Topical application of the anti-microbial chemical triclosan induces immunomodulatory responses through the S100A8/A9–TLR4 pathway

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Introduction

Triclosan is a low molecular weight anti-microbial found in numerous consumer products such as hand soaps, lotions, and toothpaste, and is incorporated into items such as kitchen utensils, clothing, furniture, and medical devices. It is so ubiquitous, 75% of the general United States population have detectable levels in their urine (Calafat et al. 2008). Healthcare workers that use triclosan-containing hand soaps in hospitals are found to have increased urinary triclosan levels (Maclaas et al. 2014), suggesting occupational settings are a relevant site of exposure. Over the last five years, epidemiological studies have identified positive correlations between urinary levels of triclosan and sensitization to food, aeroallergens and asthma perturbations (Clayton et al. 2011; Savage et al. 2012, 2014; Bertelsen et al. 2013; Spanier et al. 2014). Our laboratory has found that topical skin application of triclosan augments allergic responses in a mouse ovalbumin (OVA) asthma model and although it does not act as an allergic sensitizer (Anderson et al. 2012, 2015), it can induce expression of the cytokine thymic stromal lymphopoietin (TSLP) in the skin tissue that augments IL-4 and IL-13 immune responses (Marshall et al. 2015). It was also recently shown that IgE and IgG1 responses after epicutaneous sensitization to peanut were augmented in mice with co-application of 1% triclosan (Tobar et al. 2016). With increasing reports that exposure to anti-microbial chemicals has the potential to induce or augment allergic disease, further studies are needed to help identify the mechanisms by which this occurs to help understand ways to prevent or circumvent these effects.

The present study sought to identify innate immune factors induced in the skin during repeated triclosan exposure that may augment subsequent adaptive allergic responses. The results indicated that repeated topical application of triclosan induced expression of S100A8 (MRP-8; calgranulin A) and S100A9 (MRP-14; calgranulin B). These small calcium-binding proteins fall under the category of alarmins, or damage-associated molecular pattern proteins (DAMP) that are released by activated or damaged cells. They can exist as both homodimers and heterodimers, and increased expression of these proteins was found in skin infiltrating leukocytes during topical application of OVA though a thymic stromal lymphopoietin (TSLP) pathway in mice. In the present study, early immunological effects following triclosan exposure were further evaluated following topical application in a murine model. These investigations revealed abundant expression of S100A8/A9, which reportedly acts as an endogenous ligand for Toll-like Receptor 4 (TLR4), in skin tissues and in infiltrating leukocytes at early time points. Increased expression of TLR4 along with TLR1, TLR2 and TLR6 in skin tissues over time with triclosan exposure; high levels of TLR4 expression were observed on skin-infiltrating leukocytes. In vivo antibody blockade of the TLR4/MD-2 receptor complex impaired local inflammatory responses after four days, as evidenced by decreased IL-6, TNFα, IL-10, IL-17, and IFNγ. These findings serve as a basis for further studies exploring the anti-microbial effects of triclosan in innate immunity as well as broad-spectrum antimicrobials.
Interestingly, some of the functional effects of S100A8/A9 have been demonstrated to involve binding and activating of Toll-like receptor 4 (TLR4)-Myd88 signaling. TLR4 is best known for recognizing lipopolysaccharide (LPS) expressed on the outer membrane of Gram-negative bacteria. It forms a complex with myeloid differentiation protein-2 (MD-2) that helps to facilitate clustering and internalization of the TLR4 receptor (Shimazu et al. 1999; Kobayashi et al. 2006). This TLR4/MD-2 complex is expressed by multiple cell types including endothelial, epithelial and myeloid cells. Because the TLR4 pathway has been shown to play a role in allergic inflammation (Miller & Modlin 2007), the current study focused on identifying a potential role for the TLR4 pathway in augmenting immunomodulatory responses following triclosan exposure.

Materials and methods

Animals

BALB/cAnNTac mice (female, 6–8 week-of-age) were purchased from Taconic (Germantown, NY). This strain has a T12 bias and is commonly used for investigations of allergic disease (Woolhiser et al. 2000). Mice were group-housed in sanitized HEPA-filtered cages containing sterilized hardwood chip bedding and nesting enrichment materials in the AAALAC-accredited NIOSH animal facility in Morgantown, WV. Mice were provided NIH-31 modified irradiated rodent diet (Harlan, Indianapolis, IN) and tap water ad libitum. Tissues were collected from the mice immediately after euthanasia by CO2 overdose. All animal procedures were reviewed and approved by the NIOSH Animal Care and Use Committee.

