


# Treatment of Allergic Asthma with Fenretinide Formulation (LAU-7b) Downregulates ORMDL Sphingolipid Biosynthesis Regulator 3 (*Ormdl3*) Expression and Normalizes Ceramide Imbalance<sup>[S]</sup>

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## ABSTRACT

Zona pellucida binding protein 2 (*Zbp2*) and ORMDL sphingolipid biosynthesis regulator 3 (*Ormdl3*), mapped downstream of *Zbp2*, were identified as two genes associated with airway hyper-responsiveness (AHR). *Ormdl3* gene product has been shown to regulate the biosynthesis of ceramides. Allergic asthma was shown to be associated with an imbalance between very-long-chain ceramides (VLCCs) and long-chain ceramides (LCCs). We hypothesized that Fenretinide can prevent the allergic asthma-induced augmentation of *Ormdl3* gene expression, normalize aberrant levels of VLCCs and LCCs, and treat allergic asthma symptoms. We induced allergic asthma by house dust mite (HDM) in A/J WT mice and *Zbp2* KO mice expressing lower levels of *Ormdl3* mRNA than WT. We investigated the effect of a novel formulation of Fenretinide, LAU-7b, on the AHR, inflammatory cell infiltration, mucus production, IgE levels, and ceramide levels. Although lower *Ormdl3* expression, which was observed in *Zbp2* KO mice, was associated with lower AHR, allergic *Zbp2* KO mice were not protected from inflammatory cell infiltration, mucus accumulation, or aberrant levels of VLCCs and LCCs induced by

HDM. LAU-7b treatment protects both the *Zbp2* KO and WT mice. The treatment significantly lowers the gene expression of *Ormdl3*, normalizes the VLCCs and LCCs, and corrects all the other phenotypes associated with allergic asthma after HDM challenge, except the elevated levels of IgE. LAU-7b treatment prevents the augmentation of *Ormdl3* expression and ceramide imbalance induced by HDM challenge and protects both WT and *Zbp2* KO mice against allergic asthma symptoms.

## SIGNIFICANCE STATEMENT

Compared with A/J WT mice, KO mice with *Zbp2* gene deletion have lower AHR and lower levels of *Ormdl3* expression. The novel oral clinical formulation of Fenretinide (LAU-7b) effectively lowers the AHR and protects against inflammatory cell infiltration and mucus accumulation induced by house dust mite in both *Zbp2* KO and WT A/J mice. LAU-7b prevents *Ormdl3* overexpression in WT allergic mice and corrects the aberrant levels of very-long-chain and long-chain ceramides in both WT and *Zbp2* KO allergic mice.

## Introduction

Allergic asthma causes global health burdens because it affects millions of people worldwide. Being a complex disease, in addition to environmental exposures, several genetically controlled factors greatly influence the predisposition and severity of allergic asthma (Burbank et al., 2017). Genome-wide association studies have highlighted the 17q21 locus, which contains several susceptibility genes, including zona pellucida binding protein 2 (*ZPBP2*) and ORMDL sphingolipid biosynthesis regulator 3 (*ORMDL3*) on human chromosome 17 and their orthologs *Zbp2* and *Ormdl3* on mouse chromosome 11, as a well established susceptible locus for

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**ABBREVIATIONS:** AA, arachidonic acid; AHR, airway hyper-responsiveness; CF, cystic fibrosis; DHA, docosahexaenoic acid; HDM, house dust mite; *Il-4*, *Interleukin-4*; KO, knockout; LCC, long-chain ceramide; MCh, methacholine; MD, malondialdehyde; NT, nitrotyrosine; *Ormdl3*, ORMDL sphingolipid biosynthesis regulator 3; PAS, periodic acid-Schiff; Penh, enhanced pause; VLCC, very-long-chain ceramide; WBP, whole-body plethysmography; WT, wild-type; *Zbp2*, zona pellucida binding protein 2.

airway hyper-responsiveness (AHR) (Moffatt et al., 2007). The importance of 17q21 locus was evaluated in an ethnically diverse population, but differences based on ethnicity were not found (Kothari et al., 2018).

ZBP2 localizes to the sperm acrosome, facilitating the binding of the spermatozoa to the oocyte's zona pellucida, so it is highly expressed in testes in both mice and humans (Torabi et al., 2017). ZBP2 expression has also been detected in lower levels in human somatic airway epithelial cells (Moussette et al., 2017). Recently, our group (Kanagaratham et al., 2018) and Miller et al. (2018) have published that *Zbp2* gene deletion attenuated AHR in C57BL/6 mice; nevertheless, the functional impact of *Zbp2* on allergic asthma remains undiscovered. *Ormdl3*, which is involved in modulating the biosynthesis of ceramides, has been mapped downstream of *Zbp2*, and it was recently shown that the deletion of *Zbp2* in C57BL/6 mice led to a decrease in the level of *Ormdl3* expression (Miller et al., 2018). Compared with ZBP2, more studies have been published pertinent to ORMDL3 and allergic asthma. It has been shown that increased expression of *Ormdl3* positively correlates with airway hyper-reactivity (Davis et al., 2018). Transcriptomic and metabolomic analyses have linked the forced expiration volume in the first second / forced vital capacity (FEV1/FVC) ratio with ORMDL3 and dysregulated lipid metabolism in a study of asthmatic children (Kelly et al., 2018).

Moreover, elevated levels of ORMDL3 inhibit sphingolipid biosynthesis and result in inhibition of long-chain ceramides (LCCs; C16:0) and four species of very-long-chain ceramides (VLCCs; C22:0, C24:0, C24:1, and C24:2) (Kiefer et al., 2019). Debeuf et al. (2019) reported that transgenic mice over-expressing *Ormdl3* had significantly lower levels of VLCCs (C24:0 and C24:1); by contrast, loss of *Ormdl3* in knockout (KO) mice resulted in elevated levels of C24:0 and C24:1. Besides allergic asthma, several studies showed that the *ORMDL3* locus is also associated with other numerous pathologies (Das et al., 2017), e.g., allergic rhinitis (Tomita et al., 2013), type 1 diabetes (Saleh et al., 2011), primary biliary cirrhosis (Mells et al., 2011), rheumatoid arthritis (Kurreena et al., 2012), ulcerative colitis (Anderson et al., 2011), Crohn disease (Franke et al., 2010), and ankylosing spondylitis (Laukens et al., 2010).

Fenretinide, a vitamin A derivative, corrects the aberrant inflammatory responses and improves lung functions in both cystic fibrosis (CF) (Guilbault et al., 2008, 2009, Youssef et al., 2016, 2020; Garić et al., 2017) and allergic asthma (Kanagaratham et al., 2014; Youssef et al., 2020). In CF, our laboratory has shown that Fenretinide improves lung function by elevating levels of VLCCs and lowering levels of LCCs (Garić et al., 2017). Recently, a novel clinical oral formulation of Fenretinide (LAU-7b) with excellent bioavailability was tested using daily single oral capsules (100, 200, and 300 mg) in patients with CF in a phase Ib clinical trial study (NCT02141958). The results of a phase I clinical trial demonstrated very promising pharmacokinetics and pharmacodynamics of LAU-7b. Currently, LAU-7b is being tested in a phase II clinical trial in patients with CF (NCT03265288). However, LAU-7b capsules have not yet been tested in patients suffering from allergic asthma.

