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## Aire is not essential for regulating neuroinflammatory disease in mice transgenic for human autoimmune-diseases associated MHC class II genes HLA-DR2b and HLA-DR4

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### Abstract

The human autoimmune disease-associated HLA alleles HLA-DR2b (DRB1\*1501) and HLA-DR4 (DRB1\*0401) are strongly linked to increased susceptibility for multiple sclerosis (MS) and rheumatoid arthritis (RA), respectively. The underlying mechanisms are not fully understood, but these MHC alleles may shape the repertoire of pathogenic T cells via central tolerance. The transcription factor autoimmune regulator (AIRE) promotes central T cell tolerance via ectopic expression of tissue-specific antigens (TSAs). Aire deficiency in humans causes autoimmune polyendocrinopathy syndrome type 1 (APS1), and *Aire* knockout mice (*Aire*<sup>-/-</sup>) develop spontaneous autoimmune pathology characterized by multi-organ lymphocytic infiltrates.

Here, we asked whether impaired TSAs gene expression in the absence of Aire promoted spontaneous MS- or RA-like autoimmune pathology in the context of human HLA alleles in HLA-DR2b or HLA-DR4 transgenic (tg) mice.

The results show that reduced TSAs gene expression in the thymus of Aire-deficient HLA-DR2b or HLA-DR4 tg mice corresponded to mild spontaneous inflammatory infiltrates in salivary glands, liver, and pancreas. Moreover, Aire-deficiency modestly enhanced experimental autoimmune encephalomyelitis (EAE) in HLA-DR tg mice, but the animals did not show signs of spontaneous neuroinflammation or arthritis. No significant changes were observed in CD4<sup>+</sup> T cell numbers, T cell receptor (TCR) distribution, regulatory T cells (Treg), or antigen-induced cytokine production. Abrogating Treg function by treatment with anti-CTLA-4 or anti-CD25 mAb in Aire-

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deficient HLA-DR tg mice did not trigger EAE or other autoimmune pathology. Our results suggest a redundant role for Aire in maintaining immune tolerance in the context of autoimmune disease-associated human HLA alleles.

## Keywords

EAE; T cells; Aire

## Introduction

While the exact cause for most human autoimmune diseases is unknown, it is believed that autoreactive T cells are key mediators of pathology in many conditions, including multiple sclerosis (MS) and rheumatoid arthritis (RA) [1–3]. Central tolerance is a key mechanism involved in eliminating autoreactive T cells in the thymus during the process of negative selection of thymocytes via presentation of self-antigens in the thymic medulla [4, 5]. In the thymus, autoimmune regulator (AIRE in humans, Aire in mice) promotes central tolerance by inducing ectopic transcription of tissue specific antigens (TSAs) normally expressed in peripheral sites [6, 7]. AIRE is primarily expressed in the thymus, but it can also be expressed in secondary lymphatic tissues. Within the thymus, it is expressed mostly by thymic medullary epithelial cells (mTECs), but also by dendritic cells (DCs), although at much lower levels [6–8]. In secondary lymphoid tissues, Aire is expressed in extrathymic Aire-expressing cells (eTACs), which may help to enforce peripheral T cell tolerance [9]. The limited overlap between Aire-dependent gene expression in eTACs and mTECs suggests that peripherally expressed Aire may regulate the expression of a unique set of self-antigens.

In humans, it has been suggested that failure in T-cell tolerance caused by mutations in AIRE gene may result in the autoimmune condition observed in autoimmune polyendocrine syndrome type 1 (APS1) [10, 11]. It is also known as autoimmune-polyendocrinopathy-candidiasis-ectodermal dystrophy (APECED) and is characterized by spontaneous multi-organ failure and chronic mucocutaneous candidiasis due to immune cell destruction and dysfunction [12]. Aire-deficient mice show loss of T-cell tolerance, mimicking the phenotype of human APS1. Spontaneous autoimmune disease characterized by lymphocytic infiltrates in multiple organs and tissues has also been observed in these mice [13–15]. Moreover, human gene association studies have reported that SNPs in the AIRE gene are associated with susceptibility to RA [16–18].

Major histocompatibility complex (MHC) class II molecules (human leukocyte antigen (HLA) in humans) are expressed by antigen-presenting cells (APCs), where they function in antigen processing and presentation to CD4<sup>+</sup> T helper (Th) cells. HLA-DR2b (DRB1\*1501) and HLA-DR4 (DRB1\*0401) alleles are associated with MS and RA, respectively [15, 19–21]. HLA-DR2b and HLA-DR4 tg mice are useful to test the role of these molecules in processing and presentation of autoantigens, shaping the T cell repertoire in the thymus and the periphery, and to interrogate their function in the pathogenesis of human autoimmune diseases [22–24]. Since these animals are devoid of endogenous murine MHC class II molecules (Ia), antigen presentation is solely the property of the human HLA-DR2b and

HLA-DR4 molecules [15, 20, 25]. We showed previously that T cell tolerance was maintained in these mice and depended on the expression of self-antigen [26]. To determine if Aire was critical in maintaining T cell tolerance in the context of HLA-DR2b and HLA-DR4 molecules, we developed HLA-DR tg *Aire*<sup>-/-</sup> mice on the experimental autoimmune encephalomyelitis (EAE)-susceptible C57BL/6 background [27, 28]. Specifically, we asked if the absence of Aire promoted the development of spontaneous organ-specific autoimmune pathology akin to MS or RA in HLA-DR tg mice. Of note, Aire deficiency on the conventional C57BL/6 background displays only a mild autoimmune phenotype in mice [29, 30].

Our studies revealed a decrease in gene expression of TSAs in thymus, which was paralleled by mild, spontaneous inflammatory infiltrates in a small number of *Aire*<sup>-/-</sup> HLA-DR2b or HLA-DR4 tg mice in salivary glands, liver, and pancreas, as well as serum autoantibodies against gastric tissue, in particular in older mice. Similarly, we noted a mild decrease in thymic gene expression of neuroantigens, which corresponded to modestly enhanced active EAE, but no spontaneous neuroinflammatory disease or autoimmune arthritis was observed. No significant changes were observed in the numbers and distribution of CD4<sup>+</sup> T cells, TCR distribution, percentages of regulatory T cells, and antigen-induced cytokine production. Furthermore, abrogation of T regulatory (Treg) function by treatment with anti-CTLA-4 and anti-CD25 mAb did not trigger EAE or autoimmune arthritis in these mice. Our results support a role for the transcription factor Aire in modulating autoimmune pathology, but argue against its essential role in regulating autoimmunity in the context of the human HLA-DR2b and -DR4 alleles associated with MS and RA, respectively.

## Materials and Methods

### Mice

*Aire*<sup>-/-</sup> C57BL/6 mice [6] were originally purchased from Jackson lab (stock no. 004743) and were then maintained on the C57BL/6 background. For the experiments described here, *Aire*<sup>+/-</sup> mice were crossed to *Ia*<sup>-/-</sup> C57BL/6 mice (Jackson lab; stock no. 003584) and the progeny were then interbred to obtain *Aire*<sup>+/-</sup>*Ia*<sup>-/-</sup> animals devoid of all endogenous murine MHC II. The HLA-DR2b and HLA-DR4 tg mice have been described previously [23, 26, 31, 32]. *Aire*<sup>+/-</sup>*Ia*<sup>-/-</sup> mice were crossed to HLA-DR2b or HLA-DR4 tg mice to obtain *Aire*<sup>+/-</sup>*Ia*<sup>-/-</sup> HLA-DR2b or *Aire*<sup>+/-</sup>*Ia*<sup>-/-</sup> HLA-DR4 tg mice. F1 offspring of *Aire*<sup>+/-</sup>*Ia*<sup>-/-</sup> HLA-DR2b or *Aire*<sup>+/-</sup>*Ia*<sup>-/-</sup> HLA-DR4 tg mice were then interbred to obtain *Aire*<sup>+/-</sup>*Ia*<sup>-/-</sup>, *Aire*<sup>+/+</sup>*Ia*<sup>-/-</sup>, and *Aire*<sup>-/-</sup>*Ia*<sup>-/-</sup> HLA-DR2b or HLA-DR4 tg mice, respectively. All mice were bred at The University of Texas at San Antonio (UTSA) under specific pathogen-free conditions. All studies were performed per UTSA Institutional Animal Care and Use Committee guidelines and approved protocols.