Animal procedures

Triclosan (Calbiochem; CAS number 3380-34-5) was dissolved in acetone (Sigma, St. Louis, MO) as a percent solution (weight/vol-ume). Concentrations and timepoints used in these studies were based on observations from previous studies (Marshall et al. 2015). Triclosan is typically incorporated into consumer products in the range from 0.03 to 1.0 pg/ml. Thus, the present study used a lower dose (0.75%) that falls within this range but does not produce overt skin irritation, as well as a high dose (3%) that produces transient irritation and more significant measurable effects.

Here, 25 μl of solution was pipetted onto the dorsal surface of each ear at 24-h intervals for the durations indicated. Purified azide-free anti-mouse TLR4/MD2 (clone MTS/510; BioLegend, San Diego, CA) or rat IgG2a isotype control antibodies (BioLegend) were diluted in phosphate-buffered saline (PBS, pH 7.4; USP grade) and injected intra-peritoneally (200–250 μg/injec-tion) at 2- or 3-day intervals as indicated.

Cell isolation

Lymph nodes

Superficial cervical (auricular) lymph nodes (LN) were collected, homogenized between frosted microscope slides, and viable cell counts performed on a Cellometer according to acridine orange/propidium iodide staining (Nexcelom, Lawrence, MA). The LN cells (5 × 10⁶) were then seeded into 96-well U-bottom plates with soluble anti-CD3 (5 μg/ml) and anti-CD28 (2 μg/ml) in complete RPMI media containing l-glutamine and HEPES (Cellgro, Manassas, VA) and supplemented with 7.5% fetal bovine serum (Hyclone, Logan, UT), 50 μM 1-mercapto-ethanol and penicillin-streptomycin (Sigma, St. Louis, MO). Cells were then incubated at 37°C for 24 h in a humidified 5% CO₂/95% air incubator before well supernatants were harvested and analyzed for cytokine expression using Flowcytomix bead assay kits (formerly eBioscience, San Diego, CA).

Ears

One ear from each mouse was mechanically disrupted on a TissueLyser II in digestion media containing collagenase and DNase (STEMCELL Technologies, Vancouver, Canada). Homogenized tissues were then digested in a shaking 37°C water-bath (45 min). Tissues were then processed through 70-μm cell strainers and re-suspended in complete RPMI media. After counting, aliquots (2 × 10⁶ cells) were seeded into wells of 96-well U-bottom plates and incubated for 24 h in a 5% CO₂/95% air humidified incubator.

Ear protein analyses

One ear per mouse was mechanically disrupted using a TissueLyser II (Qiagen, Valencia, CA) in T-PER protein extraction reagent (Thermo Fisher, Grand Island, NY). The soluble protein fraction was collected and quantified by BCA protein assay (Thermo Fisher).

SDS-Page and Western blotting

For SDS-PAGE, 50 μg protein from each sample and Precision Plus Protein™ All Blue Prestained Protein Standards (Bio-Rad, Hercules, CA) were separated over 15% polyacrylamide gels and detected with Imperial™ protein stain (Bio-Rad). For Western blotting analyses, ear tissue proteins (5 μg/lane) were separated by SDS-PAGE and transferred overnight onto nitro-cellulose membranes. The membranes were then incubated in Tris-buffered saline (TBS) blocking buffer containing 0.05% Tween-20 and 5% dry milk to mitigate nonspecific inter-actions. Thereafter, membranes were incubated with either anti-mouse S100A8 antibody or anti-mouse S100A9 antibody (R&D Systems, Minneapolis, MN) at a manufacturer-recommended concentration of 1 μg/ml blocking buffer. Membranes were then washed in TBS blocking buffer containing 0.05% Tween-20 and incubated with anti-rat horseradish peroxidase (HRP) antibody (R&D Systems; 1:1000 dilution in blocking buffer). Ultimately, the membranes were developed using ECL™ Western blotting detection reagents (GE Healthcare, Pittsburgh, PA) and immunoreactive bands then detected by exposing the membranes to CL-XPosure™ film (Thermo Fisher).