In our study, we developed KO mice for *Zbp2* gene on the A/J genetic background. Unlike C57BL/6, the A/J strain of mice is not only atopic but also expresses genes regulating

AHR. Allergic asthma developed in this strain of mice is characterized by a very strong inflammatory response, a significant increase in airway hyperplasia, and an intense AHR, which can be measured after exposure to methacholine (MCh). Therefore, we have used this strain of mice to investigate the protective effects of LAU-7b treatment (10 mg/kg per day for 9 days) against HDM-induced asthma. We aimed to validate the importance of the *Zbp2* gene in allergic asthma, so we hypothesized that A/J KO mice with the *Zbp2* gene deletion would be associated with decreased AHR measured after exposure to MCh. Because the *Ormdl3* gene product regulates the biosynthesis of ceramides, we expected that changes in *Ormdl3* gene expression will correlate with changes in the distribution of specific species of ceramides. We hypothesized that the allergic asthma-induced augmentation of *Ormdl3* can be prevented by treatment with LAU-7b in the A/J hyper-responsive strain of mice. Furthermore, we hypothesized that there is an association between the LAU-7b-induced improvement in lung physiology and the normalization of relative abundance of VLCCs and LCCs in the lungs.

## Materials and Methods

**Animal Model.** Heterozygous B6.129S7-*Zbp2*<sup>tm1Zuk</sup>/J mice (Lin et al., 2007), which carry a deletion of *Zbp2* exons 1–3, were purchased from the Jackson Laboratory (Bar Harbor, ME) and backcrossed for more than 10 generations (N10) to A/J inbred mice purchased from Jackson Laboratory. N10 *Zbp2* KO mice were intercrossed to generate homozygous KO A.129S7-*Zbp2*<sup>tm1Zuk</sup> mice, referred to from this point on as KO. All experimental procedures were approved by the Animal Care Committee of McGill University Health Center, Montreal, QC, Canada. More information is provided in the Supplemental Material.

**Mouse Groups and Batches.** After sensitization (more information is provided in the Supplemental Material), the mice were split into the following groups: allergic, PBS-treated, unchallenged mice (HDM-PBS-PBS); allergic, PBS-treated, challenged mice (HDM-PBS-HDM); and allergic, LAU-7b-treated, challenged mice (HDM-LAU-7b-HDM). For simplification, the three groups are presented as “PBS,” “HDM,” and “LAU-7b,” respectively. Eight different experiment batches using *Zbp2* WT and KO mice on the A/J background ( $n = 217$ ) were tested. The results collected from all eight experiments were pooled together to make the final graphs.

**AHR Measurements.** AHR was measured using a Buxco plethysmograph system, ventilators, and nebulizers (Harvard Apparatus, Holliston, MA) as previously described (Kanagaratham et al., 2014, 2018). For the noninvasive whole-body plethysmography (WBP), the baseline enhanced pause (Penh) values were measured without anesthesia at the age of 8 weeks before the first sensitization. The invasive AHR measurements were done when sacrificing the mice on the day of the harvest. A nebulizer was used in both noninvasive and invasive lung experiments to administer ascending doses of MCh (acetyl  $\beta$ -methyl choline, Cat: A2251; Sigma Aldrich, Saint Louis, MO).

**Lung Histology Analysis.** H&E staining was used to assess lung tissue recruitment of different inflammatory cells around airways. Quantifying LAU-7b effects was done by counting the number of infiltrated inflammatory cells among at least four airways/mice, averaged, and normalized versus the perimeter of the airway basement membrane as previously described (Kanagaratham et al., 2014). Periodic acid–Schiff (PAS) stain was used to visualize goblet cell hyperplasia in the lungs. PAS-positive cells in the airways were counted and normalized by dividing the counts by the perimeter of the basement membrane. Per mouse, at least four airways were counted as previously described (Kanagaratham et al., 2014). Airway smooth

muscle mass was stained using specific  $\alpha$ -smooth muscle actin antibody as previously described by our laboratory (Camateros et al., 2007). Masson trichrome stain was used to assess collagen production as previously described (Chen et al., 2015).

**Lipids and Markers of Oxidation Analysis.** Lipid analysis was done using 25 mg of macerated lung tissue from each mouse. Classic isolation of lipids was done as previously described by Folch et al. (1957), and the levels of different lipid species were measured using high-performance liquid chromatography tandem mass spectrometry as previously described (Guilbault et al., 2008).

**Gene Expression Measurements.** RNA was extracted from snap-frozen lungs using the RNeasy Mini kit (Qiagen). In total, 500 ng of RNA was reverse transcribed into cDNA for each analyzed sample using the QuantiTect Reverse Transcription kit (Qiagen). Levels of expression of *Ormdl3* mRNA were measured by using the CFX384 Touch Real-Time PCR Detection System and SsoFast EvaGreen Supermix (BioRad). Primers were designed using the primer Basic Local Alignment Search Tool (BLAST) online software, National Center for Biotechnology Information (NCBI). More information is provided in the Supplemental Material.

**IgE Measurements in Serum.** Plasma IgE levels were measured using an ELISA IgE mouse kit (BD OptEIA Biosciences) following the manufacturer's instructions.

**Statistical Analysis.** Data were pulled out from at least three independent experiments for each analysis. Data were analyzed using GraphPad Prism 6 (version 6.01; GraphPad Software, Inc., San Diego, CA). An ANOVA test was used for analyzing the results of more than two groups, and a *t* test was used for analyzing the results of two groups. *P* values of less than 0.05 were considered statistically significant. The number of mice (*n*) used for each analysis is written in each figure caption.

## Results

### Gene Expression and Basal Lung Function Analysis.

We confirmed that, by removing exons 1–3 of the 5' region of *Zbp2*, the mRNA gene expression was abolished (Fig. 1, A and B) using lung tissue collected from the A/J *Zbp2* KO mice we developed. It has been previously reported (Miller et al., 2018) that deletion of *Zbp2* has downregulated *Ormdl3* expression in lung epithelial cell lines. Therefore, we wanted to evaluate whether *Ormdl3* gene expression might be decreased in the lungs of *Zbp2* KO A/J mice. In WT control mice, PBS-challenged mice demonstrated very low gene expression of *Ormdl3* compared with the gene expression of HDM-challenged mice (Fig. 1, A and B). Furthermore, in *Zbp2* KO mice, *Ormdl3* gene expression increase even after HDM challenges was also abolished (half-fold expression in *Zbp2* KO mice compared with 2- to 4-fold expression in WT mice) (Fig. 1, A and B). Then, we treated the WT mice with LAU-7b, and we measured *Zbp2* and *Ormdl3* gene expression. Our data show that LAU-7b treatment significantly decreased the expression of *Ormdl3* in both WT males and females compared with HDM-challenged and PBS-treated mice (Fig. 1D).