### Genotyping

Genomic DNA from tail snips was extracted using DNeasy Blood and Tissue kit (Qiagen). PCR primers were custom-ordered from Invitrogen. Standard PCR parameters were employed using Go Taq Green polymerase mix (Promega) on Thermal Cycler 2720 (Applied Biosystems). Primers used in the study are as following: forward primer for *Aire*

gene, 5'-GTC ATG TTG ACG GAT CCA GGG TAG AAA GT- 3'; reverse primer for *Aire* gene, 5'-AGA CTA GGT GTT CCC TCC CAA CCT CAG- 3'; forward primer for mouse MHC II (I-A) gene, 5'-GGG GTG GAA TTT GAC CTC TT- 3'; reverse primer for mouse MHC II (I-A) gene, 5'-TGG AGA CAT TGG CCA GTA CA- 3'; forward primer for HLA-DR2b gene, 5'-GTT TCC TGT GGC AGC CTA AGA GG- 3'; reverse primer for HLA-DR2b gene, 5'-TCC ACC GCG GCC CGC GC- 3'; forward primer for HLA-DR4 gene, 5'-CGT TTC TTG GAG CAG GTT AAA CA- 3'; reverse primer for HLA-DR4 gene, 5'-AAG CGC ACG TAC TCC TCT TGG TG- 3'.

### Real-Time PCR

Total RNA was extracted from the thymus of *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b or HLA-DR4 tg mice using RNeasy fibrous tissue mini kit (Qiagen). The RNA was reverse transcribed to cDNA using the cDNA reverse transcription kit (Life technologies). Real-time PCR was performed with the CFX96 Touch Deep Well Real-Time PCR Detection System (BioRad) using RT<sup>2</sup> SYBR Green master mix (Qiagen) per manufacturer's instructions. The amplification program included an initial denaturation step at 95 °C for 10 min, followed by denaturation at 95 °C for 15s and annealing and extension at 60 °C for 1 min for 40 cycles. SYBR Green fluorescence was measured after each extension step, and the specificity of amplification was evaluated by melting curve analysis. RT<sup>2</sup> qPCR primer assay for mouse *Tff3* (NM\_011575), *Ins2* (NM\_001185083), *Spt1*(NM\_009267), *Rbp3* (NM\_015745), *Muc6* (NM\_181729), *Csn1s1* (NM\_007784), *Igf2* (NM\_001122736), *Krt8* (NM\_031170), *Mog* (NM\_010814), *Pip1* (NM\_011123) and *Mbp* (NM\_001025245) from Qiagen were used. Gene expression was normalized by *Keratin 8* expression, which is specifically expressed in the thymic epithelial cell fraction and is not influenced by *Aire* [6, 33].

### Treg depletion

*Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice (3 to 9 months old) were divided into groups and injected with mAb as indicated. *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR4 tg mice were injected i.p. with 800 µg of anti-CTLA-4 mAb (Bio X cell; UC10-4F10) and *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b tg mice were given i.p a combination of 100 µg of anti-CTLA-4 mAb and 1 mg of anti-CD25 mAb (Bio X cell; PC-61.5.3) once a week for three continuous weeks.

### Immunofluorescence (IF) staining

Mouse organs (liver, pancreas, lung, stomach, and intestine) from the studies were immersed in OCT (Fisher Scientific) and frozen in -80 °C, then cryosectioned (10 µm) onto slides. Slides were fixed and stained with Alexa Fluor 488-conjugated anti-mouse CD4 mAb (eBioscience; GK1.5), APC-conjugated anti-mouse CD8 mAb (eBioscience; 53-6.7), followed by microscopic examination and imaging of positive staining cells (Olympus microscope with DP72 camera, CellSens Standard 1.5 software). The scoring system is described in Table 1. For indirect immunofluorescence staining to detect IgG autoantibodies, sera from *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice were incubated with tissue sections of organs from C57BL/6 SCID mice, followed by staining with FITC goat anti-mouse IgG Ab (Jackson Immunoresearch).

## Histology

Tissues (adrenal glands, salivary glands, testis and ovaries) from mice were harvested and fixed by immersion in 10% buffered formalin overnight. They were then rinsed in PBS, dehydrated through increasing concentrations of ethanol, cleared and embedded in paraffin. Each tissue block was cut into 3 sections, with each section 20  $\mu\text{m}$  apart and mounted on slides. The sections were stained with hematoxylin and eosin (H&E) followed by microscopic examination for cellular infiltrates (Olympus microscope with DP72 camera, CellSens Standard 1.5 software).

## Cytokine ELISPOT

ELISPOT plates (Millipore; Multiscreen IP) were coated with mouse IFN- $\gamma$ -specific (eBioscience; AN-18), IL-17-specific (Bio X Cell; 17F3) or GM-CSF-specific (MP1-22E9) capture mAbs diluted in PBS. The plates were blocked with 1% BSA in PBS for 1 h at room temperature and then washed four times with PBS. Splenocytes ( $1 \times 10^6$  cells/well) were collected and incubated with or without Ag for 24 h at 37  $^{\circ}\text{C}$ . The plates were washed three times with PBS and four times with PBS-Tween (0.05%) and incubated with mouse IFN- $\gamma$ -specific biotinylated detection mAb (eBioscience; R4-6A2), IL-17-specific biotinylated detection mAb (TC11-81-14) or GM-CSF-specific biotinylated detection mAb (MP1-31G6) at 4  $^{\circ}\text{C}$  overnight. The plates were washed four times with PBS-Tween (0.05%) and incubated with streptavidin-alkaline phosphatase (Invitrogen) for 2 h at room temperature, followed by four washes with PBS. Cytokine spots were visualized by 5-bromo-4-chloro-3-indolyl phosphate/nitroblue tetrazolium phosphatase substrate (Kirkegaard & Perry Laboratories). Image analysis of ELISPOT assays was performed on a Series 2 ImmunoSpot analyzer and software (Cellular Technology).

## EAE induction and disease evaluation

Active immunization EAE was induced in HLA-DR2b and HLA-DR4 tg mice (6–10 weeks of age) by s.c. injection of 300  $\mu\text{g}$  of myelin oligodendrocyte glycoprotein (MOG)<sub>35–55</sub> peptide and 200  $\mu\text{g}$  of MOG<sub>97–108</sub> peptide respectively, in 50  $\mu\text{l}$  of emulsion with complete Freund's adjuvant (CFA). CFA was prepared by adding 5 mg/ml of *Mycobacterium tuberculosis* H37RA (Difco laboratories) in incomplete Freund's adjuvant (IFA). Mice also received i.p injections of 400 ng of pertussis toxin (PTX) on day 0 and day 1 post immunization.

Mice were monitored and graded daily for clinical signs of EAE using the following scoring system [34]: 0, no abnormality; 1, limp tail; 2, moderate and hind limb weakness; 3, complete hind limb paralysis; 4, quadriplegia or premonitory state; 5, death.

## Flow Cytometry Analysis- Cell Surface Staining

Murine splenocytes ( $1 \times 10^6$  cells/sample) and thymocytes ( $1 \times 10^6$  cells/sample) were collected and stained for various cell surface markers after washing and blocking with anti-mouse CD16/32 to block binding through the Fc-receptor (eBioscience). The samples were then analyzed by Becton Dickinson (BD) FACS-Aria II using BD FACS Diva Software. All antibodies to cell surface markers were purchased from eBioscience or BD: PE-Cy7 anti-CD4 mAb (GK1.5), APC anti-CD8 mAb (53-6.7), PE anti-CD11c mAb (N418), Alexa Fluor

488 anti-CD11b mAb (M1/70), Alexa Fluor 488 anti-CD25 mAb (eBio7D4), FITC anti-HLA-DR mAb (LN3), PE anti-Ia mAb (AF6-120.1, BD) and PE-Cy7 anti-CD45 mAb (GK1.5). The FITC labeled mAbs for 14 V $\beta$  and 3 V $\alpha$  families were purchased from BD.

### Flow Cytometry Analysis- Intracellular Staining

Murine splenocytes were collected, fixed and permeabilized in 96-well round bottom plates ( $1 \times 10^6$  cells/well), and stained for the intracellular marker using PerCP-Cy5.5 anti-Foxp3 mAb (eBioscience; FJK-16s). The samples were then analyzed by BD FACS-Aria II using BD FACS Diva Software.

### Statistical analysis

For comparisons using two experimental groups, the *t* test was used. Analysis of variance (ANOVA) was used for statistical analysis involving multiple groups followed by Bonferroni posttest. Comparisons of all analyses were performed using Sigma Plot 12.5. A difference was considered statistically significant when  $p < 0.05$ .