Mass spectrometry

Select protein bands were first excised from SDS-PAGE gels. Spots were de-stained, reduced, alkylated and digested with por-cine trypsin (Sigma) according to the method of by Shevchenko et al. (2006). Specifically, digested proteins were subjected to ultra-performance liquid chromatography (UPLC) on a nanoAcquity system (Waters, Milford, MA). Eluent from the nanoUPLC system was directed to the positive nanoelectrospray (+nanoESI) source of a Synapt quadrupole time-of-flight mass spectrometer (qTOF MS). Multiplexed tandem mass spectra were then acquired in “MS” fashion and data were analyzed using ProteinLynx Global Server v 2.5.1 (Waters, Milford, MA) to search the entire non-redundant SwissProt protein database.
(www.uniprot.org). The search was constrained by using 20 ppm mass accuracy (m/Δm), carbamidomethyl cysteine as a fixed modification, trypsin as enzyme and a minimum of two peptides to match. Unmatched de novo peptide sequences were scanned against SwissProt using the PAM30MS scoring matrix.

**Luminex**

For this assay, 25 μg tissue lysate or supernatants of leukocytes isolated from the ear tissue (diluted 1:5) were analyzed for S100A8 and S100A9 expression by a luminex assay according to manufacturer instructions (R&D Systems) on a Luminex 200 system (Millipore, Billerica, MA). Results are presented as mean fluorescence intensity (MFI) for tissue lysates or pg/ml for supernatants.

**RT-PCR**

Tissues were mechanically disrupted on a TissueLyser II in Buffer RTL (ears) or Qiazol (LN) (Qiagen, Valencia, CA). Total RNA was extracted using Qiagen RNasey mini-spin column kit with DNase treatment (ears) or a Qiagen miRNeasy kit (LN) on a QIAcube work station. RNA levels and purity were analyzed on a NanoDrop spectrophotometer (Thermo Fisher). The cDNA (1–2 μg) was prepared on an Eppendorf Mastercycler using a High Capacity Reverse Transcription kit (Thermo Fisher). The cDNA was used as template for real-time PCR reactions containing TaqMan PCR Master Mix with gene-specific primers (Thermo Fisher) on a 7500 Real-Time PCR System (Thermo Fisher). Relative fold-gene expression changes (2−ΔΔCT) was determined compared to acetone controls unless otherwise indicated and normalized for expression of housekeeping gene Actb.

**Flow cytometry**

Cells were re-suspended in staining buffer (PBS +1% bovine serum albumin +0.1% sodium azide) containing Fc receptor blocking antibody anti-mouse CD16/32 (clone 2.4G2; BD Biosciences, San Jose, CA). Cells were next suspended in staining buffer containing fluorochrome-conjugated antibodies recognizing cell surface epitopes in different combinations that included: CD45-Allophycocyanin, CD8-V500 (BD Biosciences), CD45R-Alexa Fluor® 488, MHC II-Alexa Fluor® 700, CD11b-PerCP-Cy5.5 (eBioscience, San Diego, CA), CD4-Brilliant Violet 421®, TLR4-Phycoerythrin and IL-4Rα-Phycoerythrin (BioLegend, San Diego, CA). Cells were then washed in staining buffer and fixed in Cytofix Buffer (BD Biosciences). For intracellular Phospho-STAT6-Phycoerythrin staining, cells were first cultured in complete media for 90 min with or without recombinant interleukin (IL)-4 (5 ng/ml; BD Biosciences) and then immediately fixed in Phos-flow fix buffer (37°C) for 15 min and permeabilized over night in Perm Buffer III (–20°C) (BD Biosciences) prior to staining. For each evaluation, 100,000 events were analyzed on an LSR II flow cytometer (BD Biosciences). Compensation controls were prepared with OneComp eBeads (eBioscience). All data analyzes were performed using FlowJo 7.6.5 software (TreeStar Inc., Ashland, OR).

**Statistical analyses**

One- or two-way analysis of variance (ANOVA) of log-transformed data followed by Fisher’s Least Significant Difference (LSD) or Tukey’s HSD post-hoc tests were performed as indicated using JMP software (SAS Institute, Cary, NC) with an α level of 0.05. All data shown are the means ± SEM.

