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## Retinoids enhance the expression of cathelicidin antimicrobial peptide during reactive dermal adipogenesis

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### Abstract

A subset of dermal fibroblasts undergo rapid differentiation into adipocytes in response to infection and acutely produce the cathelicidin antimicrobial peptide gene *Camp*. Vitamin A and other retinoids inhibit adipogenesis, yet can show benefit to skin disorders such as cystic acne that are exacerbated by bacteria. We observed that retinoids potently increase and sustain the expression of *Camp* in preadipocytes undergoing adipogenesis despite inhibition of markers of adipogenesis such as *Adipoq*, *Fabp4* and *Rstn*. Retinoids increase cathelicidin in both mouse and human preadipocytes, but this enhancement of antimicrobial peptide expression did not occur in keratinocytes or a sebocyte cell line. Preadipocytes undergoing adipogenesis more effectively inhibited growth of *S. aureus* when exposed to retinoic acid. Whole transcriptome analysis identified hypoxia-inducible factor 1 $\alpha$  (HIF-1 $\alpha$ ) as a mechanism through which retinoids mediate this response. These observations uncouple the lipid accumulation element of adipogenesis from the innate immune response and uncover a previously unsuspected mechanism that may explain therapeutic benefits of retinoids in some skin disorders.

### Introduction

Retinoids comprise a large group of chemical compounds that are vitamin A (retinol) analogues and/or derivatives, and can influence multiple physiological processes including vision, cell proliferation, differentiation, and immunity (1, 2). Within the skin, retinoids have been shown to affect multiple cell types, modulating several processes including wound healing, pigmentation, and immune defense (3–9) and have been used pharmacologically to treat chronological and photoaging, psoriasis, and acne (9–13). There are few cases in which retinoids have exhibited direct antimicrobial activity (14, 15), however under most

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circumstances immune modulation by retinoids is thought to only occur indirectly (5–7, 9, 16–18). For example, the mechanism by which retinoids are therapeutically useful in acne, a disorder that is driven in part by the bacterium *Cutibacterium acnes*, is thought to occur through their effect to decrease sebum production and alter epidermal function rather than through an influence on immune defense against bacterial growth (9).

It has been recently observed that dermal fibroblasts of the preadipocyte lineage will undergo rapid, local proliferation and differentiation in response to bacterial infection of the skin (19, 20). This process, dubbed reactive adipogenesis, results in an increase in the expression of the cathelicidin gene *CAMP*, an antimicrobial peptide (AMP) that peaks during this process as the local adipocytes mature (20). Reactive adipogenesis is essential for optimal innate immune defense since impairment of the adipogenic process decreases total skin cathelicidin expression and increases the severity of bacterial infections (19–21).

Retinoids have been shown to inhibit adipogenesis and promote lipid breakdown (22–27), and therefore would be predicted to decrease cutaneous antimicrobial defense capacity. In this study, we sought to understand the effects of retinoids on AMP production during adipogenesis in order to better understand the therapeutic mechanism of action of retinoids. Despite exhibiting the anticipated anti-adipogenic effects, we unexpectedly observed that retinoids enhanced and sustained cathelicidin levels in developing preadipocytes, ultimately translating to increased antimicrobial activity. This process was found to depend in part on hypoxia-inducible factor 1- $\alpha$  (HIF1 $\alpha$ ), a broadly active transcription factor within the *CAMP* gene promoter. These observations reveal an uncoupling between adipogenesis and antimicrobial expression that could be exploited as a novel means of modulating innate immune defense.

## Materials and Methods

### Antibodies, chemicals, and reagents

Vitamin A (retinol), all trans-retinal, all trans-retinoic acid, 9 cis-retinoic acid, 13 -cis-retinoic acid, 1 $\alpha$ ,25-Dihydroxyvitamin D3, L-Mimosine, tazarotene, IBMX (3-Isobutyl-1-methylxanthine), dexamethasone, indomethacin, and recombinant human insulin were purchased from Sigma-Aldrich (St. Louis, MO). Acryflavine hydrochloride was purchased from Thermo-Fisher Scientific (Pittsburgh, PA). AM580 was purchased from Abcam. MALP-2 was purchased from Enzo Life Sciences. Rabbit anti-CRAMP antibodies were made in our lab as previously described (28). Mouse anti-FABP4 was purchased from Santa Cruz Biotechnology (Santa Cruz, CA).

### Cell Culture

3T3-L1 primary mouse preadipocytes were purchased from ATCC (CL-173). Neonatal primary dermal fibroblasts were isolated by our lab as previously described (19). Cells were grown in DMEM supplemented with 10% FBS, glutamax, and antibiotic-antimycotic (Thermo Fisher Scientific) in a humidified incubator at 5% CO<sub>2</sub> and 37°C under sterile conditions. To induce differentiation, two-day post-confluent 3T3-L1 cells were switched to adipocyte differentiation medium containing 2  $\mu$ M dexamethasone, 250  $\mu$ M IBMX, 200