We also tested the gene expression of four of the T helper 2 (Th2) immune pathway-associated genes (Fig. 1, E–H): *Interleukin-4*; *Il-4* (differentiates naïve T cells; Th0 into Th2 cells (Steinke and Borish, 2001)), *Il-5* (activates eosinophils (Farne et al., 2017)), *Il-13* that regulates the production of IgE (Rael and Lockey, 2011), and *C-C Motif Chemokine Ligand 11*; *Ccl11*, also known as *Eotaxin-1* (enhances the infiltration of eosinophils into the airways (Williams, 2015)). Our results show that after HDM sensitization and challenge, the expression of *Il-5*, *Il-13*, and *Eotaxin-1* was significantly elevated in

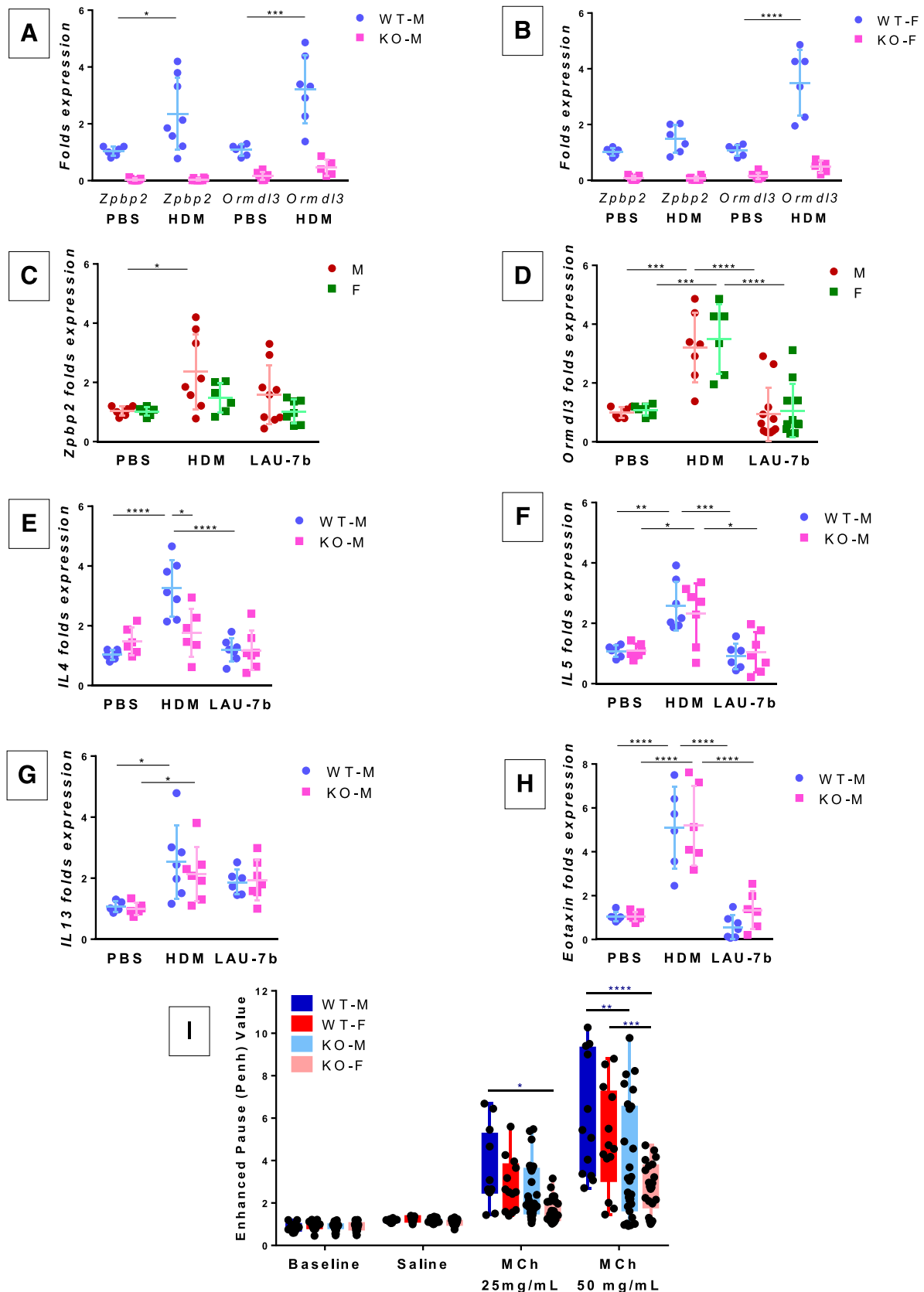
WT and *Zbp2* KO mice; meanwhile, the expression of *Il-4* was significantly elevated in WT but not *Zbp2* KO mice. LAU-7b-treated WT mice have significantly lower expression of *Il-4*, *Il-5*, and *eotaxin-1* genes but not *Il-13* compared with littermate WT controls. Noninvasive WBP was used to measure baseline respiratory functions before starting HDM sensitization and without sacrificing the mice (Fig. 1I). At a MCh dose of 25 mg/ml, both WT males and females have significantly higher lung Penh values than KO males and females, respectively.

**Assessment of AHR and IgE Levels.** The results of WBP were confirmed by classic invasive measurement of lung resistance on the day of harvest (Fig. 2, A–D). HDM challenge significantly increased the lung resistance values of all mouse groups (males, females, WT, and KO) at MCh doses of 50 and 100 mg/ml compared with PBS-challenged mice. Consistent with the previously published data (Card et al., 2006; Antunes et al., 2010), our data demonstrate that WT males have higher lung resistance than WT females; similarly, KO males have higher lung resistance than KO females (Fig. 2, A and B). Lung responsiveness to nebulized MCh revealed that WT (male and female) and KO (male) mice gavaged with 10 mg/kg LAU-7b for 9 days had significantly lower lung resistance than PBS-treated mice (Fig. 2, C and D). Allergic KO females display a very low increase in lung resistance after HDM challenge (nonsignificant from PBS), and the treatment does not improve the lung physiology any further, which under these circumstances, is not surprising in this group of mice.

The protective effects of LAU-7b against increased airway hyper-responsiveness after allergen challenge prompted us to evaluate its potential against IgE production, which is also associated with allergic asthma. After HDM challenge, the titer of IgE was increased by 4- to 10-fold in WT and KO mice (Fig. 2E). IgE measurements show that WT and KO male and female groups were not statistically significant from one another. Nonetheless, LAU-7b treatment did not lower the levels of IgE caused by HDM challenge in WT and KO male and female mice compared with PBS-treated and HDM-challenged mice.

**Evaluation of Inflammatory Cell Infiltration, Mucus Production, Smooth Muscle Mass, and Collagen Production in the Airways.** To visualize and quantify the inflammatory cells' infiltration into the airways after HDM challenge, we used H&E staining (Fig. 3). The lung sections of both WT and KO mice showed significant incoming inflammatory cells after HDM challenge compared with PBS-challenged mice (Fig. 3, A–L). We also quantified our H&E staining results by counting and normalizing the recruited cells around the lung airways (Fig. 3M). Our data demonstrated that there was equally strong inflammatory cell infiltration into the airways of both WT and KO male and female mice after HDM allergen challenge, even though KO mice had displayed much less lung resistance than WT mice. As shown in the representative pictures from each mouse group, LAU-7b treatment significantly lowered the recruitment of inflammatory cells in both WT and KO male and female mice.

