

ORIGINAL ARTICLE

A limited CpG-containing oligodeoxynucleotide therapy regimen induces sustained suppression of allergic airway inflammation in mice

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ABSTRACT

Background CpG-containing oligodeoxynucleotides (CpG-ODNs) are potent inhibitors of T helper 2 mediated allergic airway disease in sensitised mice challenged with allergen. A single treatment has transient effects but a limited series of treatments has potential to achieve clinically meaningful sustained inhibition of allergic airway disease.

Objective To optimise the treatment regimen for sustained efficacy and to determine the mechanisms of action in mice of an inhaled form of CpG-ODN being developed for human asthma treatment.

Methods We set up a chronic allergic-asthma model using ragweed-sensitised mice exposed weekly to intranasal ragweed. Using this model, the effects of a limited series of weekly intranasal 1018 ISS (CpG-ODN; B-class) treatments were evaluated during treatment and for several weeks after treatments had stopped but weekly allergen exposures continued. Treatment efficacy was evaluated by measuring effects on lung T helper 2 cytokines and eosinophilia, and lung dendritic cell function and T-cell responses.

Results Twelve intranasal 1018 ISS treatments induced significant suppression of bronchoalveolar lavage eosinophilia and interleukin 4, 5 and 13 levels. This suppression of allergic T helper 2 parameters was maintained through 13 weekly ragweed exposures administered after treatment cessation. Subsequent experiments demonstrated that at least five treatments were required for lasting suppression. Although CpG-ODN induced moderate T helper 1 responses, suppression of allergic airway disease did not require interferon γ but was associated with induction of a regulatory T-cell response.

Conclusions A short series of CpG-ODN treatments results in sustained suppression of allergic lung inflammation induced by a clinically relevant allergen.

INTRODUCTION

Current treatments for allergic asthma provide temporary relief but do not address the causes underlying allergic disease.¹ While allergen-specific immunotherapy provides long-term relief for allergic rhinitis, it is less effective for asthma.^{2–4} Thus, the need to develop treatment alternatives for asthma that have long-lasting, disease-modifying effects is apparent.⁵

Key messages

What is the key question?

- Can CpG-containing oligodeoxynucleotide (CpG-ODN) treatments induce lasting suppression of allergic airway disease?

What is the bottom line?

- A limited weekly series of intranasal CpG-ODN treatments induced a T helper 2 to T helper 1 and regulatory T-cell shift in the airway and long-lasting suppression of allergic airways disease despite ongoing allergen exposures.

Why read on?

- Delivery of CpG-ODN directly to the lungs represents a potential disease-modifying therapy for extended remission from allergic asthma.

The potential of CpG-containing oligodeoxynucleotides (CpG-ODNs) as an immunomodulating therapy for allergic asthma has been demonstrated in animal models. CpG-ODNs stimulate innate immune responses through toll-like receptor 9 (TLR9) expressed principally by plasmacytoid dendritic cells (pDCs) and B cells in humans.⁶ In allergen challenge models, prophylactic or therapeutic CpG-ODN administration inhibited airway hyperresponsiveness, mucus secretion, eosinophilia and T helper 2 (Th2) cytokines.^{7–10} Three to six-month treatment models in mice and monkeys demonstrate inhibition of lung inflammation and remodelling with continued CpG-ODN dosing.^{11–14} However, enduring suppression of Th2 responses in the presence of frequent allergen exposures long after completion of a limited CpG-ODN treatment course, a model more relevant to clinical application, has not been demonstrated in animal models of allergic asthma.

Evidence that brief treatment with CpG-ODN could produce long-term suppression of allergic disease comes from clinical studies using the CpG-ODN 1018 ISS conjugated to the ragweed (RW) allergen *Amb a 1*. Studies showed that six



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doses lowered nasal and systemic Th2 responses with reduction in allergy symptoms for two RW seasons after treatment.^{15–17}

CpG-ODN therapy thus has potential for disease-modifying effects. The optimum strategy for treating allergic asthma, usually triggered by multiple allergens, is application of non-conjugated CpG-ODN directly to the respiratory tract, the site of allergen exposure. However, treatment optimisation requires an understanding of the regimens and mechanisms needed to produce long-term allergic disease suppression. Thus, we developed a chronic mouse model focused on therapeutic delivery of free 1018 ISS directly to the airways via intranasal administration. We demonstrate that a short series of weekly treatments induces months-long inhibition of RW-induced lung Th2 responses. We assessed the contributions of dendritic cell function and T-cell responses to maintenance of this CpG-ODN-induced disease modification.

MATERIALS AND METHODS

Chronic allergic model and CpG-ODN therapy

Female BALB/c mice (6–8 weeks old; Charles River) were sensitised intraperitoneally with 15 µg RW (*Ambrosia artemisiifolia*;

Greer Laboratories) in alum on days 0 and 7. Starting day 14, mice received, under isofluorane anaesthesia, weekly intranasal low-dose RW (0.5 µg) exposures for 1–26 weeks, were rested for 2 weeks and given a final high dose (5 µg) RW challenge 24 h before sacrifice. 20 µg 1018 ISS (B-class; 5'-TGACTGTGAACGTTTCGAGATGA), synthesised as previously described,¹⁸ was co-administered with RW to indicated groups (Tx protocol 1; figure 1A). In certain experiments, RW + 1018 ISS administration was limited to a set number of weeks, after which mice received only weekly RW exposures (Tx protocol 2; figure 2A) before rest and final RW challenge.

Tissue harvest and cell isolation

Serum was isolated using separation tubes (BD Biosciences). Bronchoalveolar lavage fluid (BALF) was processed and cytospins prepared for differential counting as previously described.¹⁹ Single-cell suspensions from treatment group pooled lung tissue, spleen or lymph nodes were also prepared. Peripheral blood mononuclear cells (PBMCs) were isolated from heparinised mouse blood using Lympholyte (CedarLane) according to manufacturer's instructions.

