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REVIEW ARTICLE

Metabolism, fibrosis, and apoptosis: The effect of lipids and their derivatives on keloid formation

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Abstract

Keloids, pathological scars resulting from skin trauma, have traditionally posed significant clinical management challenges due to their persistence and high recurrence rates. Our research elucidates the pivotal roles of lipids and their derivatives in keloid development, driven by underlying mechanisms of abnormal cell proliferation, apoptosis, and extracellular matrix deposition. Key findings suggest that abnormalities in arachidonic acid (AA) synthesis and non-essential fatty acid synthesis are integral to keloid formation. Further, a complex interplay exists between lipid derivatives, notably butyric acid (BA), prostaglandin E2 (PGE2), prostaglandin D2 (PGD2), and the regulation of hyperfibrosis. Additionally, combinations of docosahexaenoic acid (DHA) with BA and 15-deoxy-Δ12,14-Prostaglandin J2 have exhibited pronounced cytotoxic effects. Among sphingolipids, ceramide (Cer) displayed limited proapoptotic effects in keloid fibroblasts (KFBs), whereas sphingosine 1-phosphate (S1P) was found to promote keloid hyperfibrosis, with its analogue, FTY720, demonstrating contrasting benefits. Both Vitamin D and hexadecylphosphorylcholine (HePC) showed potential antifibrotic and antiproliferative properties, suggesting their utility in keloid management. While keloids remain a prevalent concern in clinical practice, this study underscores the promising potential of targeting specific lipid molecules for the advancement of keloid therapeutic strategies.

KEYWORDS

apoptosis, fibrosis, keloid, lipid metabolism, prostaglandins

Abbreviations: cAMP/PKA, Cyclic Adenosine Monophosphate/Protein Kinase A; FAs/FasL, Fas Cell Surface Death Receptor/FAs Ligand; PPAR-γ, Peroxisome Proliferator-Activated Receptor Gamma; PKC/IP3, Protein Kinase C/Inositol Triphosphate; TGF-β1/Smad, Transforming Growth Factor Beta 1/Sma- and Mad-related protein; MMP2/MMP9, Matrix Metalloproteinase-2 and Matrix Metalloproteinase-9; P38-MAPK, P38 Mitogen-Activated Protein Kinase; JNK/ERK, c-Jun N-terminal Kinase/Extracellular signal-Regulated Kinase.

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Key Messages

• Wound healing often involves scar formation, with keloids being pathological scars that are currently difficult to cure effectively. This review introduces a novel perspective on the intricate interplay between lipids, their derivatives, and keloid formation. By synthesising existing research, we offer insights into biochemical pathways and therapeutic implications. Our findings enhance understanding and may aid future research in scar treatment and dermatology.

1 | INTRODUCTION

Keloids represent a pathological scarring that occurs following the healing of skin trauma, typically extending beyond the boundaries of the original wounds and per-sisting without natural regression.^{[1](#page-7-0)} Manifesting as raised, hard-textured stripes or flaky lumps on the skin's surface, they pose a complex clinical management challenge due to notably high recurrence rates.^{[2](#page-7-0)} Most studies suggested that keloid formation is intricately tied to mechanisms of abnormal cell proliferation, apoptosis, and excessive extracellular matrix (ECM) deposition.^{3[–](#page-7-0)5} Numerous recent lipidomics studies underscored the crucial roles of lipids and their derivatives in keloid development, implicating ongoing inflammation, progressive fibrosis, and irregular cell proliferation and apoptosis in keloids. $6,7$ We embarked on a meticulous and systematic investigation into the correlations between fatty acids and their derivatives, sphingomyelin derivatives, phospholipid analogues (such as HePC), and sterol derivatives (such as Vitamin D) and keloids. Our study aims to further illuminate the functions of lipids and their derivatives in keloid scarring and related mechanisms, providing a pivotal reference for devising new clinical interventions for keloids.