**Results**

**Repeated topical application of triclosan induces S100A8/A9 expression in skin tissue**

Proteins in the soluble fractions of lysates prepared from ear tissue of mice treated daily for 10 days with acetone control or 3% triclosan and resolved them by SDS-PAGE gel electrophoresis were evaluated for initial investigations into the innate immune factors with repeated exposures. An abundant protein band with an approximate molecular mass of 13 kDa was identified in the skin tissue of triclosan-treated mice (Figure 1(A), black rectangle). Tandem mass spectrometric analysis identified this band to be calcium-binding protein S100A9. Two immunoreactive bands were further resolved by Western blot analysis for S100A9, likely reflecting different post-translational modifications (Figure 1(B)). A third band was also detected at ≈27 kDa, the predicted size of intact S100A9 homodimers or heterodimer. As S100A9 can form heterodimers with S100A8, immunoblotting for S100A8 revealed a band of the expected 10 kDa size in triclosan-treated mice. These results demonstrate that repeated topical skin application of triclosan can induce the expression of S100A8/A9 proteins.

Expression kinetics of S100A8/A9 in the skin tissue during repeated triclosan exposure was next evaluated. S100a8 and S100a9 transcripts were measured over the course of 12 days of exposure.

![Figure 1](image-url)
daily topical application of 0%, 0.75% or 3.0% triclosan. Both $S100a8$ and $S100a9$ transcripts were significantly increased by Day 4 (3%) or Day 6 (0.75%) in triclosan-treated mice compared to in 0% acetone controls. Relative levels of transcript continued to increase dose-dependently over time, peaking at levels >1000-fold for the 3% dose [compared to 0% acetone control] by Day 9 (Figure 2(A)). These effects were limited to the skin as we were not able to detect increases in $S100a8$ or $S100a9$ in the blood of mice exposed to triclosan at any time point (data not shown). Luminex® multiplex analysis of soluble proteins extracted from the ear tissue revealed $S100A8/A9$ protein expression was already significantly increased in the skin after only four days of 3% triclosan exposure (Figure 2(B)) and remained significantly elevated through Day 12.

Leukocytes, isolated from skin tissue, were evaluated to determine if infiltrating cells were responsible for the increased expression observed at the early timepoints. Approximately half of the cells recovered were determined to be neutrophils (CD11b+ Ly6g+) after eight days of 3% triclosan exposure (data not shown). After 24h of ex vivo culture, dose-dependent increases in $S100A8$ and $S100A9$ proteins were detected in supernatants (Figure 2(C)), suggesting infiltrating leukocytes are one of the sources of $S100A8/9$ in skin tissue during repeated triclosan exposure.

**Changes in TLR expression in skin tissue with repeated triclosan exposure**

Expression kinetics of TLR in skin tissue were examined as the $S100A8/A9$ heterodimer has been described to act as an endogenous ligand for TLR4 (Figure 3(A)). Interestingly, the family of surface receptors that recognize bacterial ligands including $Tlr1$, $Tlr2$, $Tlr4$ and $Tlr6$ had similar expression kinetics, with significant increases over time with 3% triclosan exposure relative to acetone control. Small but significant increases in $Tlr2$ expression was also observed at the 0.75% concentration. Although some significant changes in gene expression following triclosan exposure were observed, the expression kinetics were different for the endosomal nucleic-acid sensing family of TLRs (TLR3, TLR7, TLR8, TLR9) suggesting these were differentially regulated in the skin tissue compared to the cell surface family of TLR. TLR4 was also highly expressed on skin-infiltrating CD45+ leukocytes, including neutrophils, with significant increases in expression following exposure to 3% triclosan (Figure 3(B)). No increases in TLR2 expression on leukocytes were detected and the expression of TLR4 and TLR2 were not increased on cells in the LN or spleen (data not shown).