A RW + 1018 ISS Tx Protocol 1

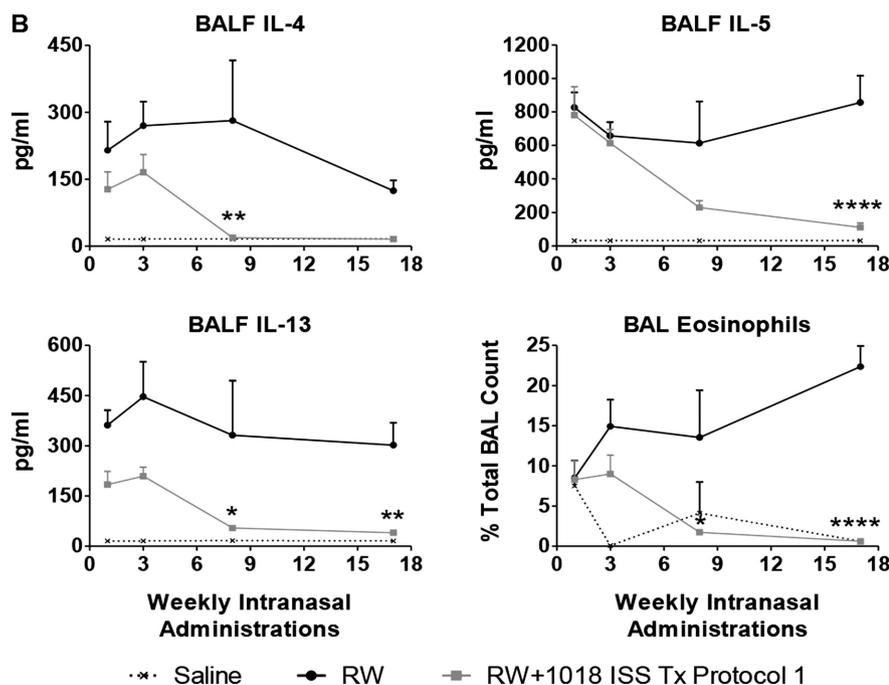
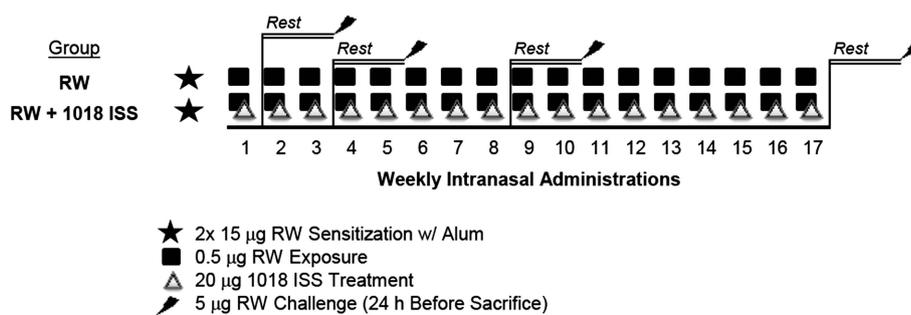
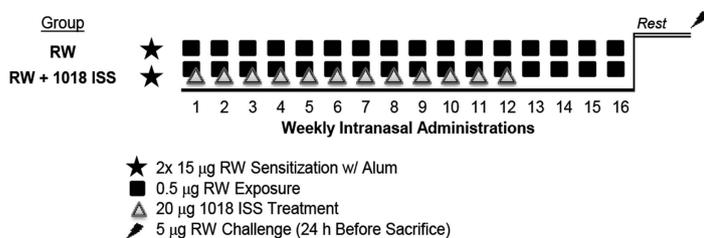
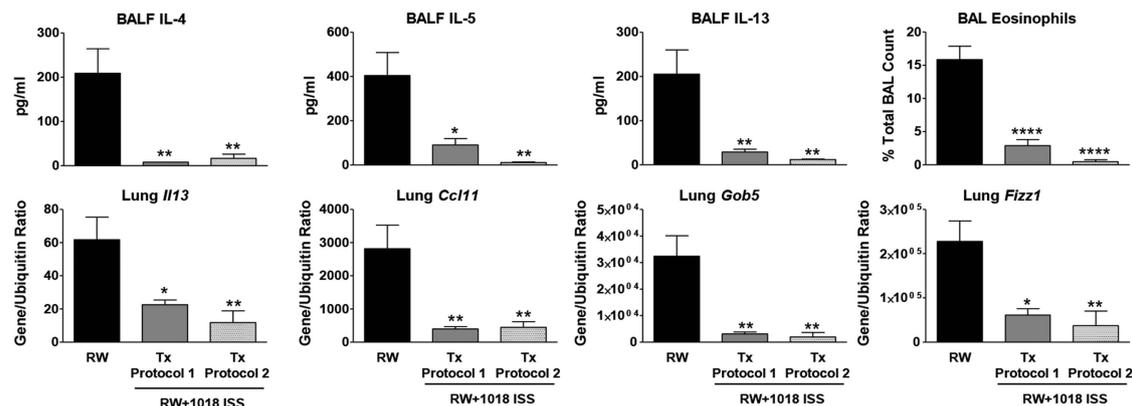


Figure 1 1018 ISS treatments progressively suppress airway T helper 2 responses. (A) Tx protocol 1 schematic: mice received 1, 3, 8 or 17 weekly intranasal saline, 0.5 µg ragweed (RW), or RW+20 µg 1018 ISS treatments, were rested 2 weeks, and then given a final 5.0 µg RW challenge 24 h before sacrifice. (B) Bronchoalveolar lavage fluid (BALF) cytokines levels and eosinophil numbers (mean±SEM, 5 mice/group). * $p < 0.05$, ** $p < 0.01$, **** $p < 0.0001$ compared with the RW only exposed group. Data are representative of four independent experiments. IL, interleukin.

A RW + 1018 ISS Tx Protocol 2



B Tx Protocol 1 versus Tx Protocol 2 (+4 weekly RW exposures post 1018 ISS treatments)



C Tx Protocol 1 versus Tx Protocol 2 (+13 weekly RW Exposures post 1018 ISS treatments)

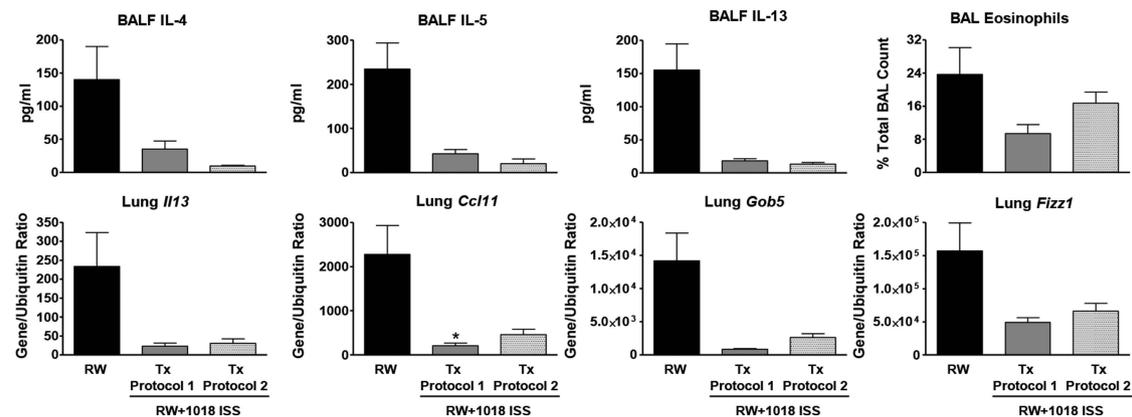


Figure 2 A limited series of 1018 ISS treatments induces lasting suppression of airway T helper 2 responses. (A) Tx protocol 2 schematic: groups were similar to Tx protocol 1 except that 1018 ISS was given for 12 weeks, followed by either 4 (B) or 13 (C) weekly ragweed (RW) only exposures followed by rest and final RW challenge. Bronchoalveolar lavage fluid (BALF) cytokines, eosinophils and lung gene expression are shown (mean \pm SEM, 5 mice/group). * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.0001$ compared with the RW only exposed group. Data are representative of four independent experiments. IL, interleukin.

ELISA

BALF cytokines and serum antibodies were quantified using matched antibody pairs (BD Biosciences) or ELISA kits (R&D Systems or Alpha Diagnostics International) according to manufacturer's instructions. Lower limits of detection were 8, 16, 8, 8 and 13 pg/mL for interleukin (IL)-4, IL-5, IL-13, IL-10 and interferon γ (IFN γ), and 250 and 100 000 ng/mL for IgE and IgG2a, respectively.

Gene expression analysis

Real-time PCR of lung tissue RNA was performed as previously described.^{18–19} Data are normalised as gene/ubiquitin ratio. Microarray analysis was performed by Expression Analysis, Inc. (Durham, North Carolina, USA).

Bone marrow-derived dendritic cell generation and adoptive transfer

Bone marrow was harvested from naïve BALB/c mouse femurs and marrow-derived dendritic cells (BMDCs) were generated as per Lutz et al's protocol.²⁰ After 9 days of maturation, BMDCs were RW pulsed (100 μ g/mL) overnight and washed prior to adoptive transfer. BMDCs were typically 80–85% I-A^{d+} and CD11c⁺. Mice received 2×10^6 RW-pulsed or non-pulsed BMDCs intratracheally as a final challenge 24 h before sacrifice, following Tx protocol 2. Control mice received RW (5 μ g) or saline intratracheally.

T-cell analysis

Enriched lung T cells were stained with surface markers. Intracellular cytokine responses of group-pooled lung T cells

were measured using de novo CD154 expression to identify antigen-specific T cells *ex vivo*,²¹ unless otherwise specified. Fixed cells were analysed on a FACSCaliber or LSRII flow cytometer (BD Biosciences). *Ex vivo* activity of lung CD25⁺ T_{Reg} cells was assessed by co-incubation with CD25⁻ cells and measurement of anti-CD3-induced proliferation by thymidine incorporation.

Statistical analysis

Individual sample data are expressed as mean±SEM. All study groups were tested for equal variance and compared with the respective RW-exposed group using one-way or two-way analysis of variance, the Mann–Whitney U test or t test. A p value of <0.05 was regarded as statistically significant.

Additional details on the methods used are presented in the online supplementary data file.