## Results

### Downregulation of Aire-dependent TSAs gene expression in thymus of Aire-deficient HLA-DR2b and HLA-DR4 tg mice

To test the effect of Aire-deficiency in the context of the human HLA-DR2b and HLA-DR4 alleles we generated Aire-deficient (*Aire*<sup>-/-</sup>) Ia<sup>-/-</sup> HLA-DR2b and HLA-DR4 tg mice as described in *Materials and Methods*. The deletion of *Aire* and Ia and the presence of HLA-DR2b and HLA-DR4 were confirmed by PCR (Fig. 1A). The expression of HLA-DR2b, HLA-DR4 and absence of Ia was corroborated by flow cytometry (Fig. 1B, C).

Quantitative RT-PCR (qRT-PCR) analysis of thymic tissue was performed to confirm the effect of *Aire* deletion on the expression of known Aire-dependent TSA genes in HLA-DR tg mice (Fig. 1D, E), including trefoil factor-3 (Tff3), insulin-2 (Ins2), mouse salivary protein-1 (Spt1), interphotoreceptor retinoid-binding protein (Rbp3), mucin 6 (Muc6), casein alpha (Csn1s1) and insulin like growth factor II (Igf2) as described previously [6, 30, 35, 36]. Specifically, mRNA expression for Ins2, Spt1, Igf2 was substantially decreased in *Aire*<sup>-/-</sup> HLA-DR tg mice as compared with *Aire*<sup>+/+</sup> HLA-DR tg littermates (Fig. 1D, E). Our findings agree with previous reports showing a similar reduction of these genes in *Aire*<sup>-/-</sup> mice on other genetic backgrounds, i.e. BALB/c [6, 36, 37].

### Aire-deficiency does not affect immune cell distribution in HLA-DR tg mice

To begin to investigate the effect of Aire-deficiency on the immune system in the HLA-DR tg mice, particularly in the T cell compartment, we determined the distribution and numbers of T cells, DCs, and monocytes/macrophages in thymus and spleen of 6 – 9-week-old *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR tg mice by flow cytometry.

As shown in Fig. 1F, G & H, the percentages of single positive (SP) CD4<sup>+</sup> T cells, SP CD8<sup>+</sup> T cells, or double positive (DP) CD4<sup>+</sup>CD8<sup>+</sup> T cells were comparable in thymus of *Aire*<sup>+/+</sup> versus *Aire*<sup>-/-</sup> HLA-DR2b or HLA-DR4 tg mice. No significant gender differences were



observed (not shown). Furthermore, the overall percentage and the absolute numbers of CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells, DCs (CD11c<sup>+</sup>), and macrophages (CD11b<sup>+</sup>) were similar in the spleens of *Aire*<sup>-/-</sup> mice as compared to *Aire*<sup>+/+</sup> mice, irrespective of whether the mice were on the HLA-DR2b or HLA-DR4 background (Fig. 1I, J). We also investigated the thymus architecture in our mice by H&E staining and did not note abnormalities on histology (Supplemental Fig. 1A–D). Thus, the results showed that Aire did not have a notable effect on the numbers and distribution of T cells, monocytes, and DCs in naïve HLA-DR tg mice.

### Mild spontaneous inflammatory tissue infiltrates in HLA-DR tg mice

Aire-deficient mice have been reported to exhibit spontaneous lymphocytic tissue infiltrates [14, 15]. Therefore, we investigated the HLA-DR tg mice for clinical or histopathological signs of spontaneous autoimmune pathology in other tissues.

Tissue sections from representative organs from *Aire*<sup>-/-</sup> HLA-DR2b and HLA-DR4 tg mice up to 15 months of age were procured and investigated by immunofluorescence microscopy for the presence of CD4<sup>+</sup> and CD8<sup>+</sup> T cells, and CD11c<sup>+</sup> cells. Of note, the results showed a small percentage (~20%) of *Aire*<sup>-/-</sup> HLA-DR2b and HLA-DR4 tg mice 12 months and older with inflammatory infiltrates, mostly confined to liver and pancreas, with an average immune cell infiltration score of 1 – 2 (Fig. 2, Table 1 & 2). The inflammatory infiltrates in *Aire*<sup>-/-</sup> HLA-DR tg mice consisted predominantly of CD4<sup>+</sup> T cells (Fig. 2B, D), with very few CD8<sup>+</sup> T cells present (data not shown). CD4<sup>+</sup> T cell infiltrates were not observed in stomach, intestine and lungs (data not shown). A mild gender bias was observed, with older females showing more CD4<sup>+</sup> T cell infiltrates as compared with older males (Table 2). This observation is in line with the literature, where immunopathology was more pronounced in older female *Aire*<sup>-/-</sup> mice as compared with male *Aire*<sup>-/-</sup> mice [38]. However, in contrast, we did not observe splenomegaly in our *Aire*<sup>-/-</sup> HLA-DR tg mice and the number of splenocytes was comparable between the *Aire*<sup>-/-</sup> HLA-DR tg and *Aire*<sup>+/+</sup> HLA-DR tg mice (data not shown), conceivably due to the younger average age of the mice in our studies. Finally, no immune cell infiltration was noted in organs of *Aire*<sup>+/+</sup> HLA-DR tg littermates.

The mild gender bias noted in our studies prompted us to examine reproductive organs from these mice. However, no inflammatory infiltrates were noted in ovaries or testis procured from *Aire*<sup>-/-</sup> HLA-DR tg mice or *Aire*<sup>+/+</sup> HLA-DR tg littermates (data not shown).

In APS1 the endocrine manifestations include Addison's disease [39], which lead us to examine the adrenal glands of the animals in our studies. However, in agreement with previous studies, we did not detect inflammation of the adrenal glands in *Aire*<sup>-/-</sup> HLA-DR tg mice or littermate controls (data not shown) [13, 38].

Aire-deficient mice have been reported to exhibit lymphocytic infiltration in salivary glands [6, 40]. Similarly, we observed inflammatory infiltrates in salivary glands in 3 out of 4 *Aire*<sup>-/-</sup> HLA-DR2 tg mice, but not in *Aire*<sup>+/+</sup> HLA-DR2 tg mice by H&E examination (Fig. 2I, J). Of note, our qRT-PCR results showed a significantly reduced expression of Spt1 in *Aire*<sup>-/-</sup> HLA-DR2 and – DR4 tg mice as compared with *Aire*<sup>+/+</sup> littermates (Fig. 1D, E), which would be consistent with increased induction of autoimmunity directed towards salivary gland tissue.

The occurrence of autoantibodies in the sera of APS1 patients often precedes the onset of clinical disease and signals ongoing autoimmune responses [41]. Therefore, to examine the presence of autoantibodies, sera were obtained from *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice (1 to 16 months of age) and tissue sections from SCID mice were used to test for the presence of IgG autoantibodies. Of note, IgG autoantibodies were present in sera of 22% of *Aire*<sup>-/-</sup> HLA-DR tg mice 10 months and older, which were directed against gastric tissue, but not against any other tissues (Fig. 2F, H; Table 3); this corresponded to the reduced expression of stomach specific gene *Muc6* in the thymus. Minimal IgG autoantibodies were detected in sera of *Aire*<sup>+/+</sup> HLA-DR tg littermates (Fig. 2E, G).

Despite the presence of autoantibodies, *Aire*<sup>-/-</sup> HLA-DR2b and HLA-DR4 tg mice did not show obvious clinical signs of pathology and maintained similar body weights as compared with age and gender matched *Aire*<sup>+/+</sup> HLA-DR2b or HLA-DR4 tg littermates (Supplemental Fig. 2A–D). Moreover, no signs of spontaneous neuroinflammatory disease (e.g. EAE) were noted in *Aire*<sup>-/-</sup> HLA-DR2b tg mice or their *Aire*<sup>+/+</sup> HLA-DR2b tg littermates (data not shown). Similarly, *Aire*<sup>-/-</sup> HLA-DR4 tg mice or their *Aire*<sup>+/+</sup> HLA-DR4 tg littermates did not show joint swelling or increased articular inflammation suggestive of autoimmune arthritis (data not shown).

