual livers from untreated control mice or mice exposed to TCE (0.5 mg/ml) for 16 weeks (mean  $\pm$  SD).



#### **Figure 6. TCE liver pathology correlates with loss of hepatic** *Il-6r* **expression**

A. Liver pathology based on immune cell infiltration and inflammation was assessed in mice exposed to TCE (0.5mg/ml) for 28, 34 or40 weeks. B. Equal amounts of liver protein from an untreated mouse were separated in 4 lanes of SDS-PAGE, each of which were immunoblotted with pooled sera obtained from control MRL+/+ mice or mice treated with 0.5 mg/ml TCE for 4 or 40 weeks. C. Hepatic gene expression in from mice exposed to TCE (0.5 mg/ml) for 40 weeks was plotted against liver histopathology in the same mice. Gene expression values are shown in log scale because of right skewness. Regression p-values were computed using an F test of the null hypothesis of horizontal slope.



#### **Figure 7. Submodel for parameter estimation**

A. An IL-6 submodel was developed for estimating dose-dependent reduction in the fraction of IL-6 expressed by the macrophage. Points and error bars represent data and uncertainty, while solid and dashed lines are the mean and 95% confidence intervals from model predictions. B. Time-course pathology scores were used to extrapolate liver pathology based on time of TCE exposure. Points and error bars represent data and uncertainty, while solid and dashed lines are the mean and 95% confidence intervals from model predictions.



#### **Figure 8. Liver unit state predictions based on the model**

Fraction of liver units in each state for the 0.1 (A) and 0.5 (B) mg/ml experimental doses. This reflects the overall health of the liver rather than specific changes in IL-6 production. Solid lines represent the H state, while vertical and dashed lines correspond to the C and I states, respectively.



#### **Figure 9. Dose response curve for current study**

Predicted dose response curves for pathology scores (PS) 40 weeks post TCE exposure. The mean values and 95% confidence intervals are shown as solid and dashed lines, respectively. The point represents the value of the benchmark dose (BMD) corresponding to the benchmark response level (BMR) described in the text.



**Figure 10. Simulations illustrate the effects of relative rates of repair and damage on liver damage**

Model predictions for varying relative levels of repair in (A) the H-C pathway ( $\kappa_{\text{H-C}}=[1, 10, 10]$ 100, 1000],  $\kappa_{C-I}$ =1) and (B) the C-I pathway ( $\kappa_{H-C}$ =100,  $\kappa_{C-I}$ =[0.1, 1, 10, 100]).  $\kappa_{H-C}$ =100 and  $\kappa_{C-I}$ =1 are the values that that relate to the pathology scores from the current study. The two dotted lines represent the kappas (*k*) used in this study (*k*1=100 and *k*2=1). Dashed lines are representative of hypothetical pathways with an order of magnitude increase in repair mechanisms. Conversely, solid lines represent scenarios where external factors result in an order of magnitude decrease in repair.

#### **Table 1**

# Description of LU states