RESULTS

A series of 1018 ISS treatments induces long-lasting suppression of allergic responses in chronically RW-exposed mice

To test whether ongoing CpG-ODN treatments could achieve lasting inhibition of lung Th2 responses, we developed an allergic asthma model with a treatment and allergen exposure pattern intended to mimic clinical delivery of inhaled CpG-ODN to patients. After sensitisation to RW in alum, mice were exposed weekly to low-dose intranasal (0.5 µg) RW. Treated mice received 20 µg 1018 ISS intranasally with weekly RW exposures (Tx protocol 1, figure 1A). Treatment protocol 1 represents concurrent weekly allergen exposures and 1018 ISS treatments. Groups of mice were sacrificed after 1, 3, 8 and 17 weeks to determine how many treatments were required to suppress inflammation in the presence of ongoing allergen challenges. To assess treatment effects without the confounding effects of the direct response to 1018 ISS, mice were rested for 2 weeks before a final challenge with a high dose (5 µg) of RW. Control mice were RW sensitised, but exposed to saline thereafter.

Chronic RW exposures induced Th2-polarised lung inflammation, with large increases in IL-4, IL-5 and IL-13 protein, and BAL eosinophilia (figure 1B), which continued through 17 weekly allergen challenges. Th2 parameters were also elevated

in mice given 1–3 weekly treatments with 1018 ISS in conjunction with weekly RW exposures (Tx protocol 1), indicating that allergic airway inflammation was initially evident in these mice. By 8 weeks of RW+1018 ISS treatments there was a sharp reduction to background in all markers of Th2 activity which continued through 17 weeks, the final time point assessed (figure 1B). Eight weeks of 1018 ISS treatments were also efficacious when treatment initiation started after mice received 3 weeks of RW only intranasal exposures (modified Tx protocol 1; see online supplementary figure S1), further demonstrating that a series of 1018 ISS treatments can reverse established allergic airway inflammation.

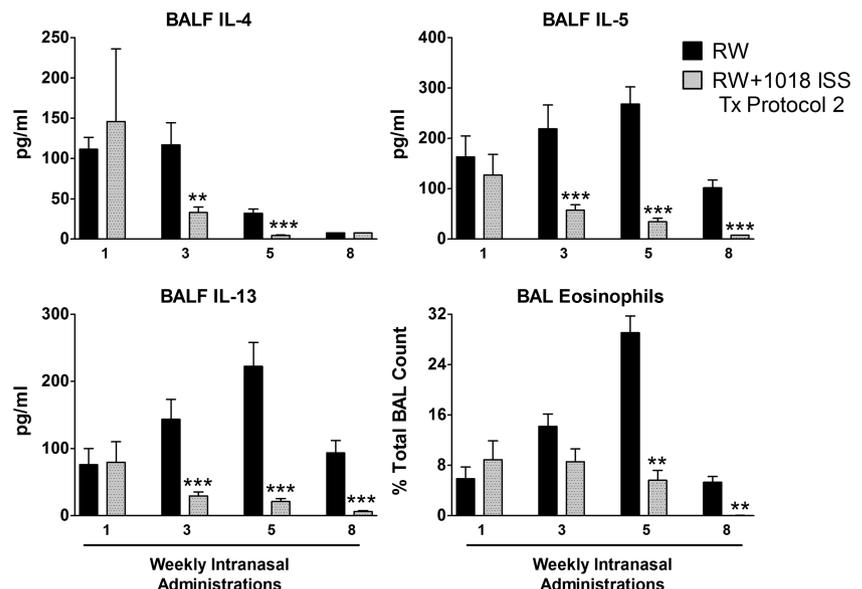
To determine whether a limited course of CpG-ODN treatment had durable effects in chronically allergen-exposed mice, mouse groups received weekly allergen exposures after completion of 1018 ISS therapy (Tx protocol 2, figure 2A). Treatment protocol 2 represents a limited course of weekly allergen and 1018 ISS exposures, followed by weekly RW exposures alone. Strikingly, allergic responses were equally inhibited by 12 weekly RW+1018 ISS treatments followed by 4 weekly RW exposures (Tx protocol 2) vs 16 weekly RW+1018 ISS treatments (Tx protocol 1). BALF Th2 cytokines, eosinophil levels and lung Th2-associated genes *Il13*, *Ccl11*, *Gob5* and *Fizz1* were suppressed equally with either protocol (figure 2B). Tx protocol 2 was also highly effective when 13 weekly RW exposures were administered after 12 weekly RW+1018 ISS treatments (figure 2C). Thus, a limited series of CpG-ODN treatments inhibited allergic lung inflammation for over 3 months after treatment in the presence of continued weekly allergen exposures.

To determine the minimum number of treatments required for disease modification, Tx protocol 2 was varied to include groups receiving 1, 3, 5 or 8 weekly RW+1018 ISS treatments, followed by 4 weekly RW exposures each. At least 5 weekly 1018 ISS treatments were required to achieve this significant, lasting reduction of Th2-mediated responses to continued allergen exposure (figure 3).

1018 ISS treatments induce a shift from a RW-specific Th2 to Th1 response

To determine whether CpG-ODN treatments induced a Th1 shift in lungs of RW allergen-exposed mice, we measured lung

Figure 3 Five weekly 1018 ISS treatments are required for durable suppression of allergen-induced T helper 2 responses in the airways. Mice were administered Tx protocol 2 with 1, 3, 5 or 8 weekly ragweed (RW) +1018 ISS treatments followed by 4 weekly RW exposures compared with mice given only weekly RW exposures. All mice were lastly rested 2 weeks and administered a final RW challenge. Bronchoalveolar lavage fluid (BALF) cytokines and eosinophils are depicted (mean±SEM, 5 mice/group). **p<0.01, ***p<0.001 compared with the RW only exposed group. Data are representative of two independent experiments. IL, interleukin.