Collectively, the results showed that a small percentage of older *Aire*<sup>-/-</sup> mice expressing the human autoimmune disease-associated MHC class II alleles HLA-DR2b and HLA-DR4 exhibited mild spontaneous inflammatory infiltrates dominated by CD4<sup>+</sup> T cells, predominantly in liver and pancreas. Moreover, a small percentage of older *Aire*<sup>-/-</sup> HLA-DR tg mice showed IgG autoantibodies directed against gastric tissue. However, no evidence of spontaneous CNS-demyelinating disease or autoimmune arthritis was observed.

### Earlier disease onset and mild increase in EAE severity in *Aire*-deficient HLA-DR tg mice

HLA-DR2b and -DR4 tg mice did not show evidence of spontaneous neuroinflammatory disease or arthritis. However, Ko et. al. reported previously that *Aire*<sup>-/-</sup> mice showed earlier EAE onset [42]. Therefore, we asked whether *Aire*-deficiency in mice transgenic for the MS susceptibility allele HLA-DR2b would lead to more severe disease in *Aire*<sup>-/-</sup> mice compared with *Aire*<sup>+/+</sup> mice. To evaluate the effect of *Aire*-deficiency on autoimmune pathology in HLA-DR2b and HLA-DR4 tg mice, EAE was induced by active immunization with MOG<sub>35–55</sub> or MOG<sub>97–108</sub> peptide, respectively, and mice were observed for clinical signs of disease for up to 4 weeks as previously described [31, 32].

Shown in Fig. 3, disease onset occurred slightly earlier in *Aire*<sup>-/-</sup> HLA-DR2b and HLA-DR4 tg mice (day 7 and day 10, respectively) as compared with *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 littermates (day 10 and day 12, respectively; Fig. 3A, B), but the results did not reach statistical significance. Disease severity (Fig. 3A, B) and incidence (Fig. 3C, D) of *Aire*<sup>-/-</sup> HLA-DR tg versus *Aire*<sup>+/+</sup> HLA-DR tg mice were comparable and no statistically significant differences were noted. Taken together, the results suggested that *Aire*-deficiency had a minor effect on the induction of EAE in mice expressing autoimmune disease-associated HLA-DR alleles and promoted a slightly earlier onset of disease.



## Aire-deficiency alters Th17 cell responses in HLA-DR tg mice but does not skew the T cell receptor repertoire

To determine whether Aire-deficiency modulated antigen-specific T cell responses in HLA-DR tg mice, IFN- $\gamma$ , IL-17 and GM-CSF production was measured by cytokine ELISPOT assay from splenocytes at EAE remission. As shown in Fig. 4A, the frequencies of T cells producing IFN- $\gamma$ , IL-17, or GM-CSF were mildly decreased in *Aire*<sup>-/-</sup> mice expressing the HLA-DR2b allele, as compared with the control mice, with only IL-17 reaching statistical significance ( $p < 0.05$ ). In contrast, frequencies of IL-17 and GM-CSF producing T cells were significantly increased in *Aire*<sup>-/-</sup> HLA-DR4 tg mice ( $p < 0.05$ ), whereas IFN- $\gamma$  was not significantly increased (Fig. 4B). Conceivably, higher frequencies of T cells in *Aire*<sup>-/-</sup> HLA-DR4 tg mice as compared with *Aire*<sup>+/+</sup> HLA-DR4 tg could have contributed to the slightly more severe EAE phenotype noted in these mice. However, differences in cytokine production could not sufficiently explain the earlier disease onset and slight increase in disease severity noted in HLA-DR2b tg mice.

Ko et. al. observed a reduction in MOG gene expression in their *Aire*<sup>-/-</sup> mouse line [42], but not in the expression of MBP or PLP. To address this question in our model we analyzed the expression of myelin specific genes MOG, PLP, and MBP by qRT-PCR. We noted decreased MOG gene expression in *Aire*<sup>-/-</sup> mice, particularly in mice expressing HLA-DR2b (Fig. 4C), whereas the expression of MBP or PLP was essentially not altered (Fig. 4C, D). Thus, decreased MOG expression may have contributed to the slightly more severe EAE observed in our model. Conceivably, altered expression of TSAs in *Aire*<sup>-/-</sup> HLA-DR tg mice could have affected the positive and/or negative selection of T cells expressing autoreactive TCRs [4, 5]. Therefore, we tested whether Aire-deficiency globally affected TCR selection in HLA-DR2b and HLA-DR4 tg mice and resulted in a skewed TCR repertoire. To address this question, we examined the repertoire of TCRV $\beta$  and V $\alpha$  families in the thymus and spleen by flow cytometry using a panel of anti-TCR monoclonal antibodies.

Shown in Fig. 4E–L, the TCR repertoire was comparable between *Aire*<sup>-/-</sup> HLA-DR tg and *Aire*<sup>+/+</sup> HLA-DR tg mice and no significant alterations were observed in the representation of TCR V $\alpha$  or V $\beta$  families. Thus, our results suggest that Aire expression did not significantly affect the overall TCR repertoire distribution in the context of human HLA-DR2b and HLA-DR4 molecules in the transgenic mice.

## Anti-CD25/CTLA-4 mAb treatment does not trigger overt autoimmune pathology in Aire-deficient HLA-DR tg mice

Natural Tregs (nTreg; CD4<sup>+</sup> CD25<sup>+</sup> Foxp3<sup>+</sup>) develop in the thymus and are a population of lymphocytes implicated in the regulation of autoimmune T cell responses, whereas inducible Tregs (iTreg) develop in the immune periphery and similarly keep aberrant T cell responses in check [43, 44]. The role of Aire in positive selection of Tregs has been controversial. However, growing evidence suggests that Aire may influence the thymic development of Treg cells [45–49]. Therefore, we asked whether the percentage of Tregs was altered in this model. However, as shown in Fig. 5A & B, the percentage of CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> Tregs in the spleen was comparable between *Aire*<sup>-/-</sup> HLA-DR tg mice and *Aire*<sup>+/+</sup> HLA-DR tg mice.

Nevertheless, while the data showed that the overall percentage of nTregs was not affected by Aire-deficiency in naïve HLA-DR2b or HLA-DR4 tg mice, conceivably, their function could still be altered in the absence of Aire. Along these lines, CTLA-4 is an inhibitory receptor expressed by regulatory and conventional T cells, which is necessary for maintaining T cell homeostasis and self-tolerance. In conventional T cells, CTLA-4 cell surface expression is induced after TCR signaling [50]. In contrast, CTLA-4 is constitutively expressed on Tregs, where it serves to control self-reactive T cells [51]. *In vivo* blockade of CTLA-4 with anti-CTLA-4 mAb can promote organ-specific autoimmune pathology [51]. Moreover, the administration of anti-CD25 mAb results in functional inactivation of Tregs [52, 53].

To test whether inhibition of CTLA-4 and/or CD25 triggered autoimmune disease in *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b or HLA-DR4 tg mice, 3–9-month-old animals were treated with a combination of anti-CTLA-4 and anti-CD25 mAbs (for HLA-DR2b tg mice), or with anti-CTLA-4 mAb alone (for HLA-DR4 tg mice) and observed for evidence of autoimmune pathology for up to 12 weeks. Of note, no clinical signs of overt autoimmune pathology, including neuroinflammatory disease or arthritis were noted in *Aire*<sup>-/-</sup> or *Aire*<sup>+/+</sup> HLA-DR2b or HLA-DR4 tg mice (data not shown). Moreover, body weight remained comparable between *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> mice (Fig. 5C, D). To corroborate the lack of clinical disease, histopathology of liver, lungs, pancreas, stomach, and small and large intestine was performed by IF staining as described in *Materials and Methods*. Summarized in Table 4, *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b tg mice injected with a combination of anti-CD25/CTLA-4 mAb did not show inflammatory infiltrates in any of the tissues examined. However, approximately half of the *Aire*<sup>-/-</sup> HLA-DR4 tg mice, and 1/3 of the *Aire*<sup>+/+</sup> HLA-DR4 tg mice injected with anti-CTLA-4 mAb showed inflammatory infiltrates, which were mostly noted in liver and pancreas, and which consisted predominantly of CD4<sup>+</sup> T cells.

Taken together, the results showed that treatment with anti-CTLA-4 and anti-CD25 antibodies accelerated the development of lymphocytic infiltrates in younger mice, whereas infiltrates in the absence of antibody treatment were only observed in older mice. However, the effect of antibody treatment was independent of the presence or absence of Aire.