2 | ASSOCIATION OF FATTY ACID-BASED LIPID MOLECULES WITH KELOIDS

2.1 | Abnormal metabolism of fatty acids in keloids

2.1.1 | Disturbances in essential fatty acid (EFA) levels may promote keloid formation

Patients with keloids exhibit alterations in plasma levels of several fatty acids $(FAs)⁸ FAs$ $(FAs)⁸ FAs$ $(FAs)⁸ FAs$, classified into saturated and unsaturated fatty acids based on structure, can also be categorised as essential and non-essential fatty acids depending on the synthesis pathway. Furthermore, unsaturated fatty acids subdivide into monounsaturated

and polyunsaturated fatty acids (PUFAs), with most of the EFAs belonging to the PUFA category.⁹

Comparative analyses have revealed disruptions in plasma levels of several PUFAs in keloid patients relative to healthy individuals, suggesting their potential as biomarkers for keloids. $10,11$ Notably, keloid skin demonstrates decreased levels of linoleic acid (LA), but elevated levels of arachidonic acid (AA) compared to normal skin. Regarding dietary intake, keloid patients consume higher than recommended levels of dietary LAs and AAs, specifically between 7 and 11 g/d .^{[8](#page-7-0)}

It is hypothesised that AA metabolism plays a pivotal role in the fatty acid metabolism of keloid patients, with high levels of AA acting as a transport hub for fatty acid metabolism and mechanisms, including inflammation and fibrosis.¹² The free AA released by phospholipase A2 (PLA2) or diacylglycerol (DAG) lipase, activates protein kinase C (PKC), which in turn stimulates PLA2 to further release AA. This process is responsible for the overproduction of pro-inflammatory prostaglandin E2 (PGE2). Moreover, LA can convert to AA, acting as a substrate for g-linolenic acid (GLA) and duomo-g-linolenic acid (DGLA), and the anti-inflammatory prostaglandin E1 (PGE1), to compensate for the over-release of AA, leading to its further reduction. $4,13$ Consequently, an excessive intake of AA stimulates the production of AA, and elevated levels of AA facilitate the production of pro-inflammatory substances, contributing to keloid formation. 14

2.1.2 | Altered synthesis of non-essential fatty acids in keloid scars

Keloids, often considered to be a tumour-like disease, showcase abnormal fibroblast proliferation, with keloid fibroblasts (KFBs) demonstrating one of their major metabolic features as increased lipid synthesis. 15 Notably, fatty acid de-novo synthesis, critical in various human tissues, primarily employs exogenous lipids for generating new structural lipids.^{[16](#page-7-0)} Typically, endogenous fatty acid synthesis is suppressed, maintaining FASN expression at low levels.^{[17](#page-7-0)}

In stark contrast, rapidly proliferating tumour cells exhibit a completely distinct process of de-novo synthe-sis of FAs.^{[18](#page-7-0)} The aberrant synthesis of non-essential FAs in keloids, particularly focusing on key enzymes involved in FA synthesis within the organism—namely sterol regulatory element-binding protein-1 (SREBP-1) and FASN—has become a focal point of research.^{[19](#page-7-0)} SREBP-1, a critical transcription factor, modulates the synthesis of steroids and lipids in the de-novo synthesis pathway, and directly regulates the expression of pivotal enzymes.^{[20](#page-7-0)} FASN, instrumental in providing the energy necessary for the survival of proliferating cells and a key enzyme in the abnormal metabolism of endogenous fatty acids, has been observed to be overexpressed in various human epithelial cancers, including those of the prostate, ovaries, colon, lung, endometrium, and stom-ach.^{[21](#page-7-0)} Subsequent experiments indicated higher mRNA expression of SREBP1 and FASN in KFBs than in normal fibroblasts (NFBs), implying that the abnormal synthesis of endogenous FAs might contribute to keloid formation.^{[11](#page-7-0)}

2.2 | Lipid-associated modulation in keloid scars

2.2.1 | Butyric acids typifying fatty acids in the inhibition of keloid fibrogenesis