$\mu\text{M}$  indomethacin, and  $10\ \mu\text{g}/\text{mL}$  recombinant human insulin. For timecourse experiments, differentiation media was refreshed at indicated times. Immortalized non-diabetic human preadipocytes (29) were a generous gift from Dr. Patrick M. Schliever (Carver College of Medicine, Iowa City, Iowa). Human preadipocytes were grown and differentiated as previously described (29) and differentiation media was refreshed on indicated days. Human epidermal keratinocytes (Life Technologies, #C-001–5C) were cultured in EpiLife medium supplemented with  $60\ \mu\text{M}$   $\text{CaCl}_2$ , EpiLife Defined Growth Supplement (Life Technologies), and antibiotic/antimycotic. SEB-1 sebocytes were cultured in Sebomed basal medium (Millipore) supplemented with recombinant human epidermal growth factor ( $5\ \text{ng}/\text{ml}$ ) (Sigma). To examine the effects of retinoids on cathelicidin expression, retinoids ( $1\ \mu\text{M}$  unless otherwise noted) or vehicle (ethanol) were added to differentiation media or cell culture media on day 0 (unless indicated otherwise). Retinoid-containing media was subsequently refreshed every 2–3 days throughout the course of the experiment. Treatment of cells with  $1\alpha,25\text{-Dihydroxyvitamin D}_3$  (Vitamin D) ( $10\ \text{nM}$ ) or vehicle (ethanol) was conducted identical to that of retinoids. To examine the role of HIF-1 $\alpha$ , cells were cotreated with L-mimosine ( $400\ \mu\text{M}$ ) or acryflavine ( $5\ \mu\text{M}$ ) or vehicle (water) for 24 hours.

### RNA isolation, cDNA synthesis, and RT-qPCR analysis

Cultured cells were lysed with Trizol reagent (Life Technologies) or PureLink Lysis Buffer (Ambion/Life Technologies) and RNA was isolated using the PureLink isolation kit. Up to  $1\ \mu\text{g}$  of RNA was reverse transcribed to cDNA using the iScript cDNA synthesis kit (Bio-Rad). Quantitative real-time PCR were performed the CFX96 Real-Time System (Bio-Rad) using SYBR Green Mix (Biomake, Houston, TX). The housekeeping gene, *Tbp* (TAT-binding box protein), was used to normalize gene expression in samples. Specific primer sequences are shown in supplemental table 1.

### Western Blots

Supernatants from treated cells were enriched for protein using Oasis HLB Plus Light Cartridges (Waters, Milford, MA) and lyophilized. Lyophilized protein from  $500\ \mu\text{L}$  of supernatant was separated on a Novex 10–20% Tricine Gel (Invitrogen) and transferred to a PVDF membrane using Trans-Blot Turbo Transfer Pack (Bio-Rad) according to the manufacturer's instructions. Membranes were blocked with Odyssey blocking buffer (LI-COR) and incubated overnight at  $4\ ^\circ\text{C}$  with primary antibodies. Immunoblots were washed and incubated with fluorescent secondary antibodies (LI-COR) for 1 hour at room temperature and subsequently imaged using the Odyssey Infrared Imager (LI-COR).

### Immunofluorescence

For immunofluorescence analysis, 3T3-L1 cells were grown, differentiated, and treated in 8-well chamber slides as described earlier. Cells were fixed in 4% PFA for 10 minutes and subsequently incubated with primary antibodies overnight at  $4\ ^\circ\text{C}$ . Cell were washed and incubated for 1 hour with secondary antibodies conjugated to Alexa Fluor 594 (Invitrogen) and nuclei were counterstained with DAPI. All images were taken with an Olympus BX41 microscope.

## Histology and immunohistochemistry

Tissue biopsies were directly embedded in OCT compound. Frozen sections were fixed in 4% PFA for 10 minutes prior to Hematoxylin/Eosin (H&E) staining or immunofluorescence staining. For immunohistochemistry, fixed and permeabilized frozen tissue sections were stained with Bodipy solution to visualize lipid droplets and nuclei were counterstained with DAPI. All images were taken with an Olympus BX41 microscope.

## *In Vitro* bacterial killing assay

Antibiotic free conditioned media was collected from differentiating dermal fibroblasts as previously described (20). Conditioned media (200  $\mu$ L) was mixed with  $10^5$  colony forming units (CFU) of *S. aureus* stain USA300 in deep-well 96-well plates and incubated at 37° with shaking at 275 R.P.M. for 12 hours prior to plating for CFU counting.

## Mice Experiments

All animal experiments were approved by the University of California, San Diego (UCSD), Institutional Animal Care and Use Committee. SKH-1 hairless mice were originally purchased from Charles River Laboratories and bred and maintained in animal facility of UCSD. Animals in all experimental models were age- and sex-matched. Skin infections were conducted as previously described (20) using *S. aureus* strain USA300 (MRSA) (*S. aureus* strain AH4807 was used for IVIS experiments) or PBS control. Mice were distally administered one subcutaneous injection with retinoic acid (10 mg/kg) or vehicle (DMSO) dissolved in olive oil on the day of infection and two additional injections on subsequent days. Mice were sacrificed on day 3 and a skin biopsy of the infected region was harvested. Skin biopsies were homogenized in PureLink Lysis Buffer (or PBS for CFU counting) with 2 mM zirconia beads in a mini-bead beater (Biospect, Bartlesville, OK) for RNA extraction. To quantify CFU, homogenized skin samples were serially diluted and plated onto Baird-Parker agar and grown for 24 hrs. to quantify the CFU per gram of tissue. For *in vivo* live bacterial imaging, mice were imaged under isoflurane inhalation anesthesia (2%). Photons emitted from luminescent bacteria were collected during a one-minute exposure using the Xenogen IVIS Imaging System and living image software (Xenogen, Alameda, CA). Bioluminescent image data are presented on a pseudocolor scale overlaid onto a gray-scale photographic image. Using the image analysis tools in living image software, circular analysis windows (of uniform area) were overlaid onto regions of interest and the corresponding bioluminescence values (total flux) were measured.

## RNA Sequencing

Purified RNA was submitted to the University of California, San Diego (UCSD) Institute for Genomic Medicine core facility for library preparation and high-throughput next-generation sequencing. Library preparation and subsequent analysis were conducted as previously described (30). Ven diagram was prepared using Venny (31).

## Statistical analysis

Experiments conducted were performed at minimal performed in triplicate with at least three technical replicates. Statistical significance was calculated using a Student's two-tailed t-test where \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , \*\*\*\* $P < 0.0001$  as indicated in figure legends.