## Discussion

In this study, we show that deficiency of Aire in mice expressing human MS and RA-associated HLA-DR2b or HLA-DR4 alleles did not lead to spontaneous neuroinflammatory disease or arthritis. However, Aire-deficiency resulted in mild spontaneous autoimmune phenomena in the HLA-DR tg mice characterized by mild inflammatory infiltrates and increased autoantibodies, predominantly in mice 12 months or older. Inhibition of regulatory mechanisms by treatment with anti-CTLA-4 and anti-CD25 mAb did not trigger neuroinflammation or arthritis in the HLA-DR2b or HLA-DR4 tg mice, irrespective of the presence or absence of Aire. Moreover, induction of active EAE in Aire-deficient HLA-DR2b or HLA-DR4 tg mice resulted in earlier disease onset, but comparable disease severity. Thus, overall, the results suggest a mild and non-essential effect of Aire on modulating autoimmune disease pathology in the context of the human MS and arthritis-associated HLA-DR2b and HLA-DR4 alleles.

Central tolerance requires presentation of self-antigens by MHC molecules in the thymus to promote negative selection of potentially autoreactive T cells. Ectopic expression of self-antigens promoted by the transcription factor Aire in mTECs is important for tolerance and elimination of autoreactive T cells [54]. Aire may also promote the selection of Tregs [8]. Deficiency of Aire can lead to breakdown of central tolerance and result in multi-organ autoimmune disease [8]. Certain MHC Class II alleles, including HLA-DR2b and HLA-DR4, are associated with human autoimmune conditions, such as MS and RA, respectively [21, 55]. However, it has remained unresolved why certain HLA alleles are associated with specific autoimmune diseases. Thus, we asked whether Aire plays a role in preventing or promoting autoimmune pathology in the context of the human HLA-DR2b or HLA-DR4 alleles in HLA-DR tg mice. Specifically, we asked whether spontaneous neuroinflammation or arthritis was observed in *Aire*<sup>-/-</sup> mice expressing these autoimmune disease-associated HLA alleles.

Of note, our studies confirmed that absence of Aire resulted in downregulation of Aire-dependent ectopic tissue antigens in thymus. Consistent with impaired central T cell tolerance, and reduced expression of TSAs, specifically Spt1, Ins2 and Muc6 in the thymus, we observed mild tissue pathology in salivary glands, liver, and pancreas of some HLA-DR2b or HLA-DR4 tg mice, and, in addition, some of the animals developed increased autoantibodies directed against gastric tissue. The pathological consequences of these autoimmune phenomena appeared to be limited, because *Aire*<sup>-/-</sup> HLA-DR2b or HLA-DR4 tg mice did not show notable loss of bodyweight or other clinical signs of autoimmune disease. Most of the mice showing inflammatory infiltrates or autoantibodies were older females, suggesting an age-related effect and gender bias.

Previously it was shown that the HLA-DR transgenes in HLA-DR tg mice behave similar to endogenous murine MHC Class II genes and instruct normal thymic development and maintain normal lymphocyte development and homeostasis in the thymus and periphery [22]. Evaluation of the immune compartment of these mice for proportion and numbers of DCs, macrophages, T helper cells, cytotoxic T cells, and nTregs did not reveal dramatic changes between *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> mice. Moreover, we did not observe alterations in thymus architecture. These observations are consistent with previous reports in the literature that Aire may have a minor effect on overall thymocyte composition and development [56]. The limited effect of Aire-deficiency on global CD4<sup>+</sup> T cell development in the context of human HLA-DR alleles was further supported by lack of significant changes in the distribution of major TCR families between the *Aire*<sup>-/-</sup> HLA-DR tg mice and their *Aire*<sup>+/+</sup> HLA-DR tg littermates. We cannot exclude that more sensitive techniques, such as gene melting spectral pattern (GMSP assay) analysis might uncover minor differences in TCR selection that were not apparent in our flow cytometry analyses due to the limited coverage imposed by the available mAbs for these studies [57]. However, based on our results we would predict minor effects on the TCR repertoire.

It is conceivable that more than one mechanisms accounted for the absence of more striking autoimmune pathology in our model. Along these lines, it is conceivable that the C57BL/6 genetic background of the HLA-DR tg mice is less conducive to development of spontaneous autoimmune phenomena associated with Aire-deficiency, which is supported by

literature showing that *Aire*<sup>-/-</sup> mice on the C57BL/6 background consistently show a very mild autoimmune disease phenotype as compared with other backgrounds, e.g. BALB/c, NOD [7, 29]. Jiang et. al. showed that *Aire*<sup>-/-</sup> NOD mice crossed to *Aire*<sup>-/-</sup> B6 displayed a mild autoimmune phenotype which, was observed only in a minority of animals [29]. Thus, the lack of Aire may be most notable in the context of a particular set of disease susceptibility background genes. However, these disease susceptibility genes do not include HLA alleles known to promote autoimmune pathology. Thus, the lack of spontaneous autoimmune pathology in our Aire-deficient HLA-DR tg mice is consistent with reports showing that HLA alleles do not strongly influence autoantibody formation in APS1 patients [58]. Similarly, Ahonen et al. and others reported that HLA-DR alleles were not associated with APS1 related pathology [58–60].

The similar numbers of Tregs in our *Aire*<sup>-/-</sup> HLA-DR tg mice compared with their *Aire*<sup>+/+</sup> littermates suggested that Treg development was not impaired. To further investigate this question, we asked whether an autoimmune phenotype was elicited by interfering with other regulatory mechanisms, for example via blockade of the regulatory CTLA-4 molecule. This was inspired by reports who observed spontaneous development of chronic organ-specific autoimmune disease after *in vivo* blockade of CTLA-4 mAb [51], and by studies showing functional inactivation of Tregs upon treatment with anti-CD25 mAb [52, 53]. Also, it has been shown that CTLA-4 has dual function in Tregs and conventional T cells to prevent multi-organ autoimmunity [61]. However, in our studies, treatment of *Aire*<sup>-/-</sup> HLA-DR2b tg mice with anti-CTLA-4/CD25 mAb did not result in significantly increased immune-mediated pathology, and treatment of *Aire*<sup>-/-</sup> HLA-DR4 tg mice with anti-CTLA-4 mAb only induced mild inflammatory infiltrates in less than half of the animals. Nevertheless, the mAb treatment accelerated development of lymphocytic infiltrates in younger mice, whereas without antibody treatment we only observed these infiltrates in a small subset of older mice. Of note, we found that the percentages of Tregs in our HLA-DR tg mice were generally lower as compared with the percentages observed in C57BL/6 Wt mice (1–4% versus 10–15%). Thus, it is conceivable that the antibody treatment may not have been as effective in further reducing the Treg compartment in the HLA-DR tg mice as compared with Wt mice.

Last, we asked whether autoimmune disease pathology was increased in Aire-deficient HLA-DR tg mice using a well-established EAE model [62, 63]. However, the effect of Aire-deficiency on EAE incidence and severity in HLA-DR2b or HLA-DR4 tg mice was also mild, and mostly characterized by earlier disease onset and slight disease increase. The results of these studies are noteworthy for somewhat divergent effects of Aire-deficiency on cytokine production by autoreactive T cells in HLA-DR2b versus HLA-DR4 tg mice. While the effect of Aire on cytokine producing T cells was marginal in HLA-DR2b tg mice, *Aire*<sup>-/-</sup> HLA-DR4 tg mice showed an increase in the frequencies of T cells producing proinflammatory cytokines, in particular for IL-17 and GM-CSF. Thus, it is conceivable that the increase in T cells producing these cytokines in the HLA-DR4 tg mice was, at least partially, compensated by regulatory mechanisms, such as Treg cells.

The interpretation of our results should take under consideration that TSA expression is not completely abolished by the absence of Aire [33]. Takaba et al. demonstrated the existence

of another transcriptional regulator, FEZF2, which is specific for and highly expressed in mTECs in humans, and, similarly, numerous TRA transcripts are downregulated in Fezf2-deficient mTECs in mice. Thus, Fezf2 has an Aire-independent and non-redundant role in promoting ectopic expression of TSAs, and this mechanism may have played a compensatory role in our model [64].