**TLR4/MD-2 antibody blockade decreases early inflammatory responses in skin with dermal triclosan exposure**

Since robust increases in $S100A8/A9$ and TLR4 expression were identified, it was further investigated whether the TLR4 pathway was mediating immunomodulatory effects with repeated triclosan exposure. A blocking antibody that recognizes TLR4 and associated MD-2 molecule or an isotype control was administered during topical application of 0% or 3% triclosan (Figure 4(A)). The 3% dose was selected for these evaluations since the majority of changes were identified at this concentration. On Day 5, effects of triclosan and anti-TLR4/MD2 on molecular and cellular responses in the skin and LN were examined. Following four days of repeated triclosan exposure transcript levels of select Tlr and $S100a8$ and
were starting to increase, therefore the Day 5 timepoint was selected to evaluate early immunomodulatory effects. Repeated triclosan application induced significant increases in pro-inflammatory cytokines in the skin, including Tslp, Il1b, Il6 and Tnfa, with no significant effects on Il25 or Il33 relative to control (Figure 4(B)). Of these, Il6, S100a8, S100a9 and Tlr2 were significantly reduced after TLR4/MD-2 blockade compared to isotype control along with the loss of significant increases in Tnfa expression compared to acetone control. The expression of Tlr1, Tlr4 and Tlr6 were also no longer significantly increased relative to acetone control after TLR4/MD-2 blockade. Taken together, these results suggested that expression of the bacterial-sensing surface TLR family of receptors could be partially regulated through TLR4 signaling in skin tissue and that activation of the TLR4 pathway promoted early pro-inflammatory responses in skin tissue during triclosan exposure.

**TLR4/MD-2 antibody blockade decreases early Th2 cytokine responses in LN with dermal triclosan exposure**

Local draining LN was also examined to determine whether TLR4/MD-2 antibody blockade impaired localized immune responses induced with repeated triclosan application. A significant decrease in total LN cellularity for triclosan-exposed mice with anti-TLR4/MD-2 treatment compared to isotype control was identified (Figure 5(A)). Additional phenotyping revealed the significant decreases were not unique to one particular cell type as there were no changes in frequencies of different cell subsets (data not shown) and numbers of CD4 T-cells, CD8 T-cells, B-cells and dendritic cells (DC) were all significantly decreased with TLR4/MD-2 blockade (Figure 5(B)).

LN T-cells were also stimulated ex-vivo to reveal effects of triclosan and anti-TLR4/MD-2 blockade on T-cell cytokine production. Early significant increases in Il-2, -6 and -17 production following triclosan exposure were not affected by TLR4/MD-2 blockade (Figure 5(C)). Additional phenotyping revealed low CD67L and CCR7 expression on the T-cells (data not shown), suggesting the population consists of a heterogeneous mixture of primary effectors and effector memory T-cells of unknown antigen specificities. At this Day 5 timepoint, IL-4 and IL-13 were significantly increased at the transcript level with triclosan exposure and interestingly, the levels of IL-4 transcript and protein were significantly decreased with TLR4/MD-2 blockade (Figure 5(C, D)). Other cytokines including interferon (IFN)-γ, IL-5, IL-10 and tumor necrosis factor (TNF)-α were...
also produced by LN T-cells but were not significantly affected by triclosan exposure or TLR4/MD-2 blockade (data not shown).

The phenotype of LN antigen-presenting cells was also examined. No differences in the expression of co-stimulatory markers CD80 and CD86 or MHC II on B-cells and DC were noted with TLR4/MD-2 blockade (data not shown). As the expression levels of IL-4 and IL-13 were significantly decreased, expression of their shared receptor subunit IL-4Rα was also examined. There was a small but significant decrease in IL-4Rα expression on DC with TLR4/MD-2 blockade, but no effect on expression on B-cells or CD4 T-cells was observed (Figure 5(E)). As shown in Figure 5(F), LN cells from triclosan-exposed mice pulsed with IL-4 expressed increased levels of phosphorylated-STAT6 which is activated downstream of IL-4Rα. Levels of phosphorylated STAT6 were equal to that of control cells suggesting that while blocking TLR4 signaling decreased the production of IL-4 and IL-13 in the LN, the cognate receptor signaling pathway was not impaired.

**Impaired Th2 cytokine responses are sustained during extended triclosan dosing and TLR4/MD-2 blockade**

Next, the effects of extended subchronic triclosan dosing with TLR4/MD-2 blockade were examined to determine whether the effects on the skin and LN could be sustained. Both an acetone control- and triclosan-treated group of mice received anti-TLR4/MD-2 antibody to control for any effects of extending blocking activity of TLR4. Mice were dosed for nine consecutive days with blocking antibody administered every third day (Figure 6(A)).