Mucus hypersecretion, and the subsequent plugging of the airways, has long been recognized as a common phenotype of allergic asthma. The mucus production was not investigated before in the *Zbp2* KO mouse model, so in our study, we wanted to examine whether the deletion of this gene would



**Fig. 1.** *Zbp2* and *Ormdl3* gene expression and basal lung function in WT vs. *Zbp2* KO mice fold-change expression of *Zbp2* and *Ormdl3* genes quantitated using qPCR for WT and *Zbp2* KO (A) male mice and (B) female mice. Fold-change expression of (C) *Zbp2* and (D) *Ormdl3* genes quantitated using qPCR for LAU-7b-treated WT male mice and female mice. LAU-7b treatment and fold-change expression of (E) *IL-4*, (F) *IL-5*, (G) *IL-13*, and (H) *Eotaxin-1* genes in *Zbp2* KO and WT mice. All mice were sensitized with HDM and challenged with either PBS or HDM. Treatment group is marked as LAU-7b. (I) Noninvasive WBP for WT and *Zbp2* KO mice. Penh values were measured in response to increasing doses of inhaled MCh in

result in any effects that could be visualized by the PAS stain (Fig. 4) and corrected with LAU-7b treatment. Our lung sections of both WT and KO mice markedly show mucus production by goblet cells after HDM challenge compared with PBS-challenged mice (Fig. 4, A–L). No significant differences between WT and KO male or female mice were observed. LAU-7b treatment significantly decreased the production of mucus by goblet cells in both WT and KO male and female mice (Fig. 4M).

As remodeling of the airways is an important aspect in the context of human asthma, we wanted to assess the smooth muscle mass and collagen production in our mouse model and evaluate the effect of treatment with LAU-7b. Our data (Supplemental Fig. 1) show that after HDM challenge, there is a slight increase in the mass of the smooth muscles of the mouse airways. Similarly, challenge with HDM results in the production of collagen (Supplemental Fig. 2), which was evident in the airways. LAU-7b treatment lowered both the mass of the smooth muscles and the collagen production in our KO and WT A/J mice.

**Analysis of Ceramides, Fatty Acids, and Markers of Oxidation.** We evaluated the levels of malondialdehyde (MD), a marker of lipid oxidation, and nitrotyrosine (NT), a marker of protein oxidation, in our mice because they are markers of cellular stress and damage, which happen after allergen challenge. After HDM challenge, both MD and NT significantly increased in male and female KO and WT mice (Fig. 5, A and B). No significant differences were observed between males and females or KO and WT mice. LAU-7b significantly normalized the levels of MD and NT in the HDM-challenged and treated mice compared with HDM-challenged and untreated mice.

Furthermore, fatty acid analysis (Fig. 5C) reveals significant elevation in the arachidonic acid/docosahexaenoic acid ratio (AA/DHA ratio) after HDM challenge, which is typically associated with inflammation. Nevertheless, this ratio is significantly lowered after treatment with LAU-7b in HDM-sensitized and -challenged mice compared with the untreated HDM-sensitized and -challenged mice. Likewise, no significant differences in the AA/DHA ratio were observed between males and females or KO and WT mice.

The lipidomic analysis of the lungs (Fig. 5, D and E; Supplemental Fig. 3) revealed that the relative levels of VLCCs (C22:0, C22:1, C24:0, C24:1, C26:0, C26:1) were diminished in HDM-challenged KO mice (females = 24.07%, lower than males = 34.45%) compared with PBS-challenged KO mice (females = 34.44%, lower than males = 37.93%). LAU-7b-treated KO mice displayed higher levels of VLCCs compared with untreated KO mice in both males and females: 40.70% and 40.24%, respectively. In KO mice, treatment with LAU-7b resulted in significantly elevated levels of C24:0, C24:1, C26:0, and C26:1 in males and C24:0 in females. Similarly, the treatment with LAU-7b resulted in significantly reduced levels of LCCs: C14:0 and C18:0 in KO males and C14:0, C16:0, and C18:0 in KO females.

Moreover, compared with KO mice, the assessment of VLCCs demonstrated that WT male and female mice have higher levels of VLCCs (Supplemental Fig. 3). Total VLCC

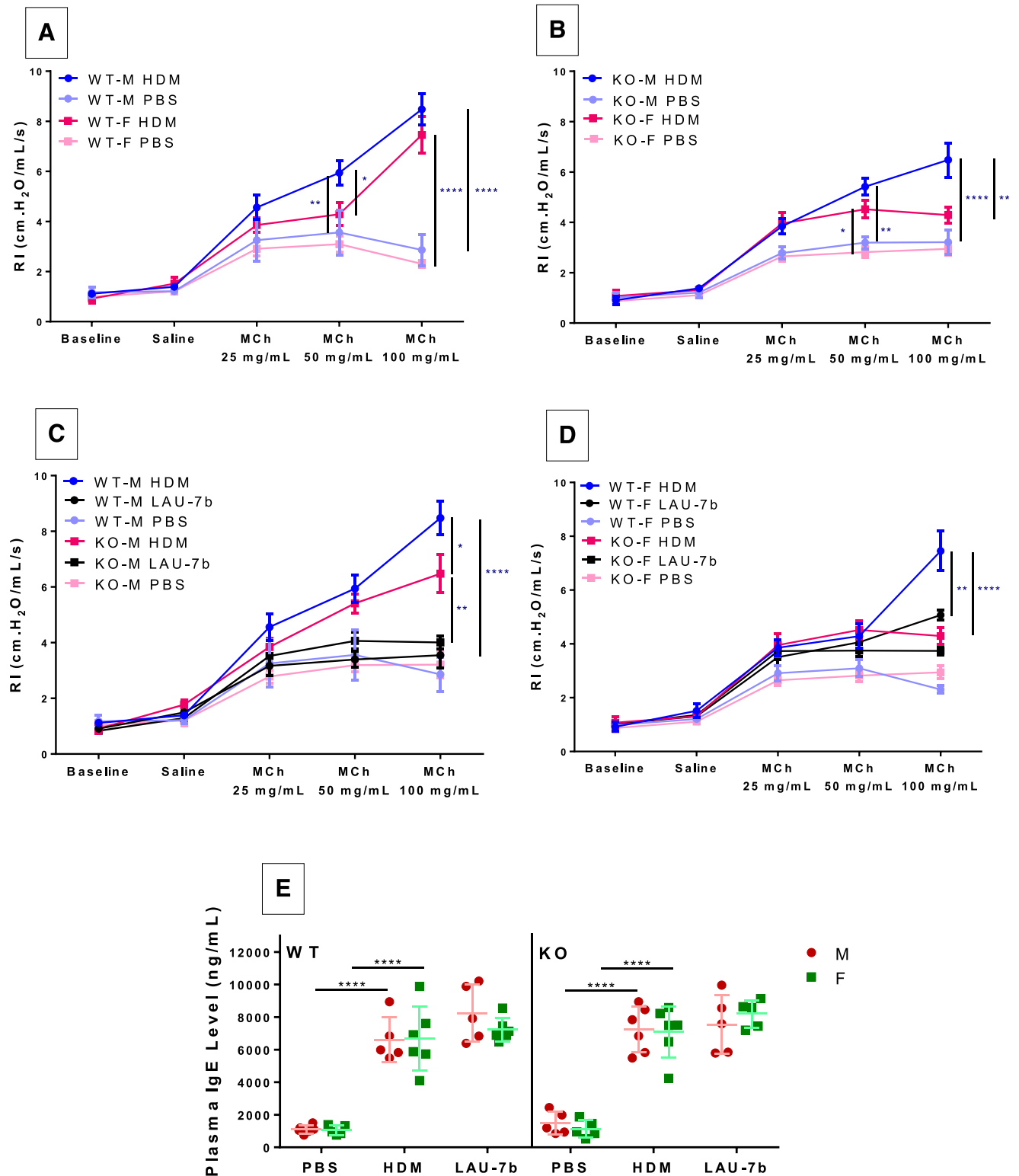
levels in PBS-challenged WT mice demonstrated a percentage of 41.85% in males and 43.07% in females. After challenge with HDM, the total VLCC levels were reduced to 39.92% in males and 31.06% in females. As shown here, the VLCC values obtained from male and female KO mice were lower than the values obtained for male and female WT mice for both PBS- and HDM-challenged groups. However, as observed in KO mice, LAU-7b treatment restored the levels of VLCCs to those typically observed in WT male and female mice—up to 42.43% and 47.89%, respectively—which is even higher than the levels of VLCCs detected before HDM challenge (in PBS mouse groups).