## Results

### Retinoids increase cathelicidin expression during adipogenesis

To determine the effects of retinoids on *Camp* expression during reactive adipogenesis, we first examined differentiation of preadipocytes *in vitro* under defined conditions. Differentiation was associated with increased *Camp* mRNA expression, and treatment with the TLR2/6 ligand, MALP-2, further stimulated *Camp* expression in both undifferentiated and differentiating 3T3-L1 preadipocytes (Figure 1A). The addition of vitamin A (retinol) greatly amplified the *Camp* response to differentiation, resulting in an increase in steady-state *Camp* mRNA levels compared to untreated 3T3-L1 cells with the greatest increase occurring with Malp-2 and retinol (Figure 1A).

Vitamin A, (AKA, retinol (ROL) is an inactive precursor that can be enzymatically converted to retinoic acid (RA) or retinal (RAL) to exhibit physiological activity. Addition of ROL, RAL or RA had similar effects on *Camp* induction when added to 3T3-L1 cultures at the start of an 8-day differentiation experiment. RA appeared most potent when added between day 2–8 after differentiation was initiated, and induction was least when added later during adipogenesis at day 5–8 (Figure 1B). These findings suggest adipocyte precursors can metabolize Vitamin A to active forms.

RA can isomerize between all-*trans* (ATRA), 9-*cis* retinoic acid (9CRA), and 13-*cis* retinoic acid (13CA) (32). Similar activity was observed by each of these isoforms on *Camp* expression (Figure 1C). Using RA analogues specific for the retinoid acid receptors (RAR), agonists for both RAR $\alpha$  and RAR $\beta/\gamma$  potently stimulated *Camp* expression to an extent similar to RA (Supplemental Figure 1A). RA demonstrated a dose-dependent induction of *Camp* at 24 hours (Figure 1D), but significantly induced *Camp* expression in as little as 6 hours and resulted in sustained high levels of expression at day 8, a prolonged increase which contrasted to the normal downregulation of *Camp* seen during late adipogenesis in the absence of a retinoid (Figure 1E). Previous observations have shown that *Camp* is the only antimicrobial peptide induced during adipogenesis whereas expression of  $\alpha$ - and  $\beta$ -defensins was not altered (20). Our current studies are consistent with these observations as addition of RA was unable to induce expression of other AMPs (Supplemental Figure 1B). Contrary to the increase in AMP gene expression that was promoted by retinoids, but consistent with prior observations of the effect of RA to inhibit adipogenesis (27, 33), we observed that RA inhibited expression of adipocyte maturation markers *Adipoq*, *Fab4* and *Rstn* during the time course of this experiment (Figure 1F–G). These observations suggest that RA promoted an uncoupling between AMP expression and classic markers of adipocyte differentiation that are associated with lipid accumulation.

In agreement with our transcriptional studies, significantly elevated levels of the murine cathelicidin protein (Cramp) precursor protein were detected in RA treated 3T3-L1 cell

supernatants (Figure 2A). Cramp immunostaining complimented these findings as more Cramp staining was seen in the cytoplasm of 3T3-L1 cells during differentiation into adipocytes in the presence of RA (Figure 2B). On day 12, numerous round areas in the cytosol that were unstained by DAPI or for Cramp are maturing lipid droplets.

Human and mouse cathelicidin genes exhibit several regulatory parallels, but also have important differences. The human *CAMP* promoter possesses an active vitamin D response element (VDR) that is lacking in mice (34, 35), and vitamin D can upregulate *CAMP* expression in several human cell types including human keratinocytes (34, 36). To determine if the observations made in murine-derived 3T3-L1 cells could be found in human preadipocytes, we investigated the effects of RA and vitamin D in a cultured human preadipocyte line (19, 20). Both 9-*cis* retinoic acid and vitamin D increased *CAMP* expression in human preadipocytes and exhibited an additive effect when cells were treated with both together (Figure 2C). However, RA was unable to stimulate *CAMP* expression under similar culture conditions in a human sebocyte cell line or in primary cultures of normal human keratinocytes (Figure 2D). Thus, we conclude that retinoids can increase expression of cathelicidin in both human and mouse preadipocytes but the effects of retinoic acid on this response are specific to cell type.

Given the significant cathelicidin upregulation observed with retinoid treatment, we next wished to determine whether these results correlate with antimicrobial activity. As our observations have shown that mouse neonatal dermal fibroblasts (MSFBs) express significantly more Cramp protein compared to 3T3-L1 cells (Supplemental Figure 2A) we selected to use MSFBs to determine antimicrobial activity. MSFB cells were stimulated to initiate adipogenesis in culture with in the presence of RA, and this culture supernatant then added to bacterial cultures of *S. aureus* (USA300). The growth of *S. aureus* was maximally inhibited from culture supernatants of cells treated with RA (Figure 3A). Despite promoting *Camp* expression and enhancing antimicrobial activity, RA's effects on cytokine and chemokine expression were comparatively minor among under the conditions we examined (Supplemental Figure 2B).

Given our *in vitro* findings, we next wished to determine how RA might contribute to antimicrobial activity during *S. aureus* infection. Systemic treatment of mice with 10 mg/kg RA for 3 days decreased dermal adipose in uninfected skin as well as in skin after challenge with *S. aureus* (Figure 3B,C). *Camp* mRNA expression was induced by *S. aureus* and RA (Figure 3D). However, no consistent change in the susceptibility to *S. aureus* survival was seen in mice treated with RA by manual CFU counting of live bacteria (Figure 3E) or as detected by IVIS analysis of a reporter strain of *S. aureus* (Figure 3F).