In comparison with physiologically normal skin, keloids discernibly exhibit divergent lipid profiles, with 27 disparate FAs and a total of 30 differentially expressed FAs identified. Specifically, eight short-chain fatty acids (SCFAs) manifest alterations, with substantial diminutions in the expressions of butyric, isobutyric, malonic, valeric, and succinic acids.^{[11](#page-7-0)}

SCFAs, produced through anaerobic bacterial fermentation of dietary fibre within the colon, especially butyrate and propionate, are known to exhibit histone deacetylase (HDAC) inhibitory activity.^{[22](#page-7-0)} Engaged in inflammation regulation, SCFAs curtail mast cell activation when in deficit, thereby limiting the production of inflammatory mediators. 23 The antifibrotic potential of butyrate and other HDAC inhibitors, as evidenced across various mesenchymal stem cells (MSCs), resides in their ability to mitigate cell proliferation, collagen synthesis, and α-Smooth Muscle Actin $(\alpha$ -SMA) expression through histone acetylation modulation.^{[24](#page-8-0)}

An in-vitro exploration involving the co-culturing of butyric acid (BA) with keloid KFBs demonstrated that BA inhibited fibroblast proliferation and type III collagen

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expression while concurrently augmenting COX-1 expression and facilitating PGE2 production. 25 This suggested that the scarcity of SCFAs in keloid scars modulates apoptosis, proliferation, and differentiation of KFBs, further exacerbating collagen deposition. Additionally, a potential therapeutic synergism for keloids was realised through the amalgamation of docosahexaenoic acid (DHA) and BA, which seemed to amplify inhibitory effects on α-SMA, collagen types I and III, TGF-β1, and type I TGF-β1 receptors, concomitantly inhibiting cell proliferation, precipitating apoptosis, and dismantling stress fibers. 25 A secondary pathway, initiated by BAactivated FAs in colonocytes, posits that a synergistic apoptotic effect is feasible through an intrinsic mitochondrial pathway[.26](#page-8-0)

The antifibrotic properties of DHA observed in human peritoneal fibroblasts encompass the inhibition of the expression of TGF-β1, VEGF, and type I collagen. Given that DHA acts as a ligand for peroxisome proliferator-activated receptor γ (PPAR γ) and DHA-derived lipid mediators robustly actuate PPARγ, extensive investigations corroborate that PPARγ expression or activation subdues the expression of type I collagen, α-SMA, and TGF-β across various fibroblast populations. $27,28$

2.2.2 | HePC: A phospholipid analogue suppressing KFBs' anomalous proliferation

Hexadecylphosphorylcholine (HePC), recognised as an alkyl phospholipid analogue, has garnered interest as an innovative antiproliferative agent, exerting its effects on the plasma membrane. 29 29 29 HePC, in vitro, substantively inhibits cellular growth across a spectrum of tumour cell lines, further manifesting antitumour activity in-vivo. Investigations elucidated that HePC disrupts the proliferation of human epidermal keratinocytes and stifles phosphatidylcholine synthesis in-vitro, establishing its efficacy in the treatment of non-neoplastic, hyperproliferative skin disorders, inclusive of psoriasis.[30](#page-8-0)

Its potential applicability in keloid therapy has been explored. In-vitro cultivation revealed that HePC impedes fibronectin synthesis and alters the constitution of the intracellular plasma membrane and membrane protein activity by obstructing phosphatidylcholine biosynthesis. This alteration culminates in marked antiproliferative characteristics by impeding cell cycle progression.[31](#page-8-0) Nevertheless, the specific role of HePC in modulating keloid fibrosis and the keloid ECM demands additional scrutiny and is primed for subsequent investigative pursuit Table [1.](#page-3-0)

TABLE 1 The regulatory roles of various molecules in the progression of keloids.