![Figure 4](https://example.com/figure4.png)

**Figure 4.** TLR4/MD-2 antibody blockade impairs gene expression changes in skin tissue with repeated topical application of triclosan. (A) Timeline depicting administration of antibodies and topical application of triclosan to the mice. (B) Relative fold-changes in gene expression in cDNA prepared from ear tissue on Day 5 from mice treated with 3% triclosan/isotype control antibody (gray) or 3% triclosan/anti-TLR4/MD-2 (black) relative to acetone control (white) (n = 5 mice). ANOVA with Tukey’s HSD statistical analyses were performed; significant differences indicated as *p < .05, **p < .01, ***p < .001.
any measurable effects. Interestingly, the differences in gene expression in the skin tissue of the triclosan-treated mice that received anti-TLR4/MD-2 compared to isotype control on Day 5 (Figure 4(B)) were no longer detected at Day 10 (Supplementary Figure 1A).

Examination of the LN at Day 10 revealed that the decreased cellularity with anti-TLR4/MD-2 treatment had also rebounded to equal numbers as isotype control in triclosan-exposed mice (Figure 6(B)), with no significant changes in frequencies or numbers of the LN cell subsets (data not shown). When the LN T-cells were stimulated *ex vivo*, the concentrations of IL-4, -13 and -17 in the supernatants were significantly elevated relative to acetone control (Figure 6(C)). Remarkably, the amounts of IL-4 and IL-13 produced by LN T-cells from mice treated with acetone/isotype or triclosan/isotype still remained decreased after TLR4/MD-2 blockade (Figure 6(C)). Taken together, although early responses in skin and LN during triclosan exposure were impaired with TLR4/MD-2 blockade, these effects eventually rebound with the exception of impaired TH2 cytokine production that continued to persist.

**Discussion**

The goal of these studies was to identify innate immune factors induced in skin tissue with repeated triclosan exposure that may...
contribute to previously identified immunomodulatory responses. Large increases in S100A8/A9 protein expression – most likely due to infiltrating neutrophils along with de novo synthesis in the skin – were identified. S100A8/A9 protein expression increased very early before elevated transcript was measured suggesting infiltrating cells, in addition to local production were potential sources. Although keratinocytes are a known source of S100A8/A9, cells of myeloid lineage can constitutively express it and neutrophils were previously found to be a major cellular infiltrate in the triclosan-exposed mouse ear skin tissue (Marshall et al. 2015). Significant increases in S100A8/A9 expression with application of 0.75% triclosan, a concentration within the range found in consumer products (0.03–1.0%) was identified. Our studies previously found that repeated application of 0.75% triclosan did not cause overt irritation to the skin as can occur transiently with 3% triclosan (Marshall et al. 2015); yet, despite this, S100A8 and S100A9 expression reached similar levels with exposure to both concentrations. It should be noted that in addition to dermal exposure, there is also the potential for oral exposure. Studies conducted by Fang et al. (2016) estimated using Elizabethan collars that 20% of a triclosan dermal dose was absorbed through oral exposure as a result of grooming. Generally, S100A8/A9 is expressed at low levels in healthy epidermis but expression can be induced by disruptions of the skin barrier (Eckert et al. 2004). It is highly up-regulated in skin tissue after exposure to solar radiation, repeated Vaseline application or tape stripping (Marionnet et al. 2003). S100A8/A9 is a well-characterized biomarker of inflammation (Ehrchen et al. 2009; Schiopu & Cotoi 2013); however, transgenic mice over-expressing activin A have high levels of S100A8/A9 in the epidermis in the absence of inflammation (Thorey et al. 2001). S100A8/A9 is thought to play an important role in skin diseases including...
psoriasis and lichen planus but is not associated with other skin conditions such as atopic dermatitis, urticaria, or granuloma annulare (Loser et al. 2010). Thus, expression of S100A8/A9 in skin tissue does not appear to be a prerequisite for inflammation—nor is it expressed with all inflammatory skin conditions, suggesting it may be involved in the immunomodulatory activity induced following dermal exposure to certain agents like triclosan that can act as allergic immune adjuvants.