### **Transcriptomic response of preadipocytes to RA identifies HIF1 $\alpha$ as mediator of antimicrobial response.**

To gain further insight into the mechanism through which preadipocytes increase *Camp* in response to RA, we performed RNA-sequencing of 3T3-L1 cells. The expression of genes that were changed by more than 2-fold when 3T3-L1 cells were differentiated in the presence of retinoic acid were identified (Figure 4A–B). An increase in genes associated with adipogenesis (*Zfp423*, *Fabp4*, *Adipoq*, *Cebpa*) was evident with differentiation, but



several genes involved in hypoxic responses were also observed to be upregulated by RA (Supplementary Figure 3A), particularly HIF1 $\alpha$  (Figure 4C). Quantitative PCR validated that *Hif1a* expression was increased in both mouse and human preadipocytes treated with RA (Figures 4F and Supplemental Figure 3B). *Hif1a* mRNA expression was also increased in skin samples from mice treated with RA and infected with *S. aureus* (Figure 4D).

To ascertain the role of HIF1 $\alpha$  in the response of preadipocytes to RA we used an established HIF1 $\alpha$  agonist (L-mimosine) and inhibitor (acryflavine). *Camp* expression was induced in preadipocytes treated with the HIF1 $\alpha$  agonist and further enhanced by RA, and the HIF1 $\alpha$  antagonist blocked *Camp* expression (Figure 4E,F). Vascular endothelial growth factor A (VEGFA), a regulatory target of HIF1 $\alpha$ , responded as expected with an increase in expression after agonist exposure and a decrease after antagonist treatment, thus serving as a control for the response to these chemical modifiers of HIF1 $\alpha$  activity (Supplemental Figure 3C). Given our observations that MALP-2 can stimulate *Camp* expression both in undifferentiated and differentiating 3T3-L1 cells, we validated the role HIF1 $\alpha$  in *Camp* regulation by cotreating cells with both acryflavine and MALP-2 under both conditions. Similarly, HIF1 $\alpha$  inhibition effectively blocked MALP-2 induction of *Camp* expression and further evidenced the regulatory role of HIF1 $\alpha$  (Supplemental Figure 3D, E).

## Discussion

Retinoids are an effective therapy in the treatment of acne and other disorders (9–13). However, the clinical response to retinoids is dependent on mode of delivery and the mechanisms responsible for the physiological effects of retinoids are incompletely understood. Retinoids indeed have been shown to exhibit both anti-inflammatory and pro-inflammatory effects on cellular immune responses of the skin (9, 37–39), but the activity of retinoids on innate host antimicrobial defense have not been well characterized. In this study we present an unexpected observation that retinol and its metabolites potentially enhance cathelicidin antimicrobial peptide expression during reactive adipogenesis. These findings suggest an alternative and previously underappreciated mechanism for the action of retinoids in the skin.

Retinoids inhibit adipogenesis and promote lipid breakdown (22–27). Inhibition of adipogenesis by inhibition of the *Ppar $\gamma$*  system will also inhibit the antimicrobial activity of dermal preadipocytes (19, 20). However, and contrary to expectations, we show in the current study that retinoids increase *Camp* expression despite exhibiting anti-adipogenic activity. This observation demonstrates an uncoupling between the processes of adipogenesis and antimicrobial peptide expression.

Previous observations have shown that adipocytes produce the cathelicidin precursor protein and that is upregulated during adipogenesis that is processed into an active antimicrobial peptide (CRAMP), however the enzyme that catalyzes this reaction in adipocytes has not been identified (19, 20). Our studies have shown that an increase in *Camp* gene expression then results in a significant upregulation of the CRAMP preprotein, yet the increase in antimicrobial activity was modest in comparison to the change in protein abundance. Whether this was indicative of a saturation of the system or modulation of CRAMP



processing from inactive precursor protein to active peptide is unclear. We hypothesize that the contribution of other cells *in vivo* during the complex response of reactive adipogenesis may contribute to the maturation of the cathelicidin precursor that is enhanced by retinoids in preadipocytes.

Excess Vitamin A results in epidermal hyperplasia and a reduction in white adipose tissue (WAT), while infection results in an expansion of dermal white adipose tissue (DWAT) (20, 27, 40). Interestingly, RA treatment attenuated the DWAT expansion associated with *S. aureus* infection, yet *Camp* was still significantly upregulated in comparison to the individual stimuli. RA improved the capacity of differentiating preadipocytes to inhibit *S. aureus*. However, despite these observations that RA was active *in vivo* and induced antimicrobial activity *in vitro*, there was no clear increase in the antimicrobial activity *in vivo* of mouse skin against *S. aureus*. This suggests that the immunomodulatory role of RA in the skin is complex and may be dependent on dose, timing or target organism. Additional work is needed to define the physiological significance of retinoid exposure on innate immune defense.

To gain further insight and understand the complexity of RA-mediated regulation, we examined the transcriptomic response to RA in the context of differentiating preadipocytes. RNA-seq analysis identified HIF1 $\alpha$  as a potential regulator of *Camp* expression. The human *CAMP* promoter possesses a HIF1 $\alpha$ -response element (HRE) and has been implicated in cutaneous immune defense as well as CRAMP expression in the colon (41). The *Hif1a* promoter possesses a retinoic acid-response element (RARE) (42) and RA has been reported to enhance HIF1 $\alpha$  expression under normoxic conditions (42–44). These past observations are in agreement with our RNA-seq and qPCR data where we found that RA induced elevated *Hif1a*. Similarly, activation of HIF1 $\alpha$  by the iron-chelator L-mimosine mimicked the effects of RA on *Camp* expression. The combinatorial effect of both activation and increased *Hif1a* expression exhibited a significant additive effect, thus providing further evidence that HIF1 $\alpha$  and RA work in concert to promote *Camp* expression. Interestingly, HIF1 $\alpha$  inhibition abolished *Camp* expression under all conditions. This may suggest *Camp* transcription is entirely dependent on HIF1 $\alpha$  or that the global effects of HIF1 $\alpha$  further attenuate other mechanisms involved in the expression of *Camp*. Support for this comes from observations that HIF1 $\alpha$  inhibition also downregulates *Cebpb* transcription and inhibits initiation of adipogenesis (45). Inhibition by acryflavine similarly downregulated *Cebpb* expression while L-mimosine downregulated *Cebpa*, the latter of which has also been proposed to regulate *CAMP* expression. Taken together this could suggest that C/EBP $\beta$  indirectly regulates *Camp* expression, however our data do not define if either C/EBP $\beta$  or C/EBP $\alpha$  directly regulate this process.