## Discussion

The importance of the 17q21 locus, containing genes such as *ORMDL3* and *ZBPB2*, was replicated in several studies (Moffatt et al., 2007; Karunas et al., 2011; Torgerson et al., 2011; Wan et al., 2012). Our main objective was 3-fold. First, we aimed to evaluate the effect of the *Zbp2* gene deletion on lung physiology. Secondly, we aimed to investigate the effects of LAU-7b treatment against HDM-induced allergic asthma in *Zbp2* KO and WT A/J mice. Our third objective was to test the expression of *Zbp2* and *Ormdl3* genes after LAU-7b treatment to evaluate whether there might be an association between the modulation of the expression of these genes and the changes in the regulation of relative ratios of VLCCs and LCCs.

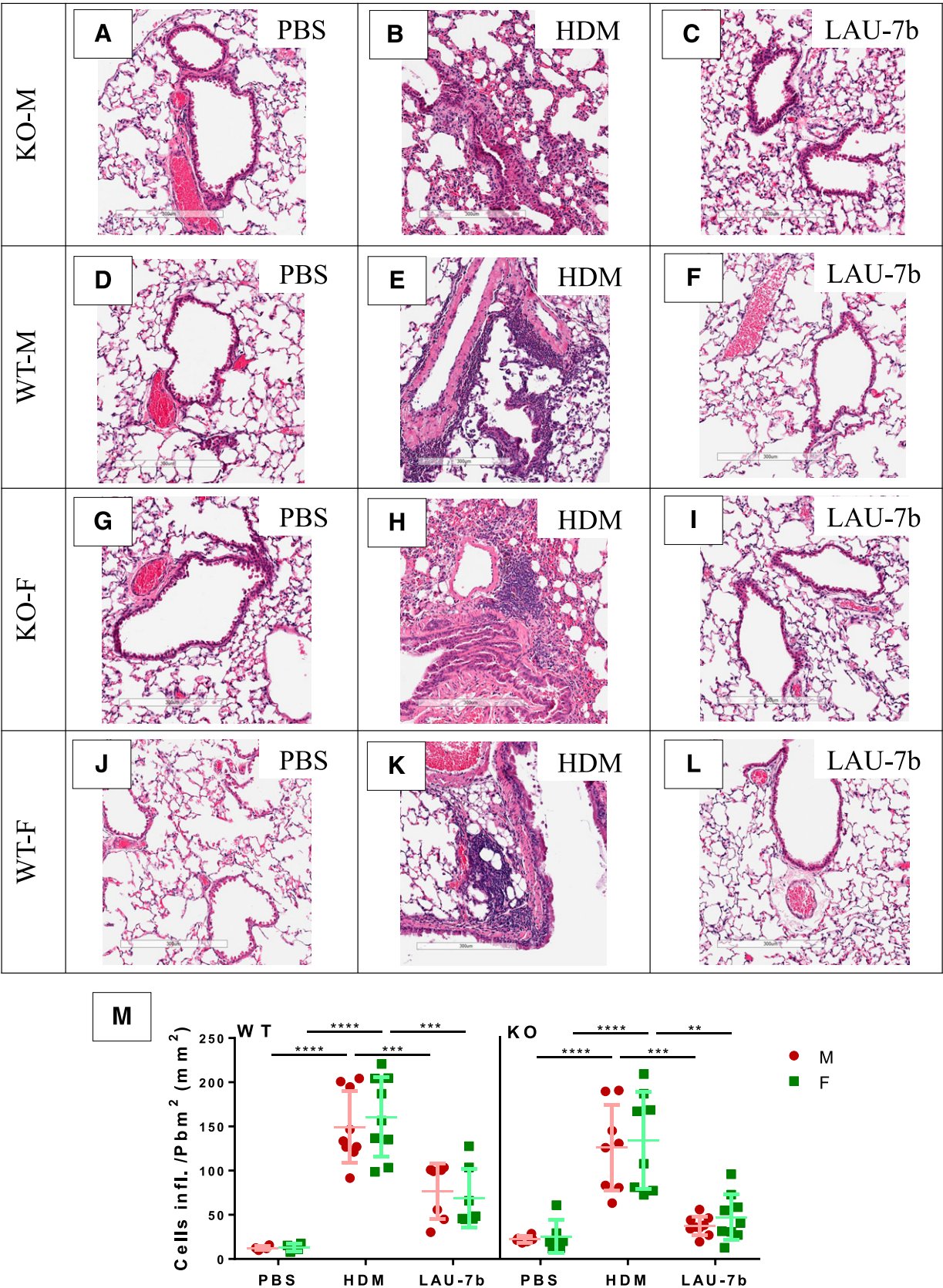
We have previously published (Kanagaratham et al., 2018) that the deletion of the *Zbp2* gene in C57BL/6 mice resulted in a reduction of AHR in females but not males on baseline levels. However, the deletion of the *Zbp2* gene in our previous study (Kanagaratham et al., 2018) did not significantly affect the AHR of male or female mice after ovalbumin sensitization and challenge, perhaps because the C57BL/6 strain of mice is genetically resistant to developing allergic asthma. Miller et al. (2018) similarly reported that *Zbp2* KO C57BL/6 male and female mice challenged with HDM had significantly reduced AHR compared with WT controls. Our results here demonstrated that deletion of the *Zbp2* gene on the A/J background has resulted in a significant reduction of baseline Penh values in male and female mice (Fig. 1I) compared with WT controls. Similarly, we observed a significant reduction in AHR (shown by the lung resistance values, Fig. 2, A and B) of *Zbp2* KO male and female mice compared with WT mice. By using an A/J hyper-responsive strain of mice and inducing allergic asthma by HDM, we have obtained a good segregation of different mouse groups (KO and WT, males and females) in terms of AHR. Our data demonstrate for the first time that in the genetically very susceptible (high AHR) and atopic mice, this gene plays an important role in the control of lung physiology in allergic asthma.