In summary, our studies showed that mice deficient for Aire and tg for human autoimmune disease-associated HLA alleles developed limited spontaneous autoimmune pathology, mostly restricted to salivary glands, liver, and pancreas, and predominantly in older animals. The animals showed no major changes in immune cell subsets, particularly CD4<sup>+</sup> T cells, and increased systemic autoimmune pathology was not triggered upon treatment with anti-CTLA-4 or CD25 mAb. Active induction of EAE resulted in earlier disease onset, but with similar disease severity in Aire-deficient mice as compared with their *Aire*<sup>+/+</sup> HLA-DR tg littermates. However, the animals did not develop spontaneous neuroimmune disease or arthritis despite the presence of the human MS- and RA-associated HLA-DR2b and -DR4 alleles.

Our results support that the transcription factor Aire plays a general role in modulating autoimmune pathology, but indicate that it does not have an essential role in regulating autoimmunity in the context of the human HLA-DR2b and -DR4 alleles associated with MS and RA, respectively.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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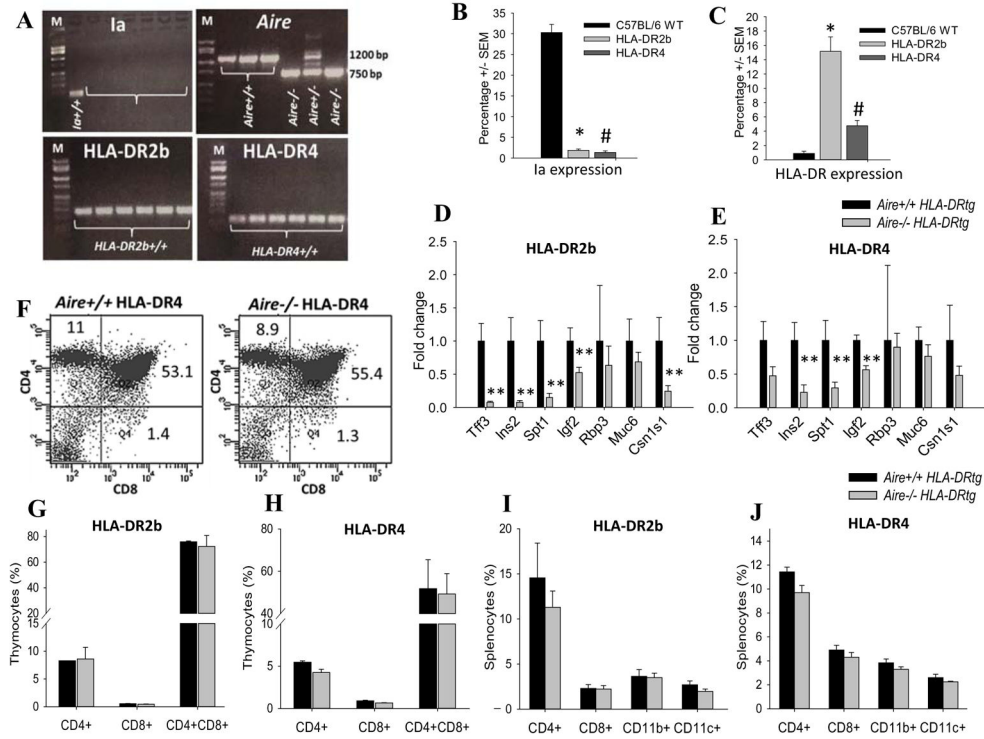
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### Highlights

- HLA-DR2b and HLA-DR4 transgenic (tg) *Aire*<sup>-/-</sup> mice were generated on the experimental autoimmune encephalomyelitis (EAE)-susceptible C57BL/6 background.
- These mice were used to investigate the role of Aire for autoimmune pathology in the context of the multiple sclerosis and rheumatoid arthritis-associated human HLA-DR2b and -DR4 molecules.
- The mice showed a decrease in thymic expression of gene expression of tissue-specific antigens in parallel with mild spontaneous inflammatory infiltrates in salivary glands, liver, and pancreas and gastric autoantibodies.
- The mice showed a modest increase in the severity of actively induced experimental autoimmune encephalomyelitis.
- However, the animals did not develop spontaneous neuroinflammatory disease or autoimmune arthritis.
- No significant effect of the Aire-deletion was noted on the immune compartment in these mice.
- The results support a role for the transcription factor Aire in modulating autoimmune pathology, but argue against its essential role in regulating autoimmunity in the context of the human HLA-DR2b and -DR4 alleles associated with MS and RA, respectively.



**FIGURE 1.** Downregulation of Aire-dependent genes and immune cell distribution in *Aire*-deficient HLA-DR2b and HLA-DR4 tg mice. Genomic DNA extracted from off-springs of *Aire*<sup>+/-</sup> HLA-DR2b or *Aire*<sup>+/-</sup> HLA-DR4 breeders were genotyped by PCR with primers for *Aire*, *Ia*, *HLA-DR2b* and *HLA-DR4* genes as described in *Materials and Methods*. (A) Shown is the representative PCR result run on 1.8% agarose gel. Arrows indicate *Aire* knockout band of 750 bp and *Aire* Wt band of 1200 bp respectively. All the mice are either positive for HLA-DR2b or HLA-DR4 and show absence of Ia. M, marker (100 bp low scale DNA ladder); *-/-*, *Aire* knockout; *+/+*, *Aire* Wt; *+/-*, *Aire* heterozygous. (B and C) Flow cytometry analysis of splenocytes from C57BL/6 Wt, HLA-DR2b, and HLA-DR4 tg mice using fluorochrome conjugated anti-CD45, anti-Ia and anti-HLA-DR mAb. (D) *Aire*<sup>+/+</sup> and *Aire*<sup>-/-</sup> HLA-DR2b tg or (E) *Aire*<sup>+/+</sup> and *Aire*<sup>-/-</sup> HLA-DR4 tg mice were euthanized, their thymi were harvested and RNA was extracted and used for performing qPCR against a panel of Aire-dependent genes as described in *Materials and Methods*. (F – J) Thymus and spleens of HLA-DR2b or HLA-DR4 mice were procured and stained with fluorochrome conjugated anti-CD4, CD8, CD11c, CD11b, CD25, Foxp3 mAbs followed by flow cytometry analysis as described in *Materials and Methods*. (F) Shown are representative flow plots of CD4<sup>+</sup>, CD8<sup>+</sup> and double positive thymocytes in (F) *Aire*<sup>+/+</sup> HLA-DR4 vs *Aire*<sup>-/-</sup> HLA-DR4 mice. (G, H) Quantification for percentages of CD4<sup>+</sup>, CD8<sup>+</sup> and double positive thymocytes in (G) *Aire*<sup>-/-</sup> HLA-DR2b and (H) *Aire*<sup>-/-</sup> HLA-DR4 mice in comparison with their *Aire*<sup>+/+</sup> HLA-DR littermates. (I, J) Distribution of CD4<sup>+</sup>, CD8<sup>+</sup>, CD11c<sup>+</sup>, CD11b<sup>+</sup> cells in the spleens of *Aire*<sup>-/-</sup> HLA-DR2b (I) and *Aire*<sup>-/-</sup> HLA-DR4 mice (J) in comparison with their respective *Aire*<sup>+/+</sup> HLA-DR littermates. Data are representative for two independent experiments (n= 3 – 4 mice/groups), (Mean ± SEM) \* indicates significant difference

between HLA-DR2b group and C57BL/6 group, # indicates significant difference between HLA-DR4 group and C57BL/6 group; \* indicates significant difference between *Aire*<sup>-/-</sup> group and *Aire*<sup>+/+</sup> group.