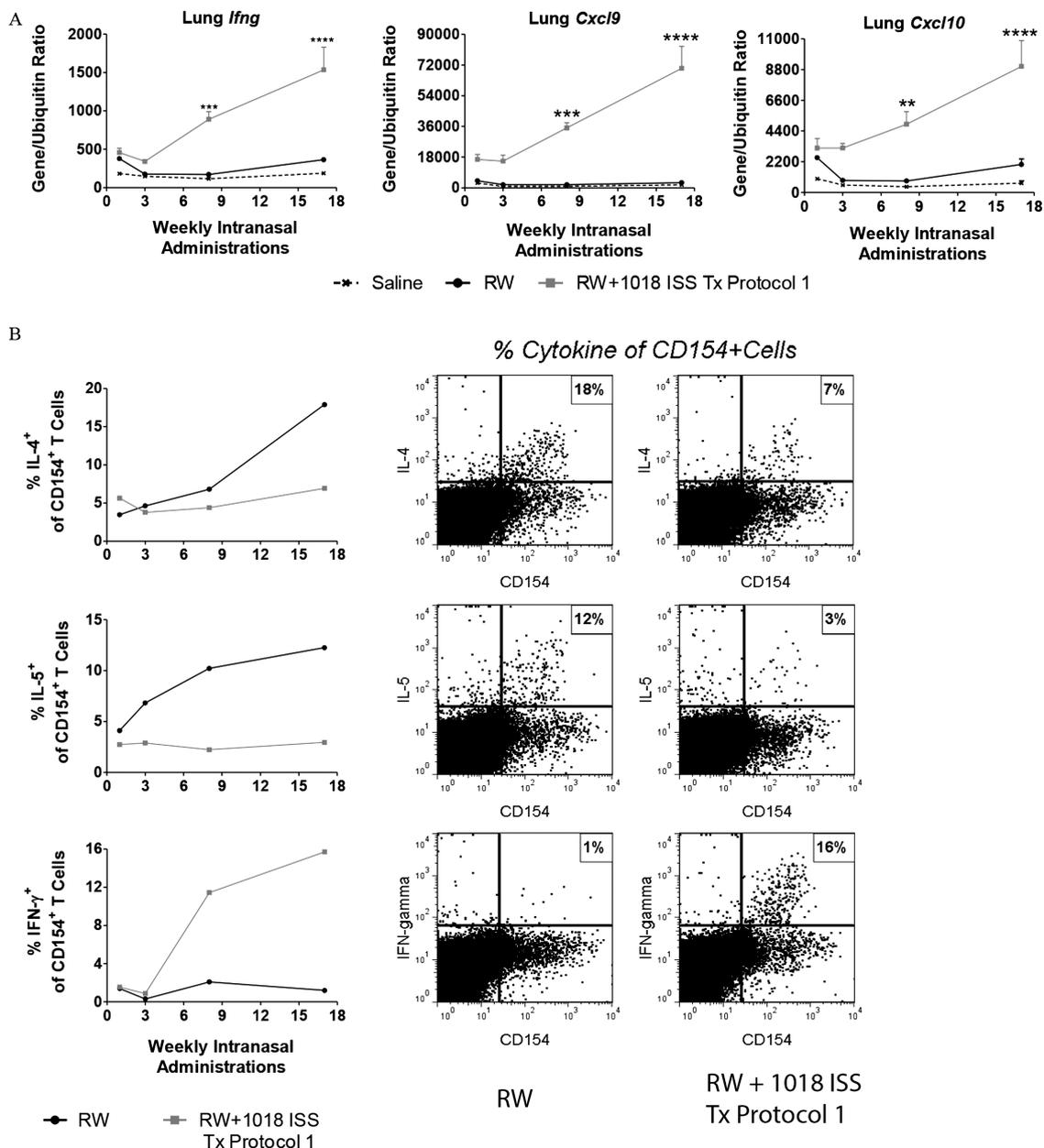


Figure 4 1018 ISS progressively induces a T helper 1 shift in lungs of treated mice. (A) Lung gene expression is shown as gene/ubiquitin ratio for mice administered Tx protocol 1 for the indicated number of weeks (mean \pm SEM, 5 mice/group), followed by rest and final ragweed (RW) challenge. ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$ compared with the RW only exposed group. (B) Kinetics of intracellular cytokine expression following ex vivo RW re-stimulation of lung-derived CD4 T cells (pooled/treatment group). Antigen-specific T cells from mice administered Tx protocol 1 for indicated weeks were identified by de novo CD154 expression. Antigen-specific intracellular cytokine expression after 17 weeks of Tx Protocol 1 is also depicted in dot plots. Data are representative of two to four independent experiments. IL, interleukin.

Th1-associated genes. RW+1018 ISS treatments progressively increased expression of Th1-associated genes, *Ifng*, *Cxcl9* and *Cxcl10* compared with expression in RW only exposed mice (figure 4A; Tx protocol 1). This Th2-like to Th1-like shift in RW+1018 ISS-treated mice was observed directly in RW-specific lung T cells (figure 4B). Following ex vivo re-stimulation with RW, lung-derived antigen-specific CD4 T cells were identified by de novo expression of CD154.^{21 22} After 17 weekly RW only exposures, cells were predominantly Th2-like, but predominantly Th1-like in mice treated for 17 weeks with RW+1018 ISS (Tx protocol 1; figure 4B). Secreted BALF IFN- γ protein was below the limit of detection (<13 pg/mL), suggesting moderate induction of Th1 responses

in 1018 ISS treated mice. Nevertheless, the lung Th1 shift was reflected systemically as indicated by reduced IgE and elevated IgG2a in serum of mice administered RW+1018 ISS (Tx protocol 1; see online supplementary figure S2).

To determine the role of IFN γ in maintenance of the Th2 to Th1 shift induced by 1018 ISS, mice were administered 13 weekly RW+1018 ISS treatments (Tx protocol 2) and then treated with anti-IFN γ antibody, isotype or no antibody during the 4 weekly post-treatment RW exposures. Once established, lasting Th2 suppression was not reversed by IFN γ neutralisation. BALF IL-4, IL-5 and serum IgE, and IL-4 or IL-5 expressing RW-specific T cells in the lung or blood remained low in anti-IFN γ treated mice (figure 5A,B). Inhibition of lung *Cxcl9*

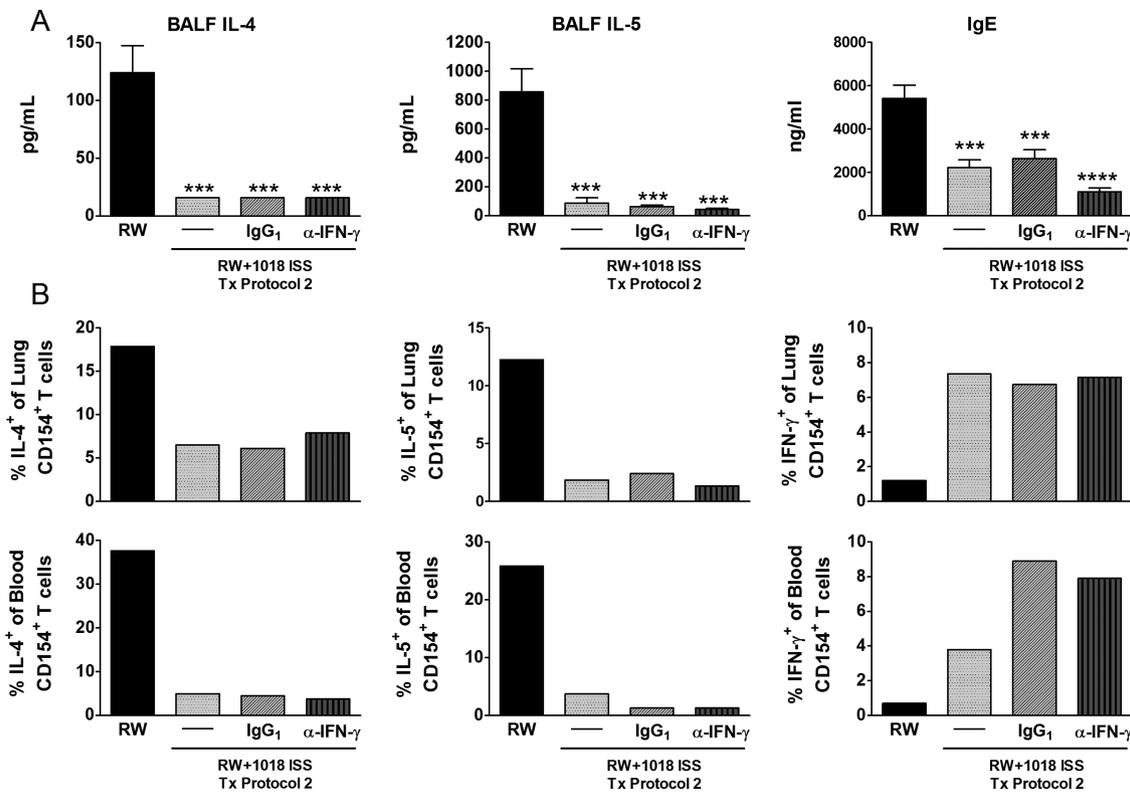


Figure 5 Interferon γ (IFN γ) is not required for maintenance of 1018 ISS-induced disease modification. Mice were administered Tx protocol 2 with 13 weekly ragweed (RW)+1018 ISS treatments, followed by 4 weekly RW exposures, rest and final RW challenge. Neutralising anti-IFN γ antibody or isotype control was administered via the intraperitoneal route concurrently with the last 4 weekly RW exposures. (A) Bronchoalveolar lavage fluid (BALF) cytokines and serum IgE (mean \pm SEM, 5 mice/group). *** p <0.001, **** p <0.0001 compared with the RW only exposed group. (B) Antigen-specific intracellular cytokine expression (tissue pooled/group). Data are representative of two independent experiments. IL, interleukin.

and *Cxcl10* expression in RW+1018 ISS-treated mice confirmed the activity of IFN γ antibodies (data not shown).

RW-pulsed dendritic cells cannot restore allergic responses in 1018 ISS-treated mice

A single CpG-ODN treatment inhibits the ability of lung antigen-presenting cells (APCs) to activate Th2 cells.¹⁹ To determine whether APC dysfunction could explain 1018 ISS-induced durable suppression, mice were challenged with exogenous APC pulsed with RW. Intratracheal challenge with allergen-pulsed BMDCs is comparable to airway allergen challenge and sufficient to induce all features of allergic airway inflammation in sensitised mice.²³ Here, mice given 5 weekly RW+1018 ISS treatments, followed by 4 weekly RW exposures (Tx protocol 2) were challenged with RW-pulsed BMDCs (RWpDCs), untreated DC, RW or saline. RW-exposed mice challenged with RWpDCs had elevated BALF IL-5, IL-13 and eosinophilia, and lung Th2-associated genes (figure 6A). However, RWpDCs did not 'restore' Th2 responses in 1018 ISS-treated mice (figure 6A), but rather induced expression of Th1-associated genes *Ifng*, *Cxcl9* and *Cxcl10* (figure 6B). Thus, allergen presented by DCs never exposed to 1018 ISS did not reinstate the Th2 response to RW in 1018-treated mice, indicating that lung APC function defects are not sufficient to explain 1018 ISS-mediated, long-lived inhibition of Th2 responses.