Induction of adipogenesis is mediated by C/EBP $\beta$ , which in turn mediates the expression of the preadipogenic factors C/EBP $\alpha$  and PPAR $\gamma$  (46, 47). While RA does not inhibit the expression of C/EBP $\beta$ , it has been shown to inhibit its function (22). Our observations have shown that initiation of differentiation, yet not final maturation of adipocytes, is sufficient and necessary to enhance *Camp* expression by retinoids. However, there appears to be a limited timeframe during which this effect can be observed as the potency of retinoid stimulation was markedly reduced when treatment was initiated at later timepoints

during differentiation. This suggests that access to the *Camp* promoter is limited during adipogenesis, such a limited access period may partially explain why acute, high dose RA administration to mice did not improve resistance to *S. aureus* infection. Under these conditions initiation of adipogenesis is also inhibited and may counteract the beneficial host defense function of RA that we have observed *in vitro*. Analysis of low dose Vitamin A exposure, as well as correlation of responses in human subjects on RA therapy, is needed.

In summary, these observations have revealed a novel disassociation between adipogenesis and the process through which fibroblasts in the dermis that become adipocytes can produce cathelicidin. Reactive adipogenesis is a recently discovered and important physiological response to infection of skin and gut (19–21, 48). These findings provide further insight into how *Camp* is regulated in adipocytes, a mechanism with important therapeutic implications.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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**Key Points**

Retinoids enhance cathelicidin antimicrobial peptide expression during adipogenesis

The action of retinoids is dependent on hypoxia-inducible factor 1 $\alpha$

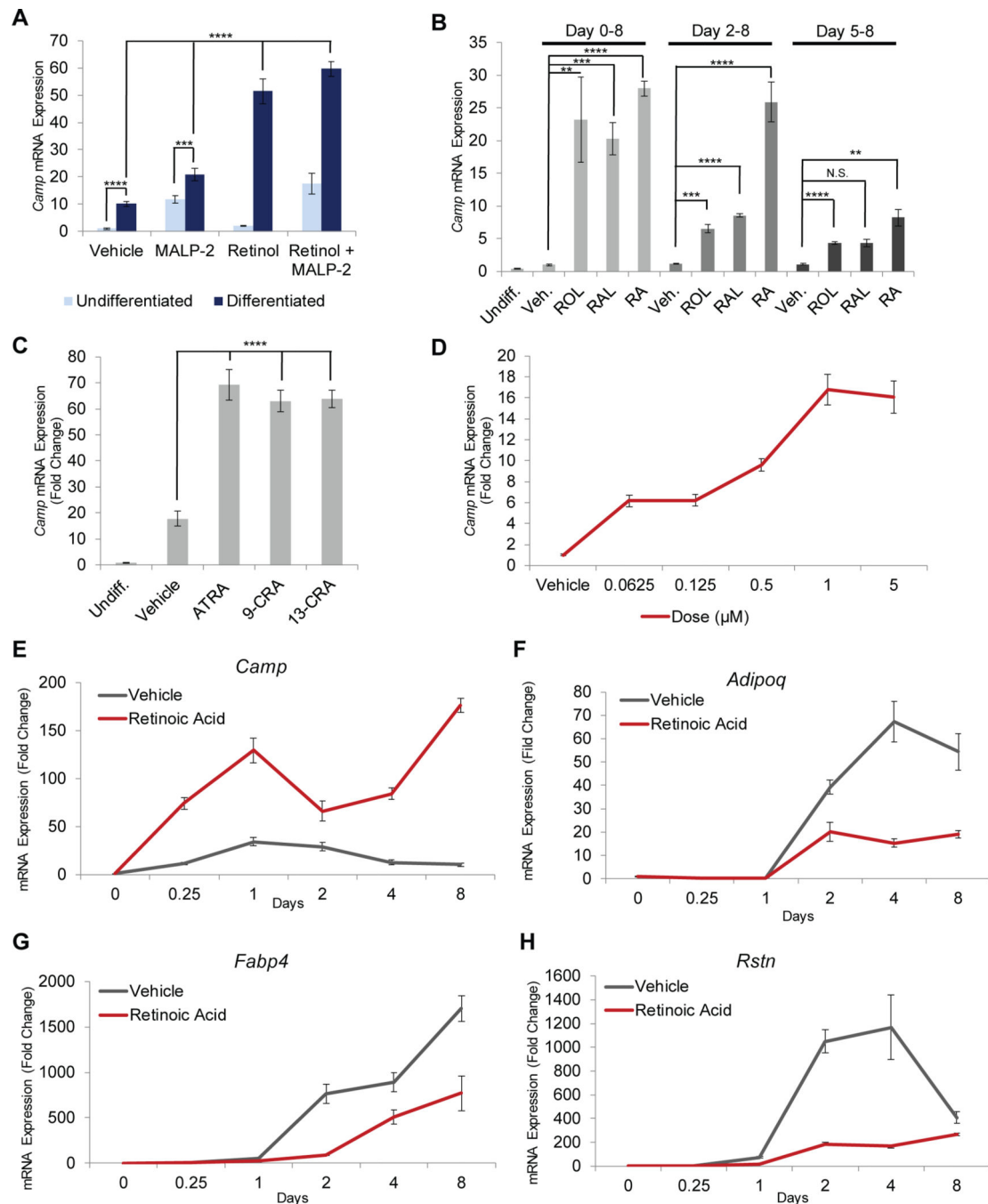
The capacity to induce cathelicidin may explain some therapeutic effects of retinoids