3 | RELEVANCE OF DERIVATIVES OF BOTH FAs AND SPHINGOLIPIDS TO KELOIDS

3.1 | Inhibition of keloid progression through FA derivatives

3.1.1 | Strategic curtailment of keloid formation facilitated by FA derivatives: A spotlight on prostaglandins

Keloids, pathologically hyperproliferative fibrotic lesions, betray a significant correlation with the multifaceted actions of eicosanoids, cytokines, and reactive oxygen species (ROS). AA, serving as a pivotal substrate for eicosanoids, including leukotrienes (LTs), prostaglandins (PGs), lipoxins (LXs), and thromboxanes (TXs), bespeaks an indelible signature in the complex pathophysiology of keloids. 32 Furthermore, notwithstanding its profound pro-inflammatory modulation in PUFAs metabolism, emerging evidence posits a consequential role for ω-3 FAs, particularly in attenuating anti-apoptotic signaling and obfuscating pro-inflammatory cues. Derivatives such as Docosapentaenoic acid (DPA) and DHA, stemming from alpha-linolenic acid (ALA), have been implicated in the mitigation of inflammatory cytokine synthesis, inclusive of interleukin 6 (IL-6) and tumour necrosis factoralpha (TNF- α).^{[7](#page-7-0)}

PGs, quintessential FA derivatives, are implicated in orchestrating a myriad of physiological and pathological

cascades. 33 Their biogenesis, a subject of meticulous enzymatic control, commences with the conversion of membrane phospholipids to AA by phospholipases, followed by the transformation of AA to prostaglandin H2 (PGH2) via cyclooxygenase (COX) pathways.^{[34](#page-8-0)} PGH2 thus serves as a progenitor for biologically salient mediators such as thromboxane A2, PGE2, and prostacyclin. 35

COX-2, an inducible isoform of COX, facilitates the synthesis of PGE2 and nitric oxide (NO) via nitric oxide synthase (NOS), particularly under the auspices of proinflammatory mediators like TNF- α , lipopolysaccharides, and interleukin-1 beta $(IL-1\beta)$.^{[36](#page-8-0)} This orchestrated inflammatory milieu begets subsequent collateral damage to adjacent tissue cells at the wound locus, thereby decelerating the wound healing trajectory.

PGE2, omnipresent among PGs within the human physiology, and a noteworthy eicosanoid fibroblast product, mitigates fibroblast functionalities through autocrine secretion.^{[37](#page-8-0)} Anomalies in the synthesis or degradation of PGE2 have been correlative with a spectrum of pathological states, encompassing persistent inflammation and tumorigenesis.^{[38,39](#page-8-0)} Moreover, PGE2 has been attributed a bifurcated role in keloid development, concurrently advancing inflammatory progression and exhibiting antifibrotic function, rendering it a nexus of contemporary research interest.

Prostaglandin D2 (PGD2), synthesised in both central nervous and peripheral tissues, safeguards pivotal roles in the modulation of inflammation and the preservation of homeostasis.[40](#page-8-0) While mast cells are preeminent

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FIGURE 1 Illustrating the abnormal synthesis process of arachidonic acid in keloids, as well as the synthesis process of prostaglandins mediated by arachidonic acid. It also covers the pathways in which prostaglandin E2 is involved in regulating fibrosis in keloids. AC, Adenylyl Cyclase; cAMP, Cyclic Adenosine Monophosphate; CREB, cAMP Response Element-Binding Protein; MMP, Matrix Metalloproteinase; PKC, Protein Kinase C; PGG2, Prostaglandin G2; PGH2, Prostaglandin H2; PGES, Prostaglandin E Synthase; PGD2, Prostaglandin D2; PGE2, Prostaglandin E2; PKA, Protein Kinase A; PGJ2, Prostaglandin J2; TIMP-1, Tissue Inhibitor of Metalloproteinases-1.

synthesisers of PGD2 in peripheral tissues, other leukocytes, such as dendritic cells (DC) and T-helper 2 cells (Th2), also harbor the capability to synthesise PDG2. $41,42$ PGD2, albeit relatively unstable with a plasma half-life \sim 30 min, undergoes subsequent metabolism to yield lipid variants such as 15 deoxy-PGJ 2, which hold particular relevance to keloids.^{[43](#page-8-0)} Additionally, PGD2 can undergo enzymatic metabolism, facilitated by aldo-keto reductase 1C3 (AKR1C3), culminating in the formation of 9α,11β-PGF2. Furthermore, 15 deoxy-PGJ 2, engendered spontaneously through PGD2 dehydration, vies in production with other PGD2-derived entities, including 9α,11β-PGF2^{[11](#page-7-0)} Figure 1.