1018 ISS induces elevated regulatory T-cell-associated genes and proteins and increased regulatory T-cell activity

Since neither IFN γ nor chronic APC suppression could fully account for persistence of 1018 ISS-induced Th2 inhibition, we

used microarray analysis for a comprehensive and unbiased approach to identify differences between RW, RW+1018 ISS-treated and control mice. To assess steady-state gene expression, no final RW challenge was administered to mice rested 2 weeks after 16 weekly RW+1018 ISS treatments (Tx protocol 1). Microarray revealed a pattern of 1018 ISS-induced upregulation of regulatory T-cell (T_{Reg})-associated genes. Genes for Foxp3, IL-10 and transforming growth factor β (TGF β) (*Tgfb1*), and T_{Reg}-associated proteins CD223 (*Lag3*), CD134 (*Tnfrsf4*) and CD25 (*Il2ra*) were all expressed at significantly higher levels in 1018 ISS-treated mice than RW-exposed or control mice (figure 7A). The expected Th2 or CpG-ODN-inducible gene expression was also evident in RW and RW+1018 ISS treated mice, respectively (see online supplementary tables S1 and S2).

Elevated T_{Reg}-associated gene expression in RW+1018 ISS-treated mice was reflected at the protein level, except for CD25, as shown by flow cytometry of lung CD3⁺CD4⁺ T cells (pooled/treatment group) from RW+1018 ISS-treated compared with RW-exposed mice (figure 7B; Tx protocol 2, 5 weekly RW+1018 ISS, followed by 4 weekly RW, no final challenge). Additionally, CD3⁺CD4⁺Foxp3⁺ T cells from lungs of 1018 ISS-treated mice (Tx protocol 1, 16 weekly RW+1018 ISS, no final challenge) demonstrated an increased proportion of CD25⁺CD127^{lo} T cells containing increased CD39⁺CD73⁺ T cells (figure 7C). CD39⁺CD73⁺ expression ranged from ~5–8% for T cells from RW+1018 ISS-treated mice to ~0–2% for T cells from RW-exposed mice. Lastly, lung CD4⁺CD25⁺ 'regulatory' T cells were incubated at different ratios with CD4⁺CD25⁻ 'effector' T cells (T_{Eff}) from the lungs of RW-exposed mice as well as irradiated spleen cells (as APCs).

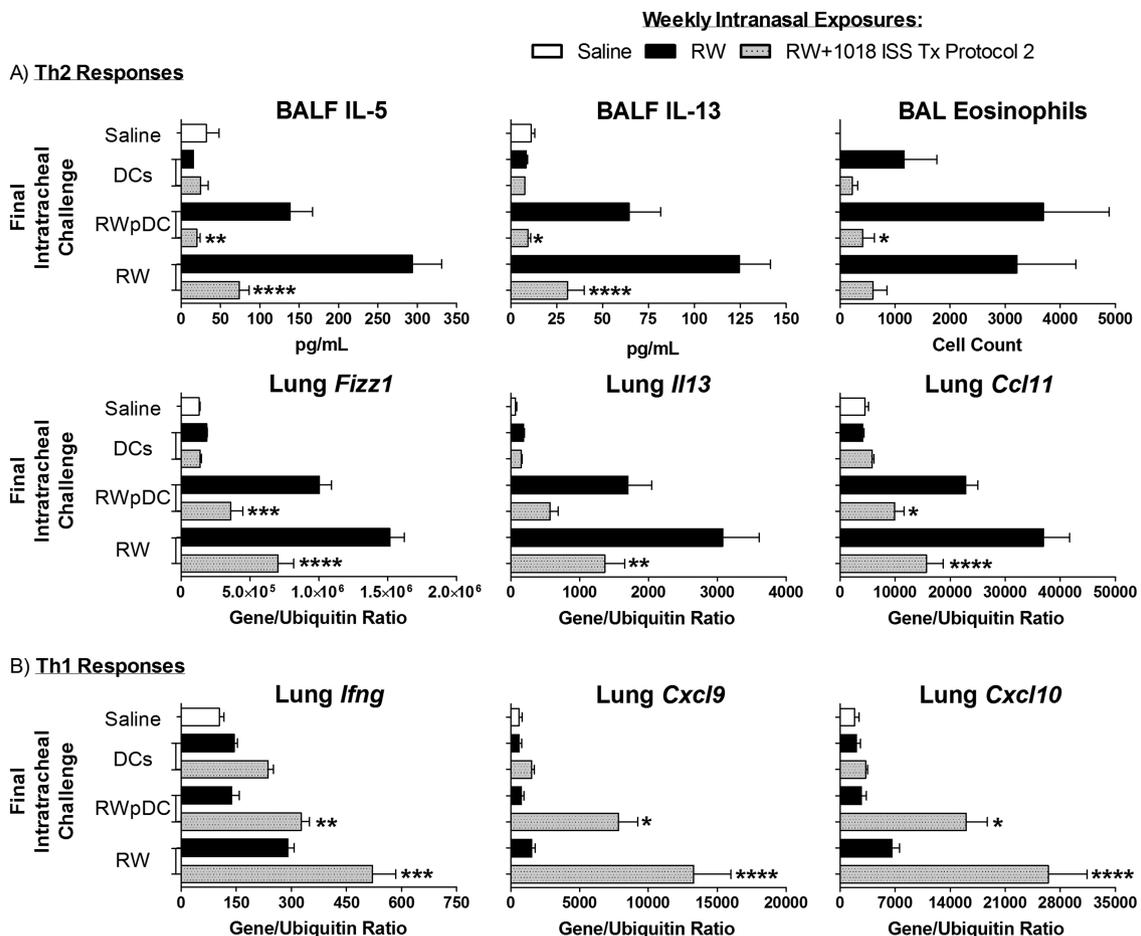


Figure 6 Ragweed (RW)-pulsed marrow-derived dendritic cells (BMDCs) do not restore T helper 2 (Th2) responses in 1018 ISS-treated mice. Mice were given Tx protocol 2: 5 weekly RW+1018 ISS treatments or RW only, followed by 4 weekly RW exposures, 2 weeks of rest and a final challenge of pulsed DCs (RWpDCs), DCs, RW or saline administered intratracheally. (A) Th2 cytokines, eosinophils and gene expression, and (B) Th1 gene expression are shown for the different treatment groups (mean±SEM, 8 mice/group). * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$ compared with the corresponding RW only weekly exposures group. Data are representative of two independent experiments. IL, interleukin.

On a per cell basis, T_{Reg} cells from RW+1018 ISS-treated mice suppressed anti-CD3 stimulated T_{Eff} cell proliferation more effectively than T_{Reg} from RW-exposed mice (figure 7D). These data demonstrate that a short course of 1018 ISS treatment induces a functional regulatory T-cell response associated with reduced allergic lung responses in this chronic asthma model.

DISCUSSION

Defining the conditions for generating durable suppression of allergic airway disease is a key step in developing a CpG-ODN treatment regimen for humans. Previous reports of CpG-ODN treatment in longer-term murine asthma models (3–6 months) examined effects within days of the last dose administered or only after a very limited number of allergen exposures following therapy, a scenario not representative of natural chronic allergen exposure.^{11–13} To model human allergic asthma as closely as possible, we developed a long-term asthma model using weekly exposures to a clinically relevant allergen (RW) administered at a dose (0.5 μ g) much lower than previously reported in limited RW allergen exposure studies.^{24–25} Lung Th2 inflammation was evident in RW and RW+1018 ISS exposed groups after the first intranasal exposure/treatment and was sustained for half a year in mice receiving weekly RW-only exposures. Using this model, we tested whether a limited course of inhaled CpG-ODN therapy would suppress allergic lung inflammation and whether

this effect would be durable in the presence of weekly allergen exposures continuing after treatment cessation. The drug was delivered directly to the lung by intranasal administration, which has been shown to be more effective than distal delivery,²⁶ and is the intended route for clinical application.

