

## 综述

# 硬脂酰辅酶A去饱和酶1在代谢性疾病中的功能及相关机制研究进展

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**摘要:** 代谢性疾病主要表现为能量稳态失调, 是全球面临的重大健康挑战。代谢性疾病人群常伴有与脂质代谢紊乱相关的并发症, 如肥胖和非酒精性脂肪性肝病。关注脂质代谢核心基因有助于代谢性疾病及其并发症的防治工作。硬脂酰辅酶A去饱和酶1 (stearoyl-CoA desaturase 1, SCD1) 是一种脂代谢关键酶, 可将饱和脂肪酸转化为单不饱和脂肪酸, 在能量稳态、糖脂代谢、自噬和炎症反应等众多生理和病理过程中发挥关键调控作用。*Scd1* 的异常转录和表观遗传激活可通过调控多条信号轴导致脂质异常积累, 从而促进肥胖、非酒精性脂肪性肝病、糖尿病和癌症发展。本文全面综述了SCD1作为代谢枢纽基因在常见生理和病理事件中的关键作用, 并进一步从转化视角出发讨论了多学科交叉背景下SCD1新型抑制剂的潜在开发路径, 以期靶向SCD1防治代谢性疾病提供新视野和新思路。

**关键词:** 代谢性疾病; 硬脂酰辅酶A去饱和酶1; 糖脂代谢; 代谢枢纽

## Advances in the function and mechanisms of stearoyl-CoA desaturase 1 in metabolic diseases

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**Abstract:** Metabolic diseases characterized by an imbalance in energy homeostasis represent a significant global health challenge. Individuals with metabolic diseases often suffer from complications related to disorders in lipid metabolism, such as obesity and non-alcoholic fatty liver disease (NAFLD). Understanding core genes involved in lipid metabolism can advance strategies for the prevention and treatment of these conditions. Stearoyl-CoA desaturase 1 (SCD1) is a key enzyme in lipid metabolism that converts saturated fatty acids into monounsaturated fatty acids. SCD1 plays a crucial regulatory role in numerous physiological and pathological processes, including energy homeostasis, glycolipid metabolism, autophagy, and inflammation. Abnormal transcription and epigenetic activation of *Scd1* contribute to abnormal lipid accumulation by regulating multiple signaling axes, thereby promoting the development of obesity, NAFLD, diabetes, and cancer. This review comprehensively summarizes the key role of SCD1 as a metabolic hub gene in various (patho)physiological contexts. Further it explores potential translational avenues, focusing on the development of novel SCD1 inhibitors across interdisciplinary fields, aiming to provide new insights and approaches for targeting SCD1 in the prevention and treatment of metabolic diseases.

**Key words:** metabolic diseases; stearoyl-CoA desaturase 1; glycolipid metabolism; metabolic hub

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代谢性疾病流行现状严峻,严重影响全球公众健康,加重医疗负担。代谢性疾病发生和发展的特征之一是脂质代谢功能的失调,以及进一步引发的慢性炎症激活、氧化应激和异常自噬等<sup>[1–4]</sup>。硬脂酰辅酶A去饱和酶1 (stearoyl-CoA desaturase 1, SCD1) 是脂质合成中的关键限速酶,通过产生单不饱和脂肪酸(monounsaturated fatty acids, MUFAs)维持细胞内脂质稳态,协调糖脂代谢、炎症和自噬,是调节代谢的中心因子<sup>[5–8]</sup>。越来越多的研究表明,SCD1的异常表达与肥胖、非酒精性脂肪性肝病(non-alcoholic fatty liver disease, NAFLD)和2型糖尿病(type 2 diabetes mellitus, T2DM)等多种代谢性疾病的风险增加有关<sup>[9–14]</sup>。进一步研究发现抑制SCD1活性能够减弱脂肪酸的再酯化和糖异生,并促进脂肪细胞中 *Atgl* 和 *Hsl* 等脂解生物标志物的表达<sup>[15]</sup>。同时,两项以 NAFLD 和非酒精性脂肪性肝炎(non-alcoholic steatohepatitis, NASH)患者为受试者的临床试验揭示了 Aramchol (Arachidyl amido cholanoic acid, 一种肝脏 SCD1 部分抑制剂)能够显著减少肝脏脂质堆积,改善肝纤维化<sup>[16–18]</sup>。其中, Aramchol 通过下调 *Scd1* 基因表达、上调 PPAR $\gamma$  表达,减少肝星状细胞中 I 型胶原的表达和分泌,最终在 NASH 患者中达到抗纤维化效果<sup>[19]</sup>。