HDM sensitization and challenge significantly enhanced the recruitment of inflammatory cells into the lungs (Fig. 3), caused hyperplasia of the lining of the airways (Fig. 3), and increased the production of mucus (Fig. 4) in *Zbp2* KO and

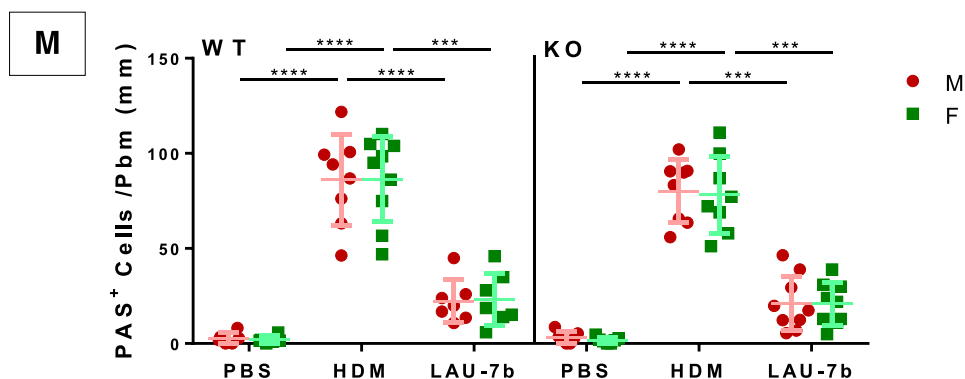
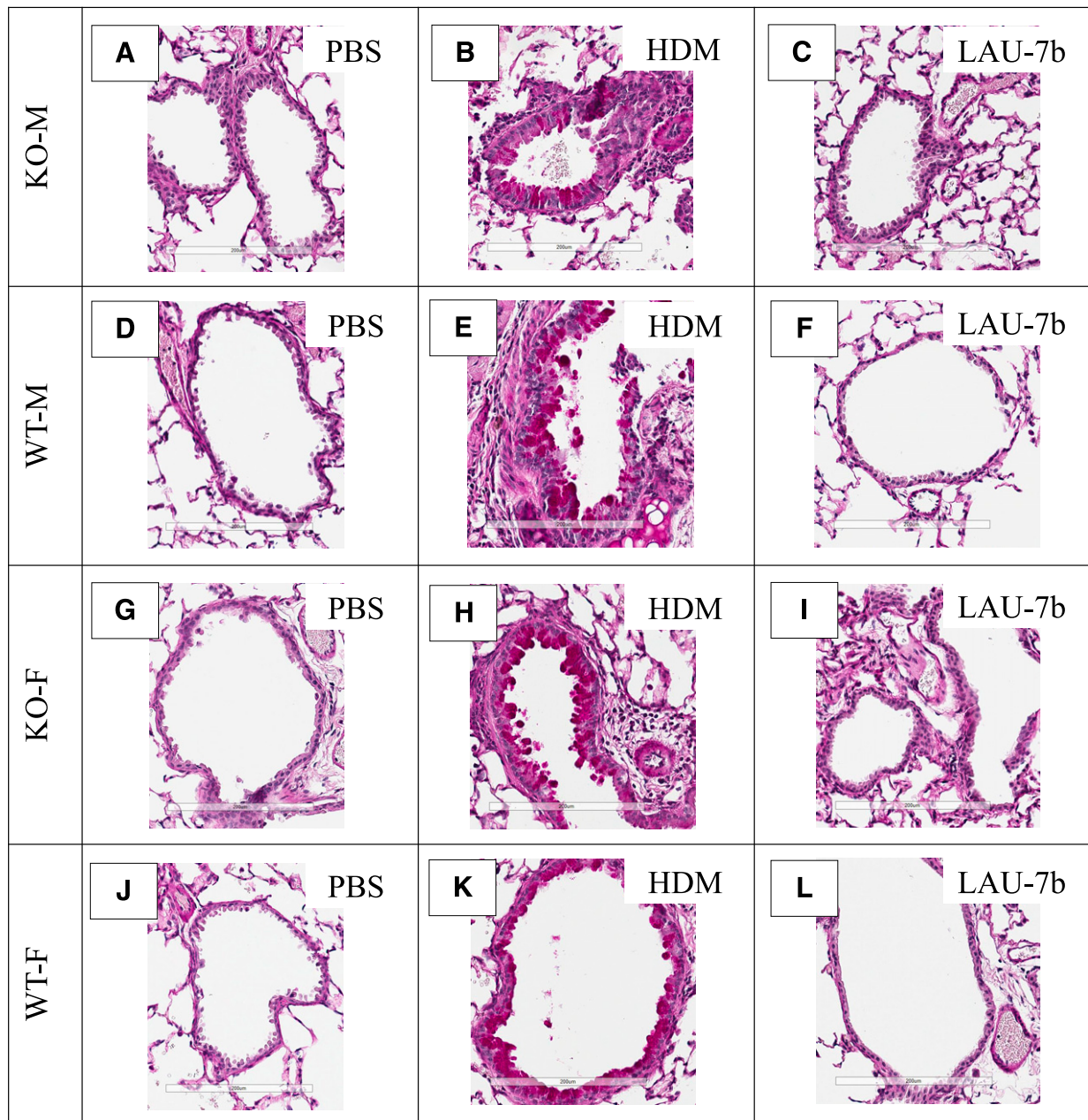


**Fig. 2.** Classic invasive measurements of lung resistance and measurements of IgE levels of *Zbp2* KO and WT mice. Classic invasive measurements of lung resistance in (A) WT and (B) *Zbp2* KO mice. All mice were sensitized with HDM and challenged with either PBS or HDM. Treatment group is marked as LAU-7b. The doses of MCh used were 25, 50, and 100 mg/ml. LAU-7b treatment and airway hyper-responsiveness in MCh nebulized male (C) and female (D) mice. Two-way ANOVA,  $n$  equals at least 11 mice in each group. (E) Treatment with LAU-7b did not affect the levels of IgE in *Zbp2* KO and WT mice. Two-way ANOVA,  $n$  equals at least five mice for each group. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\*\* $P < 0.0001$ . F, female; M, male. RI: inspiratory airway resistance.