The present study found that impairing TLR-4/MD-2 activity with in vivo administration of blocking antibody suppressed early responses in the skin tissue during triclosan exposure, including decreases in the expression of the cytokines IL-6 and TNF-α. Induction of IL-6 and TNFα along with increases in cell proliferation have also been observed in human keratinocytes stimulated with S100A8/A9 (Nukui et al. 2008). Tlr1, Tlr2, Tlr4 and Tlr6 showed similar expression kinetics with repeated topical application of triclosan, and blocking TLR4 signaling impaired their expression on Day 5 of exposure. These data suggested that regulation of this family of bacteria-sensing surface receptors was closely connected and controlled in part through the TLR4 receptor signaling pathway. The lack of expression changes at sites other than the skin, suggested factors unique to the environment at the site of triclosan application promoted the recruitment of TLR4+ leukocytes and induce expression of TLR4 in the skin tissue.

With the robust increases in S100A8/A9, we postulate this is likely an important endogenous ligand activating the TLR4 pathway. Further, although significant increases in Tlr4 expression at the 0.75% dose were not observed, given that TLRs are constitutively expressed in dermal tissue, the robust increases in S100A8/A9 at this dose would still likely engage the TLR4 signaling pathway. Other endogenous ligands of TLR4 have been described including high mobility group box 1 (HMGB1) (Park et al. 2004) and methyl-CpG binding domain protein 2 (mBD2) (Biragyn et al. 2002). Thus, we cannot rule out that blocking TLR4 interactions with other ligands may also be producing immunomodulatory effects as well.

It is important to consider that commensal bacteria that inhabit the skin can be stripped away with exposure to antimicrobials. In mice, gram-negative bacteria represent a large portion of the commensal skin population that can be recognized by TLR4, along with populations of gram-positive bacteria that can be recognized by TLR2. Skin commensals can interact with the immune system (Naik et al. 2012); as such, alterations of the skin microbiome may produce immunomodulatory effects. Interestingly, allergic contact hypersensitivity (CHS) induced by 2,4,6-trinitro-1-chlorobenzene (TNCB) was shown to be ameliorated in the congenitally absent of TLR2 and TLR4 as well as IL-12Rβ2 and TLR4 but not in the absence of TLR4 alone (Martin et al. 2008). To determine a role for interactions with commensal bacteria, TNCB was then applied to the skin of germ-free (GF) mice which showed that CHS was induced in both wild-type GF mice as well as IL-12Rβ2−/− GF mice, but not in double-deficient TLR4−/− IL-12Rβ2−/− GF mice. This suggested an important role for interactions with endogenous ligands like S100A8/A9 in promoting CHS even in the absence of skin commensals.

Here, by Day 10, measurable differences in gene expression in the skin tissue after TLR4/MD-2 blockade compared to isotype control were no longer detected in triclosan-exposed mice. The loss of these effects suggests engagement of alternative and/or redundant receptor pathways are eventually sufficient to mediate the responses induced by repeated triclosan exposure. While some of the extracellular functions of S100A8/A9 are mediated through TLR4, effects have also been described through the engagement of the receptor for advanced glycation end-products (RAGE) which activates the MAP Kinase/NF-κB pathway (Zhong et al. 2015). In the skin, although levels of RAGE were not affected by TLR4 blockade, relative levels of transcript eventually increased four-fold by Day 12 in triclosan-treated mice (Supplementary Figure 1B). However, a role for RAGE in the responses to triclosan has not yet been identified. Other pathways may also be activated. S100A9-induced airway inflammation in mice for example, was reported to occur independent of both RAGE and TLR4 signaling (Chen et al. 2015). Thus, the roles for the S100A8/A9-TLR4 axis in the modulation of immune responses are still not fully understood.

The present study found that engagement of the TLR4/MD-2 receptor helped to augment early inflammatory responses in the skin tissue as well as increases in IL-4 and IL-13 production by T-cells in the LN of triclosan-exposed mice. These findings provide additional insight into the immunomodulatory mechanisms of action following topical exposure to triclosan and suggest that components of the S100A8/A9-TLR4 pathway may be involved in previously-observed immune adjuvancy effects.

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**Disclosure statement**

The authors declare no conflicts of interest. The authors alone are responsible for the content of this manuscript.

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