Our data show that 12 weekly 1018 ISS treatments induce suppression of allergic airway inflammation lasting through 13 subsequent weekly RW exposures and a final high-dose (5 μ g) RW allergen challenge 2 weeks later (ie, suppression was maintained for a total of 15 weeks post treatment). Therefore the period of post-treatment disease inhibition lasted longer than the treatment period and likely may last even longer, as no loss of efficacy was observed. A minimum of 5 weekly 1018 ISS doses were required to suppress established Th2 responses and the suppression lasted for at least 6 weeks without loss of efficacy. Thus, CpG-ODN offers the potential to induce disease-modifying effects in allergic asthma after a relatively brief treatment regimen, possibly to multiple allergen sensitivities. In comparison, allergen-specific immunotherapy generally requires much longer treatment regimens (2–3 years) to induce sustained mono-specific remission lasting in time up to the length of the treatment regimen.²⁷

CpG-ODN are known to induce Th1 responses^{6–28} and the durable suppression of Th2 responses achieved here was accompanied by a moderate Th1-like shift in the lungs of 1018

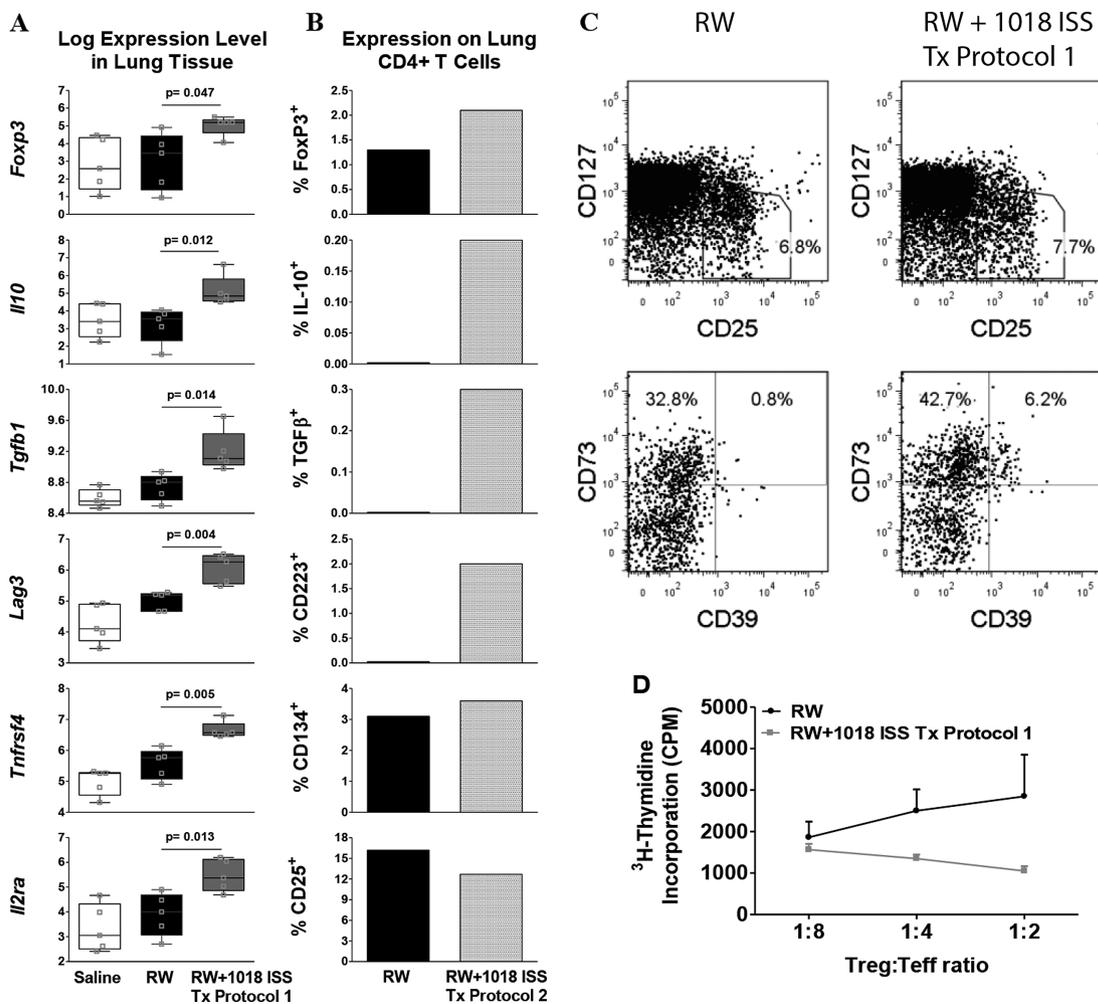


Figure 7 1018 ISS-induced disease modification is associated with induction of a lung T_{Reg} response. (A) Lung tissue expression of T_{Reg} -associated genes revealed by microarray analysis (16 weeks of Tx protocol 1, no final challenge). Student's t test p value, compared with ragweed (RW), is indicated on box-and-whisker plots (1 experiment/ $n=5$ /group). (B) Expression of T_{Reg} markers on lung CD4⁺ T cells and ex vivo anti-CD3 (10 μ g/mL) +anti-CD28 (2 μ g/mL) stimulated intracellular cytokines by flow cytometry (pooled/treatment group; 5 weeks of Tx protocol 2, then 4 weeks of RW only, no final challenge). (C) CD39 and CD73 expression on lung CD25^{hi}CD127^{lo}CD4⁺CD25⁺Foxp3⁺ T cells (pooled/treatment group; same protocol as A). (D) Proliferative responses of anti-CD3-stimulated lung T_{Eff} (CD4⁺CD25⁻) from RW-exposed mice incubated with irradiated splenic antigen-presenting cells (APCs) and lung T_{Reg} (CD4⁺CD25⁺) from RW or RW+1018 ISS treated mice at indicated ratios (mean \pm SEM of triplicate cultures, cells pooled/treatment group; 7 weeks of Tx protocol 1, no final challenge). Average for T_{Eff} in absence of T_{Reg} was 2000 CPM. Data are representative of three to five independent experiments.

ISS-treated mice. The continuing presence of a small population of lung RW-specific Th2 cells in 1018 ISS-treated mice suggests that the treatment regimen may not completely eliminate Th2 cells, but substantially compromises their function. In vivo IFN γ neutralisation after completion of 1018 ISS treatment did not reverse lung Th2 suppression (figure 5), showing that the Th1 shift is not required for continued Th2 inhibition, although it may still contribute. Sustained impairment of lung DCs cannot account for 1018 ISS-induced long-term suppression of Th2 responses as delivery of fresh antigen-loaded DCs directly to the lungs did not reignite local Th2 inflammation in 1018 ISS-treated mice (figure 6).

Animal model and clinical studies suggest that regulatory T cells have the potential to contribute to suppression of allergic airway disease. Adoptive transfer of CD4⁺CD25⁺ T_{Reg} cells into sensitised mice before or after allergen challenge reduces allergic airway inflammation^{29 30} and, conversely, T_{Reg} depletion exacerbates allergic responses.³¹ In patients on specific immunotherapy, induction of CD4⁺CD25⁺Foxp3⁺ T cells and IL-10

expression has been correlated with IgG4/IgE ratios and clinical efficacy for suppression of seasonal allergies.^{32 33} Indeed, analysis of allergen-specific T cells in healthy and allergic individuals suggests that the fine balance between T_{Reg} and Th2 cells may determine the development of allergic disease.³⁴ In the present study, expression of multiple T_{Reg} -associated genes was higher in lungs of 1018 ISS-treated mice (figure 7) and this was reflected at the protein level, except for CD25, which is also expressed on activated T cells, likely still present in the lungs at this point.³⁵ Functionally, increased T_{Reg} activity in 1018 ISS-treated mice was confirmed in an ex vivo assay. We have not defined the precise T_{Reg} mechanism(s) operating in our system, but given the upregulation of multiple T_{Reg} markers in the lungs, several mechanisms may be involved. The mixed Th1/ T_{Reg} response that resulted from 1018 ISS treatments suggests the possibility that Th1-like regulatory cells expressing IFN γ and IL-10 may form a component of this response. However, IFN γ expression by RW-specific lung T cells increased with successive 1018 ISS doses (figure 4), whereas IL-10 expression by these

cells did not differ between treatment groups at any time point (data not shown), indicating predominant induction of RW-specific Th1 cells. We did not, however, co-stain for antigen-specific T cells expressing IL-10 and IFN γ . Nevertheless, a role for IL-10 is possible as ex vivo polyclonal re-stimulation of the entire lung T-cell population revealed slightly elevated IL-10⁺CD4⁺ T cells from 1018 ISS-treated mice and lung IL-10 gene expression was also elevated in 1018 ISS-treated mice (figure 7). In this readout, other cell populations including bystander T cells may be contributing to the IL-10 response.