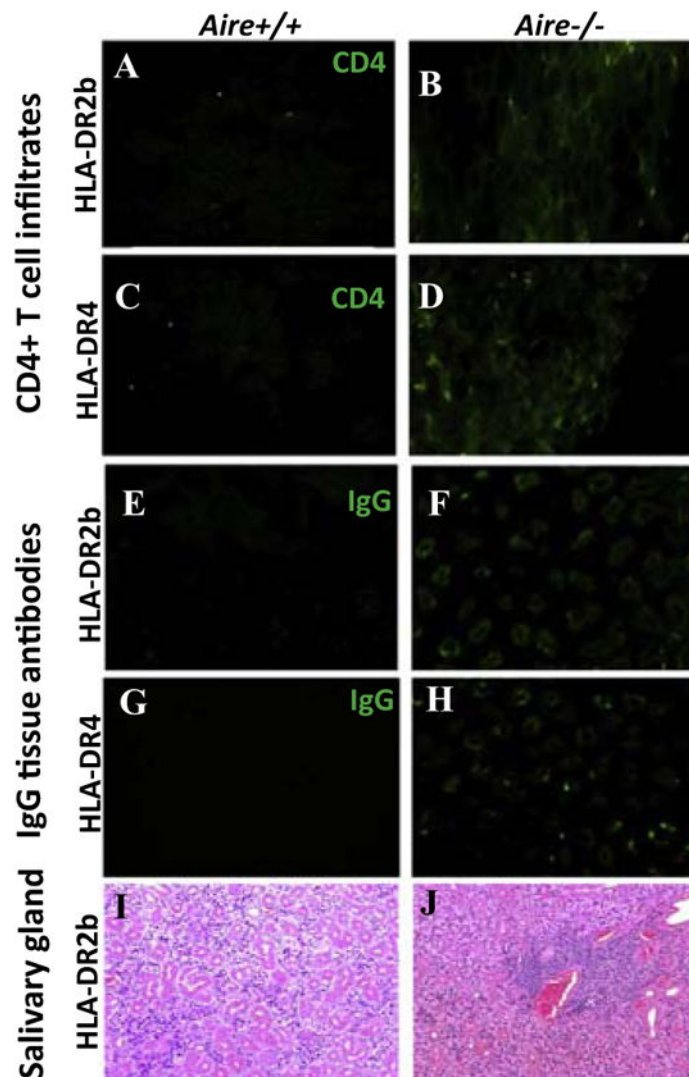
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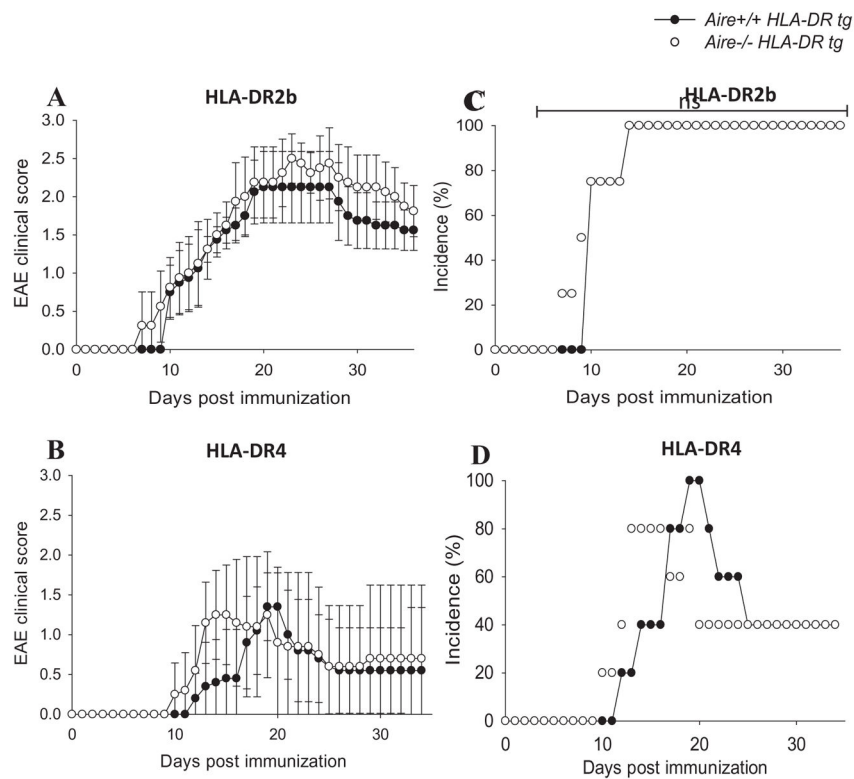
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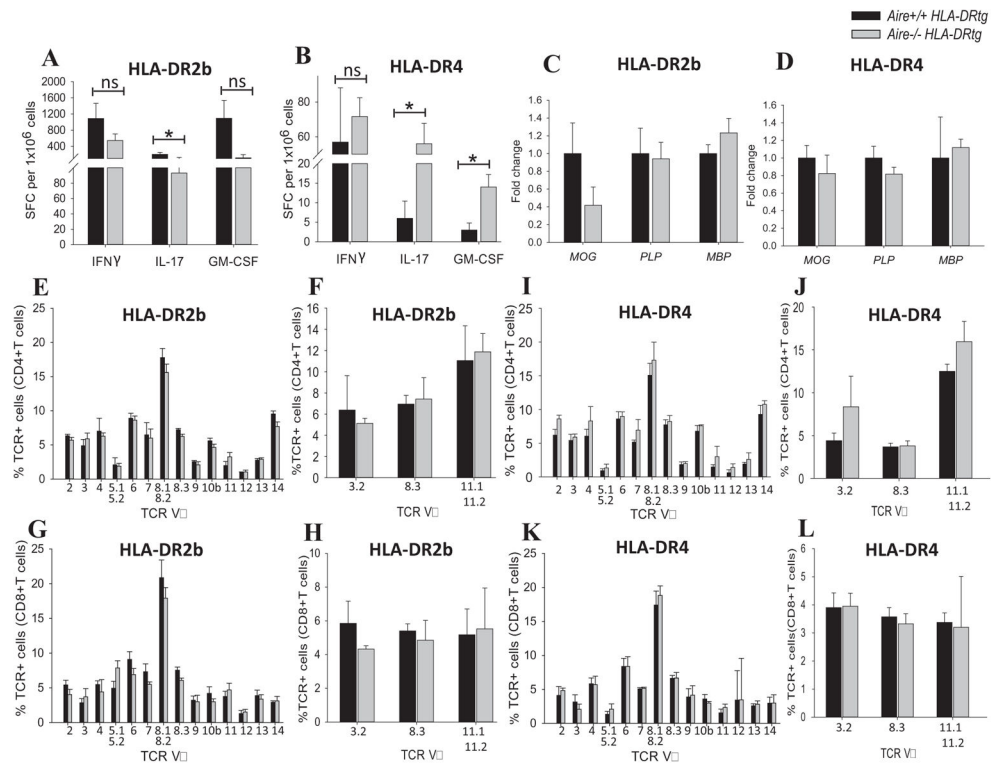


**FIGURE 2.**

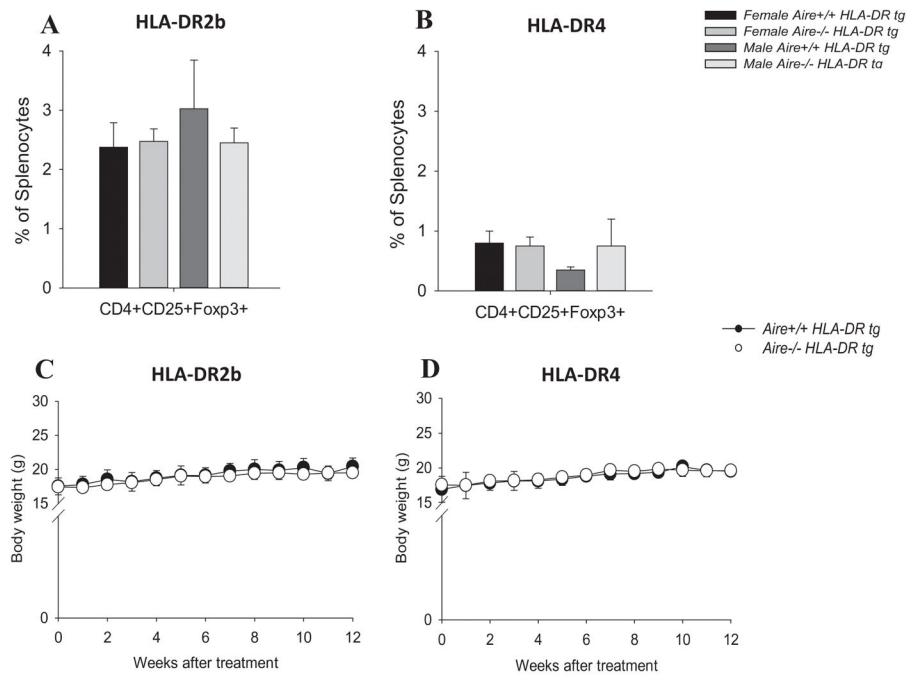
Mild spontaneous immune pathology in Aire-deficient HLA-DR2b and HLA-DR4 tg mice. (A, B) Shown are representative IF microscopy images for CD4<sup>+</sup> T cells in the liver of 13-months-old *Aire*<sup>+/+</sup> HLA-DR2b tg mice (A) versus *Aire*<sup>-/-</sup> HLA-DR2b tg mice (B). (C, D) Representative IF microscopy images for CD4<sup>+</sup> T cells in the liver of 16-months old *Aire*<sup>+/+</sup> HLA-DR4 mice (C) vs (D) *Aire*<sup>-/-</sup> HLA-DR4 mice. Magnification 200x. (E, F) Detection of autoantibodies against gastric tissues by IF microscopy images of gastric tissue sections from SCID mice in sera from 12-month-old *Aire*<sup>+/+</sup> HLA-DR2b tg mice (E) vs *Aire*<sup>-/-</sup> littermates (F), and 12-month-old *Aire*<sup>+/+</sup> HLA-DR4 mice (G) vs *Aire*<sup>-/-</sup> littermates (H). Magnification 200x. Representative H&E staining images from salivary glands of *Aire*<sup>+/+</sup> HLA-DR2b tg mice (I) vs *Aire*<sup>-/-</sup> HLA-DR2b tg mice (J). Magnification 100x.