3.1.2 | In-depth examination of prostaglandin-mediated mechanisms in alleviating keloid fibrosis and anti-apoptosis

PGE2, interfacing with its receptor EP2 within fibroblasts, instigates the cAMP/Protein Kinase A (PKA) signaling cascade, ensuing in the concomitant inhibition of cellular proliferation, migration, collagen synthesis, and myofibroblast differentiation[.44,45](#page-8-0) Consequently, PGE2 heralds an anti-fibrotic impact across an array of organ systems.^{[46](#page-8-0)} Additionally, PGE2 propagates tissue remodeling within skin tissues, while a diminution in dermal PGE2 levels has been associated with an amplification in collagen synthesis and scar ontogenesis.[47](#page-8-0) Empirical investigations have illuminated a potential correlation between a diminished proficiency of KFBs to synthesise PGE2 and the overexpression of Transforming Growth Factor-beta 1 (TGF-β1), subsequently culminating in a downtrend in the endogenous expression of COX-2 and PGE2 within KFBs.^{[48](#page-8-0)} This intimated that defects in endogenous PGE2, induced by TGF-β1, may be pivotal in keloid fibrosis. The regulatory equilibrium of Matrix Metalloproteinase/Tissue Inhibitor of Metalloproteinase (MMP/TIMP) expression is also modulated by PGE2, escalating the expression of MMP-2 and MMP-9 whilst diminishing TIMP-1 expression.^{[49](#page-8-0)} Moreover, PGE2, upon activation of EP2-integrated with the cAMP signaling cascade-augments cAMP levels, inhibiting collagen synthesis in dermal fibroblasts induced by TGF-β1. The profibrotic effects of TGF-β1 are contingent upon the Smad signaling pathway, whereas PGE2 obliterates TGFβ1-induced Smad 2/3 phosphorylation and Smad 2/3/4 complex formation, as well as nuclear translocation.

Consequently, PGE2's anti-fibrotic efficacy is ascribed to the inhibition of the TGF- β 1/Smad signaling pathway.^{[50,51](#page-8-0)}

PGD2 orchestrates physiological functionalities via its distinct receptors, DP1 and DP2-synonymously acknowledged as Chemoattractant Receptor-Homologous Molecule expressed on Th2 Cells (CRTH2).⁵² Activation of the DP1 receptor elicits an upsurge in cAMP levels and intracellular PKA activation, a phenomenon analogous to that of EP2. Albeit CRTH2's capacity to amalgamate with Gi proteins, inhibiting cAMP synthesis and enhancing intracellular Ca2+ concentration, it is imperative to note that CRTH2 is predominantly expressed within Th2 cells. 53 This revelation posits that receptor activation by PGD2 impedes the fibrotic process.

Fifteen deoxy-PGJ 2, a ligand that actuates the peroxisome proliferator-activated receptor-gamma (PPAR-γ), inhibits the NF-κB pathway and induces oxidative stress, participates in numerous biological processes.^{[54](#page-8-0)} 15 deoxy-PGJ 2 mediates its pro-apoptotic and anti-inflammatory impacts by actuating PPAR-γ and obliterating NF- $κB$.^{[55](#page-8-0)} The concentration of 15 deoxy-PGJ 2 is instrumental in determining its inflammatory impact: it amplifies chemokine-induced chemotaxis in eosinophils at suboptimal concentrations, whereas at maximal concentrations, it induces apoptosis and suppresses eosinophil survival.^{[56](#page-8-0)} Distinct from its PPAR-γ-dependent manner, various studies have discerned 15 deoxy-PGJ 2 as possessing anti-fibrotic properties.^{[57,58](#page-8-0)} Therapeutic application of 15 deoxy-PGJ 2 to KFBs facilitated an enhancement in ROS formation, P38-MAPK activation, and Superoxide Dismutase 1 (SOD1) transcription, thereby revealing cytotoxic impacts. Furthermore, aldo-keto reductase 1C3 (AKR 1C 3) – the enzyme responsible for metabolising PGD2 – was observed to be overexpressed in keloids relative to normotrophic skin. This enzyme, which metabolises PGD2 to 9α,11β-PGF 2, is thereby hypothesised as one of the defensive mechanisms of KFBs.^{[54](#page-8-0)} Consequently, inhibition of AKR 1C 3 could emerge as a potent local therapeutic strategy for keloids.