In summary, we evaluated whether repeated treatments with a TLR9 ligand, administered directly into the mouse respiratory tract, lead to a disease-modifying alteration of allergic airway inflammation caused by a common human allergen, RW pollen. Durable inhibition of the response to allergen challenge was achieved in as few as 5 weekly treatments with 1018 ISS co-administered with allergen and was associated with a shift from a Th2 to a mixed Th1/T_{Reg} response. Neither IFN γ nor long-lived inhibition of lung APC function was required for disease modification, although both mechanisms may contribute to the overall effect. These findings support development of inhaled CpG-ODN as a novel disease-modifying therapy for allergic asthma.

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Contributors JDC conducted experiments, analysed data, prepared figures, advised on study design, and wrote and submitted the final manuscript approved by all authors. SAK conducted the majority of the experiments, analysed data, prepared figures and contributed to manuscript writing. HMK, JAL, RS, MC and CRC performed some of the experimental work. HS analysed microarray data. CML advised on study design. RLC advised on study design and edited the article. EMH designed the study and supervised the project.

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Competing interests JDC, SAK, HMK, JAL, RS, MC, CRC, RLC and EMH are or were employees of Dynavax Technologies Corporation which funded the work.

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A limited CpG-containing oligodeoxynucleotide therapy regimen induces sustained suppression of allergic airway inflammation in mice

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ONLINE DATA SUPPLEMENT

SUPPLEMENTAL METHODS

Animal Procedures

Animal procedures were performed at Pacific Biolabs (Hercules, CA) or Murigenics (Vallejo, CA) and were Institutional and Animal Care Use Committee–approved following the “Guide for the Care and use of Laboratory Animals”, National Research Council (1996). All mice were acclimatized to the facilities for at least one week between shipping and initiation of experimental procedures. In certain Tx Protocol 2 experiments, intraperitoneal injections of protein G-purified anti-IFN- γ antibody (2 mg, R46A2(ATCC) or XMG1.2) were concurrent with weekly low dose ragweed exposures after completion of 1018 ISS therapy. Hybridomas secreting XMG1.2 and GL113 (isotype control) were kind gifts from Dr Stephen Stohlman (Cleveland Clinic, OH).

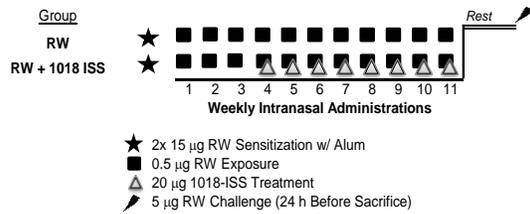
Microarray Analysis

Microarray analysis was performed by Expression Analysis, Inc. (Durham, NC). RNA was extracted from individual mouse lung samples and profiled for quality on the Agilent 2100 bioanalyzer (Agilent Technologies) and for expression on Mouse WG-6 v2 BeadChips (Illumina). Of 45,281 probes targeting transcripts, 30,221 were detected in >95% of samples. Log-2 expression levels were further analyzed using R (The Comprehensive R Archive Network), specifically to perform T-tests with Welch's correction for unequal variance and to plot data.

T Cell Analysis

For stimulation of intracellular cytokine responses, isolated and enriched lung T cells (1×10^6 /mL) were incubated in 96 well plates (200 μ L/well) with irradiated splenic white blood cells (5×10^6 /mL), anti-CD28 (1 μ g/mL) and RW (250 μ g/mL) for 6 hours with Brefledin A (5 μ g/mL) added for the final 4 hours. Intracellular cytokines were quantified as a percentage of CD154 (CD40L) antigen-reactive T cells. In general, cells were pooled per treatment group to facilitate enrichment and/or because individual mice yield low cell numbers. Non-stimulated, enriched lung T cells were surface stained with indicated antibodies (BD Biosciences or eBiosciences). Samples were collected on a FACSCaliber or a LSRII flow cytometer (BD Biosciences). Detailed analysis was performed using Flow Jo software (Tree Star, Inc) with gating through light scatter-defined lymphocytes followed by CD3⁺CD4⁺ T cells. To evaluate T_{Reg} activity in vitro, enriched lung T cells (pooled/treatment group) were stained for CD4 and CD25 and sorted on a MoFlo high speed sorter (DakoCytomation). 200 μ L triplicate cultures of CD4⁺CD25⁺ (T_{Reg}) cells from RW or RW+1018 ISS-treated mice and CD4⁺CD25⁻ (T_{Eff}) cells (5×10^4) from RW-exposed mice were stimulated with anti-CD3 (0.5 μ g/ml) in the presence of irradiated spleen cells (1×10^6) for 48 hours in 96-well U-bottom plates. Proliferation was measured following a final 6 hour pulse with 1 μ Ci of 3H-Thymidine.

A) Modified RW + 1018 ISS Tx Protocol 1



B)

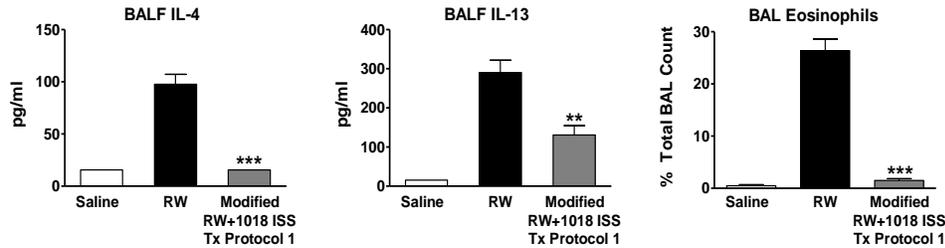


Figure S1 1018 ISS treatments initiated after 3 weekly RW exposures significantly suppress lungTh2 responses. (A) Modified Tx Protocol 1 schematic: Mice received 3 weekly intranasal 0.5 µg RW, followed by 8 additional RW, or RW+ 20 µg1018 ISS treatments, 2 weeks rest, and a final 5 µg RW challenge 24 hours before sacrifice (control mice received only saline intranasally). (B) BALF cytokines and eosinophils (mean ± SEM, 8 mice/group). **P <0.01, or ***P <0.001 compared with the RW exposures only group. Data are representative of two independent experiments.

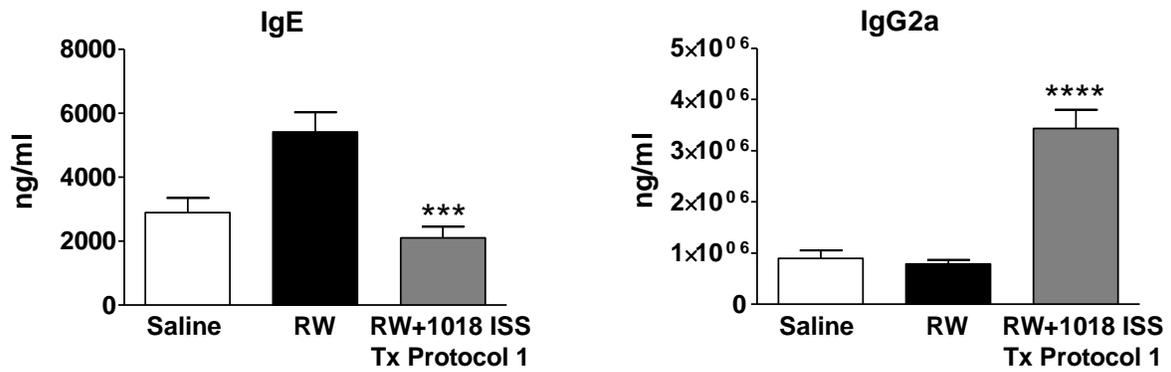


Figure S2 1018 ISS treatments induce a Th1-like shift in serum antibodies. Total serum IgE and IgG2a in mice that received 17 weeks of Tx protocol 1 RW+ 1018 ISS, RW or saline, followed by 2 rest and final RW challenge (mean \pm SEM, 5 mice/group). ***P < 0.001, ****P < 0.0001, compared with the RW exposed only group. Data are representative of two independent experiments.