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**Figure 1. Retinoids significantly increase cathelicidin expression during adipogenesis.**

3T3-L1 preadipocytes were differentiated in the presence of retinoids or control, and relative mRNA expression of *Camp* (A-E), *Adipoq* (F), *Fabp4* (G), or *Rstn* (H) was assayed by RT-qPCR. (A) 3T3-L1 cells were treated for 24 h in the presence of 1  $\mu$ M retinol and/or 100 ng/ml MALP-2 or vehicle control in the absence or presence of differentiation media (DM). (B) Cells were differentiated for 8 days and co-treatment was initiated on day 0, day 2, or day 5 in the presence of 1  $\mu$ M retinol (ROL), all-*trans* retinal (RAL), retinoic acid (RA), or control. (C) Cells were differentiated for 24 h in the presence of ATRA, 9-*cis*

retinoic acid (9-CRA), or 13-*cis* retinoic acid (13-CRA). **(D)** Cells were differentiated for 24 h in the presence of increasing concentrations of RA. **(E-H)** Cells were differentiated for 8 days in the presence of RA or vehicle control and mRNA expression was assayed at given timepoints. All data are mean  $\pm$  S.D, \*P<0.05, \*\*P<0.01, \*\*\*P<0.001, \*\*\*\*P<0.0001 using a Student's paired T test.

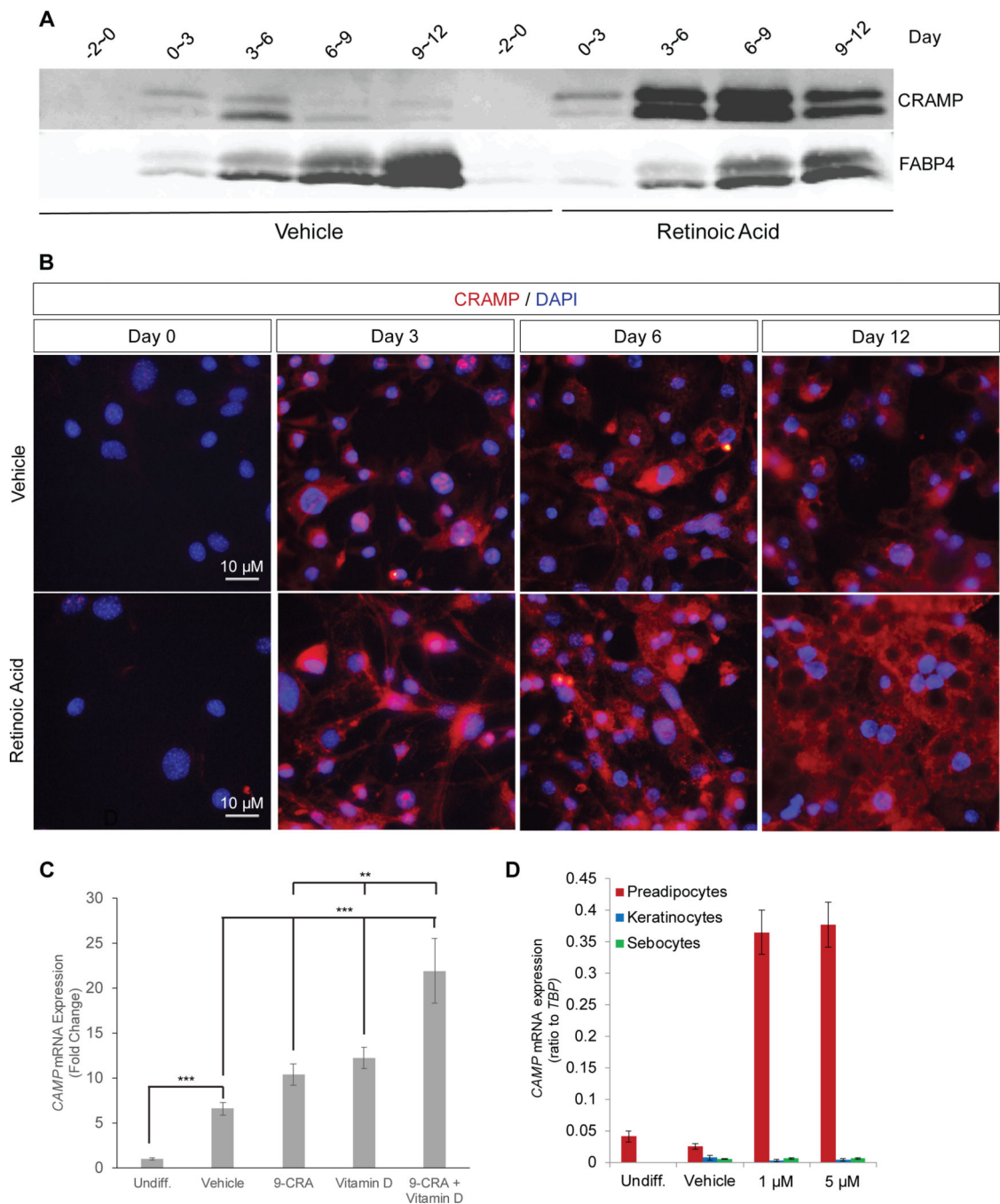
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**Figure 2. Retinoic acid enhances adipogenic innate immune response *in vitro*.**