3.2 | Sphingomyelin rheostat: Elucidating its paramountcy in keloid development

Sphingolipids, quintessential constituents of cellular membranes, pervade the plasma membranes of a plethora of mammalian cells, where their metabolites including biologically salient signaling molecules such as ceramide (Cer), sphingosine (Sph), and sphingosine 1-phosphate (S1P)—orchestrate pivotal signaling cascades implicating cell proliferation and apoptosis. The dynamic triad of Cer, Sph, and S1P has been metaphorically designated the 'sphingomyelin rheostat',

attributable to the intrinsic link of their dynamic equilib-rium to cellular vitality and apoptosis.^{[59](#page-8-0)}

3.2.1 | Interconversion within sphingolipid triangles and Sph's cytotoxicity to KFBs

Cer generation, a cornerstone of sphingolipid metabolism, is effectuated through two avenues: de-novo synthesis within the endoplasmic reticulum and sphingolipids' hydrolysis via acid sphingomyelinase (ASM).⁶⁰ Postsynthesis, Cer, in the milieu of ceramidase, undergoes deacylation, culminating in Sph production, which is subsequently shuttled to the Golgi apparatus via ceramide transporter proteins. 61 Thereupon, Sph, engaging a phosphate molecule in the presence of sphingosine kinase (SphK), begets S1P, which, through the action of sphingosine phosphate phosphatase, reverts to Sph in a reversible manner. Both Sph and Cer, imperative elements of the stress-responsive regulatory system, embody inhibitory and pro-apoptotic functionalities, whereas S1P is associated with pro-growth and anti-apoptotic activities.^{[62](#page-9-0)}

3.2.2 | Pro-apoptotic potency of ceramide: Dampened efficacy in keloids

Cer propels apoptosis, acting as a secondary messenger in the FAs-mediated apoptotic conduit, concurrently impeding fibroblast (FB) proliferation and enhancing their apoptosis.[63](#page-9-0) Moreover, gamma interferon (IFN-γ) throttles FB proliferation via the potentiation of Cer secretion. Nevertheless, in the keloid context, Cer's pro-apoptotic potency is nullified, partially due to the augmented expression of insulin-like growth factor-1 (IGF-I) in expeditiously proliferating KFBs, with keloids also evidencing overexpression of the insulin-like growth factor-1 recep-tor (IGF-IR).^{[64](#page-9-0)} Furthermore, IGF-I shields KFBs from FAs-mediated apoptosis, and an additional impediment to keloid mitigation is the obstruction of the FAsmediated apoptotic pathway, wherein signal interruption is manifested upstream of Cer, precluding its receipt of the signal and subsequent pro-apoptotic execution. $65,66$

3.2.3 | Divergent impacts of S1P and its analogue FTY720 on keloids

S1P, a crucial signaling transducer, wields substantial influence over cellular growth, differentiation, proliferation, and apoptosis.[67](#page-9-0) S1P receptors (S1PRs), G protein-coupled receptors targeted by S1P, encompass five subtypes: S1PR1–5, which modulate biological activities such as

immune cell trafficking, angiogenic endeavors, and mito-sis amid cellular proliferation.^{[68](#page-9-0)} In keloid tissue, S1PR1/ S1PR2 mRNA and protein expressions were elevated by \sim 1.5-fold and a substantive margin, respectively, compared to their normal skin tissue counterparts. By binding to S1PR, S1P oversees lymphatic cell transport and inflammatory cell recruitment to, and retention within inflamed tissues, thereby delineating a connection between S1P, collagen synthesis, and fibrosis during keloid development and progression.^{[69](#page-9-0)}