Table S1: Selected Th2-associated genes upregulated in ragweed (RW) allergic mice

Genes		With Final Challenge		Without Final Challenge	
		RW versus Saline		RW versus Saline	
Common Name	<i>mSymbol</i>	Fold Induction	p-value	Fold Induction	p-value
Anterior gradient 2 (<i>Xenopus laevis</i>)	<i>Agr2</i>	17.8	0.0017	1.6	0.4602
Arginase, liver	<i>Arg1</i>	102.6	< 0.0001	1.6	0.2370
Branched chain aminotransferase 1, cytosolic	<i>Bcat1</i>	13.0	0.0016	3.5	0.0233
Chemokine (C-C motif) ligand 7	<i>Ccl7</i>	31.7	0.0001	1.3	0.2655
Chemokine (C-C motif) ligand 8	<i>Ccl8</i>	22.5	< 0.0001	3.5	0.0024
Chemokine (C-C motif) ligand 9	<i>Ccl9</i>	4.5	0.0009	1.1	0.7067
Chemokine (C-C motif) ligand 11	<i>Ccl11</i>	20.0	< 0.0001	1.9	0.0457
Chemokine (C-C motif) ligand 17	<i>Ccl17</i>	2.2	0.0129	3.3	0.0044
Chemokine (C-C motif) receptor 8	<i>Ccr8</i>	2.8	0.0817	1.5	0.4973
CD209 antigen-like protein E	<i>Cd209e</i>	192.6	< 0.0001	2.1	0.2032
Chitinase, acidic	<i>Chia</i>	8.9	< 0.0001	1.4	0.0335
Calcium-activated chloride channel family member 3	<i>Clca3</i>	566.7	< 0.0001	449.8	0.0009
Eosinophil-associated, ribonuclease A family, member 11	<i>Ear11</i>	262.0	< 0.0001	21.8	0.0042
FXD domain-containing ion transport regulator 4	<i>Fxyd4</i>	114.1	0.0004	3.0	0.0651
Glycine amidinotransferase	<i>Gatm</i>	17.1	< 0.0001	1.6	0.2127
Interleukin 4	<i>Il4</i>	24.2	0.0014	1.3	0.6631
Interleukin 5	<i>Il5</i>	3.3	0.0123	1.7	0.4084
Interleukin 13	<i>Il13</i>	1.7	0.0949	1.2	0.5898
Interleukin 13 receptor, alpha 2	<i>Il13ra2</i>	52.5	< 0.0001	1.4	0.1674
Integrin alpha X	<i>Itgax</i>	4.0	0.0066	1.9	0.3151
Matrix metalloproteinase 12	<i>Mmp12</i>	2.2	0.0084	1.5	0.0662
Homeobox, msh-like 3	<i>Msx3</i>	185.1	< 0.0001	1.5	0.3426
Periostin, osteoblast specific factor	<i>Postn</i>	2.4	0.0572	1.4	0.4992
Resistin like alpha	<i>Retnla</i>	5.2	0.0046	6.2	0.0047
Resistin like beta	<i>Retnlb</i>	7.8	0.0006	2.5	0.0754
Serum amyloid A 1	<i>Saa1</i>	8.8	0.0223	2.2	0.2086
Serum amyloid A 3	<i>Saa3</i>	5.4	0.0338	1.3	0.3834
Scinderin	<i>Scin</i>	20.8	< 0.0001	3.3	0.0232
Selectin, platelet	<i>Selp</i>	5.4	0.0002	1.2	0.0915
Solute carrier family 26, member 4	<i>Slc26a4</i>	19.1	0.0003	4.3	0.0014
Secreted phosphoprotein 1	<i>Spp1</i>	3.8	0.0014	1.2	0.0694

Mice were sensitized to ragweed pollen extract in alum by the i.p. route and were exposed weekly to low dose (0.5 µg) ragweed administered by the i.n. route for 16 weeks. Mice were then rested for 2 weeks and given a final challenge with a single high dose (5 µg) of RW 24 hours before sacrifice or were not challenged before sacrifice. Control mice were sensitized by i.p. injection with ragweed, but exposed to saline weekly thereafter. Gene expression was analyzed using Mouse WG-6 v2 BeadChips (Illumina). Values represent fold induction in RW-exposed mice in comparison to saline-exposed mice. Significance was evaluated by student's T test.

Table S2: Selected CpG-ODN-upregulated genes in ragweed (RW) allergic mice treated with 1018 ISS

Genes	mSymbol	With Final Challenge		Without Final Challenge	
		RW+1018 ISS versus Saline		RW + 1018 ISS versus Saline	
Common Name		Fold Induction	p-value	Fold Induction	p-value
Caspase 1	<i>Casp1</i>	2.3	< 0.0001	2.1	< 0.0001
Chemokine (C-C motif) ligand 4	<i>Ccl4</i>	2.9	0.0392	7.9	< 0.0001
Chemokine (C-C motif) ligand 5	<i>Ccl5</i>	2.7	0.0001	3.3	< 0.0001
Chemokine (C-C motif) ligand 19	<i>Ccl19</i>	14.9	< 0.0001	36.3	< 0.0001
Chemokine (C-C motif) receptor 5	<i>Ccr5</i>	10.3	< 0.0001	9.1	< 0.0001
CD72 antigen	<i>Cd72</i>	5.4	0.0003	6.1	0.0018
Chemokine (C-X-C motif) ligand 9	<i>Cxcl9</i>	226.4	< 0.0001	102.9	0.0002
Chemokine (C-X-C motif) ligand 10	<i>Cxcl10</i>	13.5	0.0030	10.0	0.0001
Chemokine (C-X-C motif) ligand 13	<i>Cxcl13</i>	4.9	0.0021	5.8	0.0008
Chemokine (C-X-C motif) receptor 3	<i>Cxcr3</i>	5.8	< 0.0001	6.6	< 0.0001
Guanylate binding protein 1	<i>Gbp1</i>	2.7	0.0110	1.0	0.5198
Glycosylation dependent cell adhesion molecule 1	<i>Glycam1</i>	14.1	0.0035	25.9	0.0078
Interferon-alpha 1	<i>Ifna1</i>	3.2	0.0314	2.8	0.0870
Interferon-gamma	<i>Ifng</i>	24.2	< 0.0001	16.8	< 0.0001
Interleukin 2	<i>Il2</i>	2.8	0.0549	2.5	0.0338
Interferon regulatory factor 1	<i>Irf1</i>	1.6	0.0463	1.2	0.0109
Jun-B oncogene	<i>Junb</i>	1.6	0.0025	1.1	0.3886
26S proteasome regulatory subunit p28	<i>Psmc10</i>	1.2	0.0123	1.2	0.0026
Pentraxin related gene	<i>Ptx3</i>	1.8	0.0531	1.6	0.1309
Src-like-adaptor 2	<i>Sla2</i>	6.3	0.0050	10.8	0.0002
Signal transducer and activator of transcription 1	<i>Stat1</i>	2.5	0.0109	1.5	0.0011
Signal transducer and activator of transcription 4	<i>Stat4</i>	1.6	0.0019	1.7	0.0006
Tumor necrosis factor	<i>Tnf</i>	2.8	0.0305	6.3	< 0.0001
Tumor necrosis factor receptor superfamily, member 1b	<i>Tnfrsf1b</i>	1.9	0.0003	2.4	< 0.0001

Mice were sensitized to ragweed pollen extract in alum by the i.p. route and were exposed weekly to low dose (0.5 µg) ragweed and 1018 ISS (20 µg) administered by the i.n. route for 16 weeks. Mice were rested for 2 weeks and given a final challenge with a single high dose (5 µg) of RW 24 hours before sacrifice or were not challenged before sacrifice. Control mice were sensitized by i.p. injection with ragweed, but exposed to saline weekly thereafter. Gene expression was analyzed using Mouse WG-6 v2 BeadChips (Illumina). Values represent fold induction in RW+1018 ISS-exposed mice in comparison to saline-exposed mice. Significance was evaluated by student's T test.