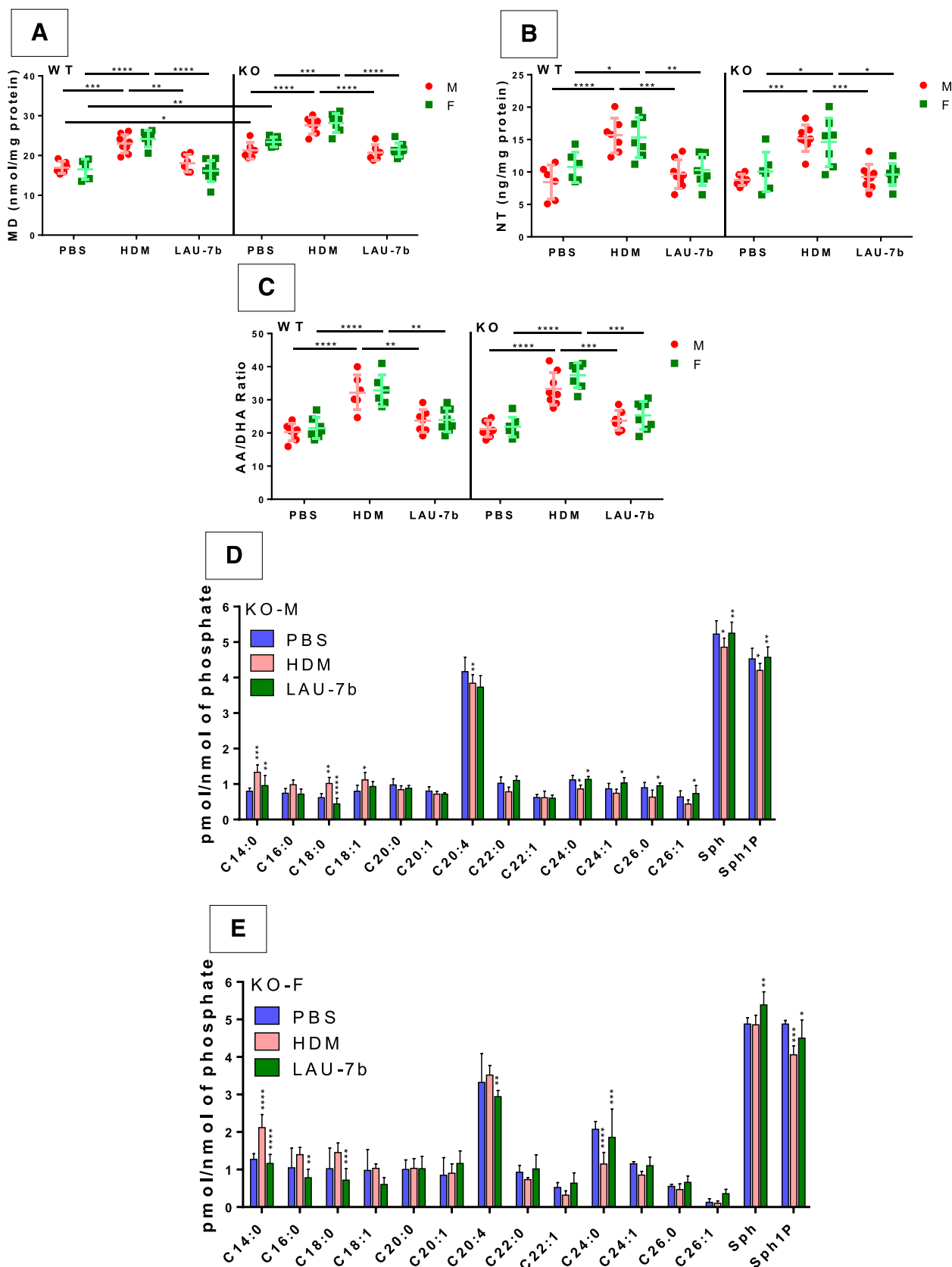


**Fig. 3.** Infiltration of airways by inflammatory cells in untreated and LAU-7b-treated mice; H&E staining. All mice were sensitized with HDM and challenged with either PBS or HDM. Treatment group is marked as LAU-7b. (A–C) *Zpbp2* KO males, (D–F) WT males, (G–I) *Zpbp2* KO females, and (J–L) WT females. (M) LAU-7b-treated mice have significantly lower lung cell infiltration compared with the lungs of untreated mice. Quantification was done by counting the number of inflammatory cells around each airway and normalizing it by division over the square of the perimeter “in millimeters” of the airway basement membrane. Measurements were done using at least four different airways for each mouse from each group. *n* equals at least seven mice for each group, two-way ANOVA. \*\**P* < 0.01; \*\*\**P* < 0.001; \*\*\*\**P* < 0.0001. F, female; M, male. Scale bar = 300  $\mu$ m.



**Fig. 4.** Mucus production by goblet cells in untreated and LAU-7b-treated mice. PAS staining. All mice were sensitized with HDM and challenged with either PBS or HDM. Treatment group is marked as LAU-7b. (A–C) *Zbp2* KO males, (D–F) WT males, (G–I) *Zbp2* KO females, and (J–L) WT females. (M) LAU-7b-treated mice have significantly lower production of mucus by goblet cells compared with untreated mice. Quantification was done by counting the number of PAS-positive cells around each airway and normalizing it by division over the perimeter “in millimeters” of the airway basement membrane. Measurements were done using at least four different airways for each mouse from each group. *n* equals at least seven mice for each group, two-way ANOVA. \*\*\**P* < 0.001; \*\*\*\**P* < 0.0001. F, female; M, male. Scale bar = 200  $\mu$ m.





**Fig. 5.** LAU-7b treatment corrected the aberrant levels of markers of oxidation, fatty acids, and sphingolipids in *Zpbp2* KO and WT mice. Analysis of (A) MD, marker of lipid oxidation; (B) NT, marker of protein oxidation; and (C) AA/DHA ratio. Analysis of sphingolipid species in (D) *Zpbp2* KO male and (E) *Zpbp2* KO female mice. All mice were sensitized with HDM and challenged with either PBS or HDM. Treatment group is marked as LAU-7b. Two-way ANOVA,  $n$  equals at least six mice for each group. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ ; \*\*\*\* $P < 0.0001$ . F, female; M, male. Sph: sphingosine and Sph1P: sphingosine-1-phosphate species.

WT mice. Interestingly, the ablation of the *Zbp2* gene, although it affected the lung resistance, did not result in the modulation of inflammatory response after allergen challenge. Although *Zbp2* KO mice have reduced AHR after HDM challenge, the inflammatory response demonstrated by the recruitment of cells into the airways and excessive mucus production was still induced. Altogether, in WT and *Zbp2* KO mice (males and females), LAU-7b treatment had resulted in significantly lower cell infiltration, no hyperplasia, and inhibited mucus production compared with PBS-treated mice of the same genotype and sex.

To determine whether *Zbp2* deficiency influenced the levels of IgE, a common marker of allergy, we quantified IgE in plasma, which was statistically elevated after HDM challenge in KO males and females to similar levels present in WT mice (Fig. 2E). Although LAU-7b treatment did not correct elevated plasma IgE levels either in KO males or in females [findings were similar to our previously published data (Kanagaratham et al., 2014, 2018)], it was still able to control the inflammatory response to the allergen in the lungs and normalize lung physiology of the allergen-sensitized and -challenged animals. These findings suggest that, even in the most severely affected patients who are not responding to either steroids or to anti-IgE therapy anymore, this type of treatment might still be effective.