Notably, FTY720, an immunosuppressant and S1P analogue, exerts anti-inflammatory impacts and influences immune cell differentiation pathways, embodying a potentially potent actor in pathological scarring treat-ment.^{[70](#page-9-0)} Experimental data showed that FTY720 imparts distinct pleiotropic impacts on hyperproliferative scarring dermal fibroblasts (HSFs) and normal fibroblasts (NFBs), implicated diminished cellular viability, G0/G1 phase cell cycle arrest, apoptosis instigation, and inhibition of migration, contraction, and α -SMA, collagen I and III expression, thereby underscore its prospective utility as an anti-fibrotic agent. 71

3.2.4 | The influence of vitamin D on fibrotic processes in keloids

Vitamin D (VD), a group of structurally analogous steroid derivatives, and its principal metabolite, 1,25 dihydroxyvitamin D3 (1,25 D), play pivotal roles in

processes such as calcium homeostasis, hormone secre-tion, and cell proliferation and differentiation.^{[72](#page-9-0)} Notably, in keloid skin, mRNA and protein expression of the functional vitamin D receptor (VDR)—a liganddependent transcription factor within the steroidogenic nuclear receptor gene family and mediator of 1,25 D effects—is significantly reduced compared to normal skin.^{[73](#page-9-0)} 1,25 D distinctly inhibits the proliferation of KFBs without triggering cytotoxicity. Ample research has pointed towards the involvement of VD in TGFβ1-related fibrosis in recognised VD target cells or organs.^{[74,75](#page-9-0)} In keloids, 1,25 D suppresses the TGFβ1-induced extracellular matrix production in KFBs, thereby reducing the expression of matrix proteins like type I collagen, Fibronectin (FN), and α-SMA. Furthermore, aside from its fundamental antiproliferative and antifibrotic actions, 1,25 D uniquely induces the manifestation and secretion of hepatocyte growth factor in keloid fibroblasts and alleviates initial fibrosis by pro-moting MMP-9 expression^{[76,77](#page-9-0)} Figure 2.

4 | CONCLUSION

This article illuminates the intricate connections between lipids, lipid derivatives, and keloids, revealing a significant link with lipid metabolism in the progression of keloids. Abnormalities in AA synthesis and the synthesis of non-essential FAs stand out as key characteristics of keloid scars. Delving into the impact of lipids on keloids,

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detrimental regulation primarily through BA, PGE2, and PGD2 is highlighted, particularly noting their role in inhibiting hyperfibrosis. Meanwhile, a combination of DHA with BA and 15-deoxy-Δ12,14-Prostaglandin J2 exhibits noteworthy cytotoxicity. In the sphere of sphingolipid metabolism, Cer, Sph, and S1P emerge as pivotal. Specifically, Cer struggled to manifest its pro-apoptotic effects in KFBs. Contrarily, S1P promoted keloid hyperfibrosis, while its analogue, FTY720, exhibited opposing effects. VD and HePC both offered antifibrotic and antiproliferative effects, thereby contributing to the negative regulation of keloid formation.

Despite being a common concern in plastic surgery and dermatology, keloids continue to be a challenge with no highly specific and effective treatment currently available; existing management typically navigates through surgical excision and radiotherapy. The explorations encapsulated in the current body of research provide a foundational understanding of the relationship between lipids, their derivatives, and keloids. Thus, targeting specific lipid molecules and derivatives opens a promising avenue for the development of novel interventions, aiming to enhance and refine therapeutic approaches to managing keloid scars.

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CONFLICT OF INTEREST STATEMENT

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DATA AVAILABILITY STATEMENT

Data sharing not applicable to this article as no datasets were generated or analysed during the current study.

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