**FIGURE 3.**

Earlier onset and mild increase in EAE severity in *Aire*-deficient HLA-DR tg mice. *Aire*<sup>+/+</sup> and *Aire*<sup>-/-</sup> HLA-DR2b tg mice (A and C) or HLA-DR4 tg mice (B and D) were immunized for EAE and scored for clinical signs of disease as described in *Materials and Methods*. Data are the representative of 3–4 independent experiments (n= 3 – 4 mice/group; Mean ± SEM).

**FIGURE 4.**

Antigen-specific cytokine production, myelin-specific gene expression, and T cell receptor repertoire unaltered in *Aire*-deficient HLA-DR tg mice. Mice immunized for EAE were euthanized during disease remission (day 34 – 36), splenocytes of *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b tg (A) or HLA-DR4 tg mice (B) were harvested and recalled with peptides MOG35–55 (HLA-DR2b tg) or MOG97–108 (HLA-DR4 tg) and tested by cytokine ELISPOT assay to quantify IFN- $\gamma$ , IL-17 and GM-CSF producing cells as described in *Materials and Methods*. Shown is mean  $\pm$  SEM of the number of cytokine forming cells. Data are the representative of 3–4 independent experiments (n = 3 – 4 mice/group; \* indicates significant difference between *Aire*<sup>-/-</sup> group and *Aire*<sup>+/+</sup> group). C) naïve *Aire*<sup>+/+</sup> and *Aire*<sup>-/-</sup> HLA-DR2b or (D) naïve *Aire*<sup>+/+</sup> and *Aire*<sup>-/-</sup> HLA-DR4 mice were euthanized, their thymus was harvested, and RNA was extracted and used for performing qPCR against myelin-specific genes (MOG, PLP, MBP) as described in *Materials and Methods*. Data are representative for two independent experiments (n = 3 – 4 mice/groups), (mean  $\pm$  SEM of mRNA expression). (E–H) Shown is the flow cytometry analysis of TCR V $\beta$  and V $\alpha$  distribution of splenic CD4<sup>+</sup> (E, F) and CD8<sup>+</sup> T cells (G, H) of *Aire*<sup>-/-</sup> vs *Aire*<sup>+/+</sup> HLA-DR2b tg mice. (I–L) Flow cytometry analysis of TCR V $\beta$  and V $\alpha$  distribution of splenic CD4<sup>+</sup> (I, J) and CD8<sup>+</sup> T cells (K, L) of *Aire*<sup>-/-</sup> vs *Aire*<sup>+/+</sup> HLA-DR4 tg mice. Flow cytometry analysis was performed using panels of mAbs for TCR families as described in *Materials and Methods*. Shown are pooled data of two independent experiments (n = 2 – 4 mice/group, Mean  $\pm$  SEM).

**FIGURE 5.**

No significant increase in autoimmune pathology in *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b or HLA-DR4 mice after CTLA-4 and/or CD25 blockade. Spleens of naïve HLA-DR2b or HLA-DR4 mice were procured and stained with fluorochrome conjugated anti-CD4, CD25, Foxp3 mAbs followed by flow cytometry analysis as described in *Materials and Methods*. (A, B) Distribution of CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup> cells in the spleens of *Aire*<sup>-/-</sup> HLA-DR2b (A) and *Aire*<sup>-/-</sup> HLA-DR4 mice (B) in comparison with their respective *Aire*<sup>+/+</sup> HLA-DR littermates. (C) *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b mice treated with anti-CTLA-4 and anti-CD25 mAbs, or (D) *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR4 mice treated with anti-CTLA-4 mAb were weighed and no significant differences were found between the body weights of *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR tg mice (t-test). Data are representative of two independent experiments (n = 3 – 4 mice/group, Mean ± SEM).

**Table 1**

Scoring table for tissue sections obtained from *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice.

Score	Average no. of lesions/section (2–3 sections per slide)
0	No lesions at all
1	1–4 lesions
2	5–15 lesions
3	15–25 lesions

The tissue sections were obtained from the *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg with 2 slides per mouse (staggered sections ensuring spanning across the organ). The scoring was performed based on the average number of lesions per section with 2–3 sections per slide.

mice were incubated with tissue sections of organs from C57BL/6 SCID mice, followed by staining with FITC goat anti-mouse IgG Ab (Jackson ImmunoResearch).

Incidence of lymphocytic infiltrates in the organs of *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice.

**Table 2**

Organ	CD4 <sup>+</sup> T cells					
	CD4 <sup>+</sup> T cells			CD8 <sup>+</sup> T cells		
	Liver	Pancreas		Liver	Pancreas	
Genotype	Male	Female	Male	Female	Male	Female
<i>Aire</i> <sup>+/+</sup> HLA-DR2b mice	0/5	0/3	0/5	0/3	0/5	0/3
<i>Aire</i> <sup>-/-</sup> HLA-DR2b mice	1/11	3/7	1/11	0/7	0/11	1/7
<i>Aire</i> <sup>+/+</sup> HLA-DR4 mice	0/7	0/4	0/7	0/4	0/7	0/4
<i>Aire</i> <sup>-/-</sup> HLA-DR4 mice	0/8	3/7	0/8	1/7	0/8	1/7

Immune infiltration was examined in *Aire*<sup>-/-</sup>, *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice by I.F. staining on tissue sections from different organs (liver, lung, pancreas, stomach and intestine) with anti-CD4 and anti-CD8 mAbs as described in *Materials and Methods*. Infiltrates were mostly observed in liver and pancreas of a few aged *Aire*<sup>-/-</sup> HLA-DR tg mice. Shown are the incidences of mice that have immune infiltrates found in indicated organs (n = 3–11).



**Table 3**

Incidence of IgG autoantibodies in the sera of *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice.

Genotype	HLA-DR2b		HLA-DR4	
	Male	Female	Male	Female
<i>Aire</i> <sup>+/+</sup> mice	1/3	0/2	0/7	1/3
<i>Aire</i> <sup>-/-</sup> mice	2/10	2/9	2/10	2/8

IgG autoantibodies detected in the sera of *Aire*<sup>+/+</sup> and *Aire*<sup>-/-</sup> HLA-DR2b or HLA-DR4 tg mice which was stained against a panel of organs (stomach, lung, liver, bowel, pancreas and kidney) from SCID mice. IgG binding was observed in the gastric tissue. Shown are the incidences of mice that have IgG auto-antibodies (n = 3–10).

Incidence of immune cell infiltrates in *Aire*<sup>-/-</sup> and *Aire*<sup>+/+</sup> HLA-DR2b and HLA-DR4 tg mice injected with anti-CTLA-4 + anti-CD25 and anti-CTLA-4 respectively.

**Table 4**

Genotype	HLA-DR2b (anti-CTLA-4 + anti-CD25)		HLA-DR4 (anti-CTLA-4)	
	Liver	Pancreas	Liver	Pancreas
	<b>CD4<sup>+</sup></b>	<b>CD8<sup>+</sup></b>	<b>CD4<sup>+</sup></b>	<b>CD8<sup>+</sup></b>
<i>Aire</i> <sup>+/+</sup> mice	0/4	0/4	1/3	0/3
<i>Aire</i> <sup>-/-</sup> mice	0/5	0/5	3/7	1/7

Staining was performed using AF488 anti-CD4 mAb and APC anti-CD8 mAb against a panel of organs (liver, lungs, pancreas, stomach and intestine). Shown are the incidences of mice (injected with anti-CTLA-4 and CD-25 mAb) that have immune infiltrates found in indicated organs (n = 3–7).