(A) Western blot analysis of concentrated 3T3-L1 cell supernatants treated with 1  $\mu$ M retinoic acid (RA) or vehicle (ethanol). (B) Immunofluorescence analysis of CRAMP expression (red) in cultured 3T3-L1 cells treated with 1  $\mu$ M RA or vehicle. Nuclei are counterstained with DAPI. Scale bar 10  $\mu$ m. (C) Human preadipocytes were differentiated for 16 days in presence of 1  $\mu$ M 9-*cis* retinoic acid (9-CRA) and/or 10 nM 1 $\alpha$ ,25-Dihydroxyvitamin D<sub>3</sub> (Vitamin D) or vehicle control and C/EBP $\beta$  mRNA expression was assayed. (D) Differentiating 3T3-L1 preadipocytes, human keratinocytes, and human

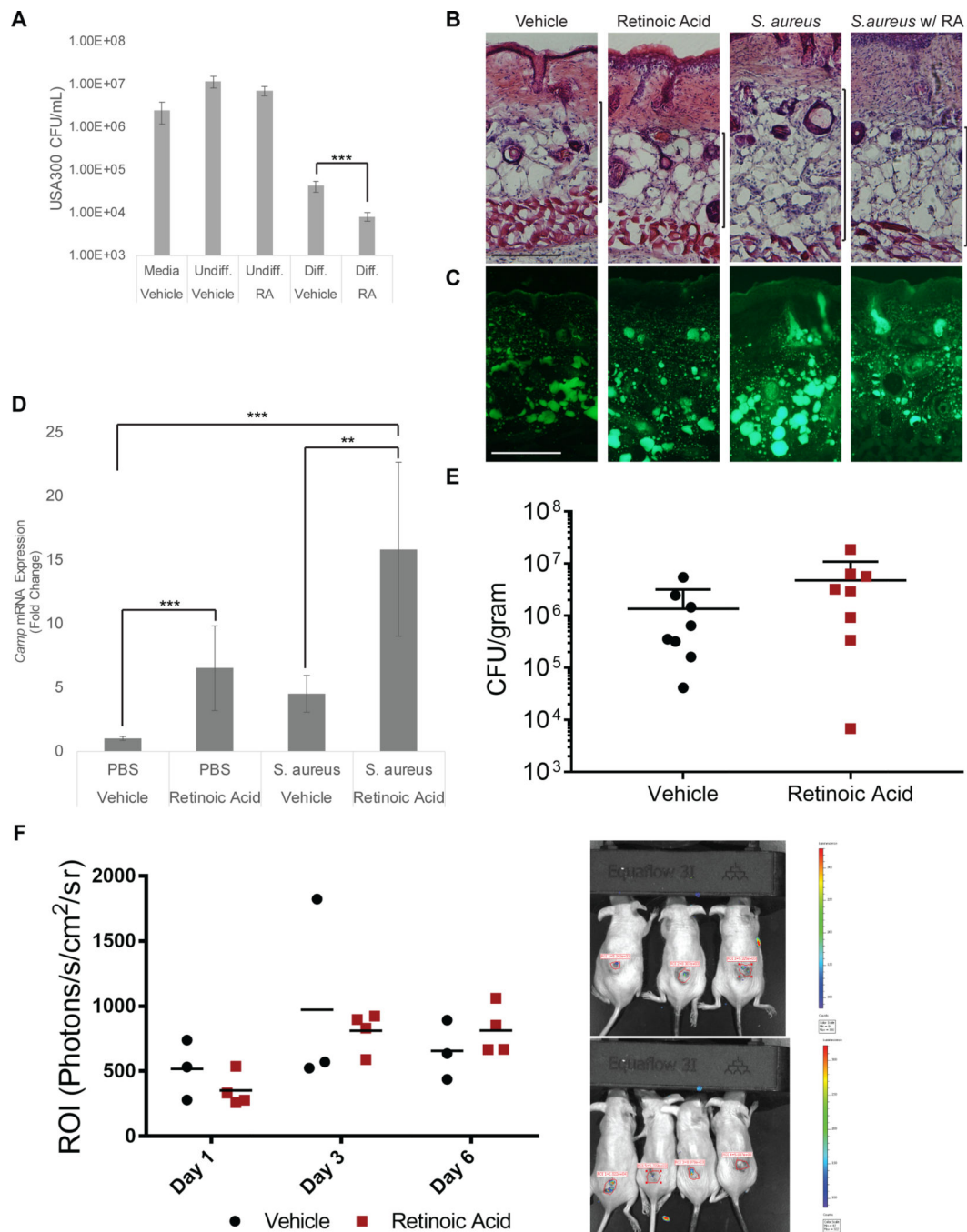
sebocytes were treated with 1 or 5  $\mu$ M RA for 5 days and *CAMP* mRNA expression was measured relative to *TBP*. All data are mean  $\pm$  S.D, \*P<0.05, \*\*P<0.01, \*\*\*P<0.001, \*\*\*\*P<0.0001 using a Student's paired T test.