*Ormdl3* plays a significant role in allergic asthma (Paulenda and Draber, 2016). In yeast, ORM proteins negatively regulate sphingolipid synthesis by forming a conserved complex with the serine palmitoyl transferase enzyme, thus inhibiting the first rate-limiting step of *de novo* ceramide, and all other sphingolipids', biosynthesis (Breslow et al., 2010). In a mouse study, myriocin, an inhibitor of the serine palmitoyl transferase enzyme, reduced the *de novo* sphingolipid synthesis and increased bronchial reactivity (Worgall et al., 2013). Likewise, increased airway responsiveness and airway remodeling has been reported in C57BL/6 transgenic mice overexpressing *Ormdl3* after ovalbumin sensitization and challenge (Miller et al., 2014). Similarly, C57BL/6 mice lacking *Ormdl3* were protected from developing AHR and airway eosinophilia induced by *Alternaria alternata* (Löser et al., 2017).

Further, the expression of *ORMDL3* is not independent of the expression of other genes present in the 17q21 chromosomal locus. It has been reported that *ZBP2* and *ORMDL3* genes are coregulated together as closely associated *cis*-haplotype elements (Verlaan et al., 2009). In our study, we report that there is significantly lower *Ormdl3* gene expression in A/J *Zbp2* KO mice than in the littermate WT controls (Fig. 1D). The finding that knocking out the *Zbp2* gene in A/J mice has markedly reduced AHR can be, at least partially, attributed to the downregulation of two hyper-responsiveness susceptibility genes (*Ormdl3* and *Zbp2*) since this strain of mice expresses genes that make them both atopic (loci on chromosome 4) and display increased AHR (loci on chromosome 12).

It has been reported that elevated levels of *ORMDL3* inhibit sphingolipid biosynthesis, resulting in inhibition of both LCCs (C16:0) and four species of VLCCs (C22:0, C24:0, C24:1, and C24:2) (Kiefer et al., 2019). By contrast, Zhang et al. (2019) had reported that *ORMDL3* gene silencing in A549 and normal human bronchial epithelial (NHBE) cell lines resulted in a marked increase in the levels of C24:0, C24:1, C26:1, and

sphingosine-1-phosphate species. Recently, Debeuf et al. (2019) reported that transgenic mice overexpressing *Ormdl3* had significantly reduced levels of C24:0 and C24:1. By contrast, loss of *Ormdl3* in mice resulted in elevated levels of C24:0 and C24:1. Interestingly, our results demonstrated that LAU-7b treatment had lowered the expression of *Ormdl3* (Fig. 1D) and protected the sensitized and challenged WT mice against allergic asthma. It is well established that *Ormdl3* overexpression inhibits *de novo* sphingolipid biosynthesis; therefore, the LAU-7b-induced downregulation of *Ormdl3* gene expression in WT mice may explain the reason why we observed an increase in the levels of VLCCs in LAU-7b-treated mice (Fig. 5). Although *Zbp2* KO mice have lower AHR after HDM sensitization and challenge, the inflammatory reaction after allergen exposure still occurred to its full capacity. It is intriguing that LAU-7b treatment was able to protect the *Zbp2* KO mice against the inflammatory reactions after HDM sensitization and challenge. This effect cannot be explained by the ability of LAU-7b to modulate *Ormdl3* expression, since our *Zbp2* KO mouse model has ablated expression of *Zbp2* and reduced expression of *Ormdl3*.

We tested four genes (*Il-4*, *Il-5*, *Il-13*, and *Eotaxin-1*) that are overexpressed by T helper type 2 cells in allergic asthma (Fig. 1, E–H). After challenge with HDM, both *Zbp2* KO and WT mice had significantly increased expression of *Il-5*, *Il-13*, and *Eotaxin-1* genes compared with PBS-challenged control mice (Fig. 1, E–H). However, for these three genes, no significant differences were observed between *Zbp2* KO and WT mice. These findings coincide with nonsignificant differences between the two mouse strains in terms of elevated IgE levels (Fig. 2E) and an increased number of inflammatory cells recruited into the airways (Fig. 3) after allergen challenge. In our *Zbp2* KO mouse model, the expression of *Il-4* was not elevated significantly after HDM challenge compared with PBS KO controls, unlike HDM-challenged WT mice, which show significant elevation of *Il-4* levels. Miller et al. (2018) reported a lack of significant increase in *IL-13* levels in *Zbp2* KO mice after HDM challenge compared with *Zbp2* KO controls. The results of the studies reported by Debeuf et al. (2019) also demonstrated the lack of a significant increase in the levels of *IL-13* or *IL-5* after HDM challenge when the mice were either overexpressing, or had displayed the loss of, *Ormdl3*. Levels of *IL-4* were not reported in these two studies cited above. Taken together, there is no evidence linking the alteration of *Zbp2* or *Ormdl3* genes with modulation of *Il-4*, *Il-5*, *Il-13*, or *Eotaxin-1* expression.

Our studies clearly demonstrate that treatment of allergen-sensitized and allergen-challenged mice with LAU-7b had lower expression of *Il-4* (WT mice), *Il-5*, and *Eotaxin-1* (WT and KO mice) compared with PBS-treated mice, which is consistent with the previously reported effect on extracellular signal-regulated protein kinases 1 and 2 (ERK1/2) phosphorylation (Lachance et al., 2013). *IL-13* is a key regulatory cytokine in the production of IgE (Rael and Lockey, 2011). Interestingly, both *Zbp2* KO and WT mice treated with LAU-7b did not show significant reduction of *Il-13* expression, which may explain why IgE levels remained elevated in both groups. Interestingly, inhibition of IgE levels was not essential for improving lung physiology in treated animals.

Overall, the preclinical data presented here for Fenretinide, the active pharmaceutical ingredient of LAU-7b capsules,

provide robust grounds that justify testing the efficacy of this dosage form in a clinical trial, especially among asthmatics whose severe asthma is no longer treatable using the currently available therapies.

## Conclusions

LAU-7b treatment in a dose of 10 mg/kg per day for 9 days protects both *Zbp2* KO mice, which have significantly reduced *Ormdl3* expression, and their WT littermate controls from HDM-induced allergic asthma. Interestingly, in WT mice, LAU-7b had significantly lowered the expression of *Ormdl3*, and hence, it can be effective in allergic asthma treatment by elevating the levels of VLCCs and decreasing the levels of LCCs.

## Authorship Contributions

*Participated in research design:* Youssef, Naumova, Radzioch.

*Conducted experiments (animal experiments, whole-body plethysmography, airway resistance measurements, histology work and IgE measurement):* Youssef. *(RNA expression analysis):* Youssef, Shah, Dumut. *(Measurements of lipids, ceramides and markers of oxidative stress):* De Sanctis, Hajdudch.

*Performed data analysis:* Youssef.

*Wrote or contributed to the writing of the manuscript:* Youssef, De Sanctis, Shah, Dumut, Hajdudch, Naumova, Radzioch.

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