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**Figure 3. Retinoic acid modulates reactive adipogenesis.**

(A) Supernatants from differentiated and undifferentiated dermal fibroblasts were grown in the presence of USA300 for 12 hrs. and CFU counts were determined. (B-D) SKH-1 wildtype mice (~8 weeks old) were distally intradermally infected with *S. aureus* on day 0 and simultaneously treated subcutaneously with RA or olive oil (control) for 3 days (n=4 per group). (B) Representative H&E histology of skin on day 3 of infection, scale bar 100  $\mu\text{m}$ . (C) Lipid droplet staining of frozen sections, scale bar 100  $\mu\text{m}$ . (D) Skin-lesions were collected and gene expression was analyzed by qPCR for *Camp* expression. (E-F)

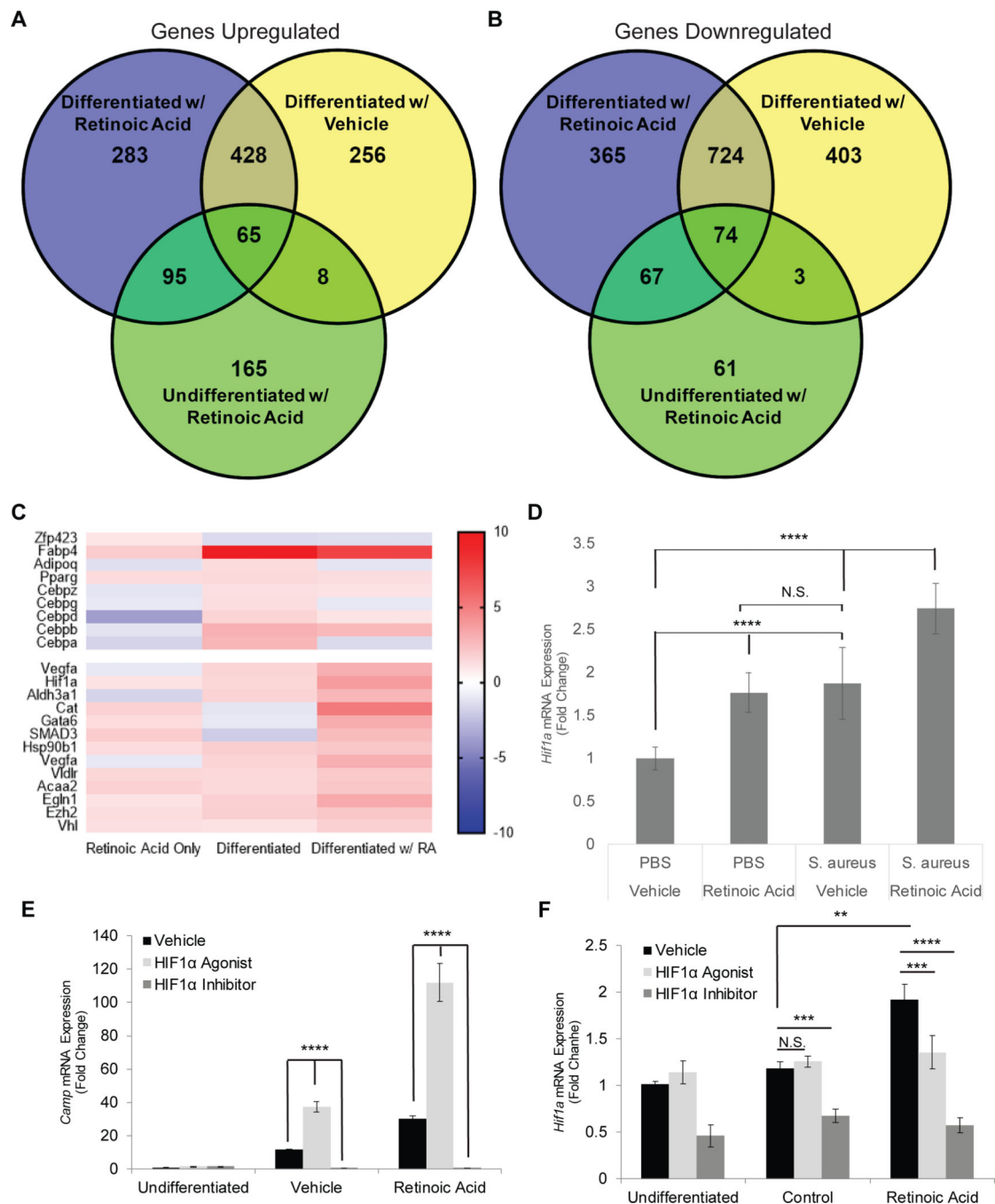
SKH-1 wildtype mice were distally intradermally infected with *S. aureus* or bioluminescent *S. aureus* as on day 0 and treated subcutaneously with RA or olive oil (control) for 3 days (Day 0, 1, and 2). (E) Measurement of bacterial CFU from skin lesion of the infected area on day 3 (N=8) (F) Representative day 3 images of vehicle (top) and RA treated (bottom) and quantification of fluorescence in region of interest (ROI) of skin from images taken by IVIS on days 1, 3, and 6 (N=3 or 4). All data are mean  $\pm$  S.D., \*P<0.05, \*\*P<0.01, \*\*\*P<0.001, \*\*\*\*P<0.0001 using a Student's paired T test.

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**Figure 4. Retinoic acid and HIF1 $\alpha$  coordinate to enhance innate immune defense.**

Transcriptomic comparisons of genes (A) upregulated or (B) downregulated in 3T3-L1 cells differentiated and/or treated with RA for 24 hours. (C) Heat map clustering of adipogenic-related genes and hypoxia response genes. (D) SKH-1 wildtype mice (~8 weeks old) were distally intradermally infected with *S. aureus* on day 0 and simultaneously treated subcutaneously with RA or olive oil (control) for 3 days (n=4 per group). Skin-lesions were collected and gene expression was analyzed by qPCR. (E-F) 3T3-L1 cells were differentiated for 24 hours in the presence of RA supplemented with a HIF1 $\alpha$  agonist

(400  $\mu$ M L-mimosine) or antagonist (5  $\mu$ M acryflavine) along with appropriate controls and mRNA expression of (E) *Camp* or (F) *Hif1a* was determined by qPCR. and mRNA expression of (E) *Camp* or (F) *Hif1a* was determined by qPCR.

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