综上,探明 SCD1 如何调控上述复杂信号网络是理解其在代谢性疾病发病机制中作用的核心。因此,本文综述了 SCD1 作为代谢枢纽基因在生理和病理事件中的关键影响,并进一步讨论了针对多种组织和器官的新型强效 SCD1 抑制剂用于治疗代谢性疾病的潜在开发路径,以期为靶向 SCD1 防治代谢性疾病提供新视野。

## 1 SCD1 分子特征

SCDs 是一类高度保守的脂肪酸去饱和酶家族,亦属于亚铁去饱和酶家族。迄今为止,在小鼠中已鉴定出具有高度氨基酸序列同源性的四种 SCD 亚型(SCD1~4),在大鼠中已鉴定出两种 SCD 同工酶(SCD1~2)<sup>[20,21]</sup>。其中,SCD1 是一种位于内质网(endoplasmic reticulum, ER)的膜蛋白,具有四个跨膜 $\alpha$ 螺旋(TM1~TM4),这些螺旋在结构上呈锥形排列<sup>[22]</sup>。在跨膜区域,大部分氨基酸是疏水性的,但在 TM4 的中心位置有一个保守的精氨酸残基(Arg249)以稳定 SCD1 结构。SCD1 的 N 端和 C 端位于细胞质侧<sup>[22,23]</sup>。在 SCD1 结构域中,高度保守的

组氨酸残基通过结合二价铁离子形成催化中心,通过电子传递参与 SCD1 的脂肪酸去饱和反应<sup>[22–24]</sup>(如图 1A)。此外,SCD1 发挥活性需要两个膜内蛋白质的辅助,即细胞色素 b5 还原酶(b5 reductase, b5R)和细胞色素 b5 (cyt b5)<sup>[22]</sup>。b5R 在其 N 端有一个单个跨膜 $\alpha$ 螺旋,以及由 N 端和 C 端两部分组成的水溶性胞质区域。N 端部分结合黄素腺嘌呤二核苷酸(FAD),C 端部分结合 NADH<sup>[22]</sup>。b5R 的主要还原因子是 cyt b5,它包含一个水溶性胞质区域和 C 末端的单个跨膜区域,其中含有一个 b 型血红素,由两个轴向组氨酸配体连接。b5R 和 cyt b5 都通过其单个跨膜区域锚定在 ER 膜上。从 NADH 开始,通过 b5R 和 cyt b5,与 SCD1 相互作用形成电子传递链,催化饱和酰基辅酶 A 上的双键形成<sup>[22,23]</sup>。

SCD1 在肝脏、脂肪、胰岛、大脑和骨骼肌等各类组织中广泛表达<sup>[25–35]</sup>(图 2)。在人类、恒河猴、大鼠和小鼠中,SCD1 的蛋白质序列同源性为 91.3% (图 1B),这表明 SCD1 通过脂质代谢调节机体内稳态的作用在进化上是保守的。SCD1 的转录受到如葡萄糖、饱和脂肪酸(saturated fatty acids, SFAs)、胰岛素、固醇调节元件结合蛋白 1 (sterol regulatory element-binding protein 1, SREBP1)、过氧化物酶体增殖物激活受体 $\alpha$  (peroxisome proliferators-activated receptor  $\alpha$ , PPAR $\alpha$ ) 和肝 X 受体(liver X-receptor, LXR)等营养、激素和环境多种因素的严格调控<sup>[36–40]</sup>。其中,瘦素、雌激素和多不饱和脂肪酸(polyunsaturated fatty acids, PUFAs)能够抑制 *Scd1* 基因转录<sup>[41–43]</sup>。比如,来自脂肪组织的 PUFAs 通过与启动子区域的 60 bp PUFAs 反应元件结合可下调小鼠肝脏 *Scd1* 启动子活性,抑制 *Scd1* 基因表达<sup>[44]</sup>。此外,激活 LXR 可上调人类间充质基质细胞中 SCD1 蛋白表达,同时防止过量 SFAs 造成的脂毒性损伤(如氧化应激和炎症反应)<sup>[45]</sup>。有趣的是,随着 SFAs 的水平升高,SCD1 的蛋白表达和活性被上调,增强机体对 SFAs 的去饱和化以维持脂质稳态<sup>[45,46]</sup>。

## 2 SCD1 的生理功能

SCD1 广泛调控体内多种生理和病理功能<sup>[47]</sup>。锚定在 ER 的整合蛋白 SCD1 通过催化 SFAs 的碳 9 和碳 10 之间形成顺式双键,将 SFAs (如硬脂酸)转化为 MUFAs (如油酸)<sup>[47,48]</sup>。SCD1 能够促进体内的脂质代谢,保护癌细胞免受 SFAs 引发的脂毒性影响,从而增强肿瘤细胞增殖和再生,提高其存活

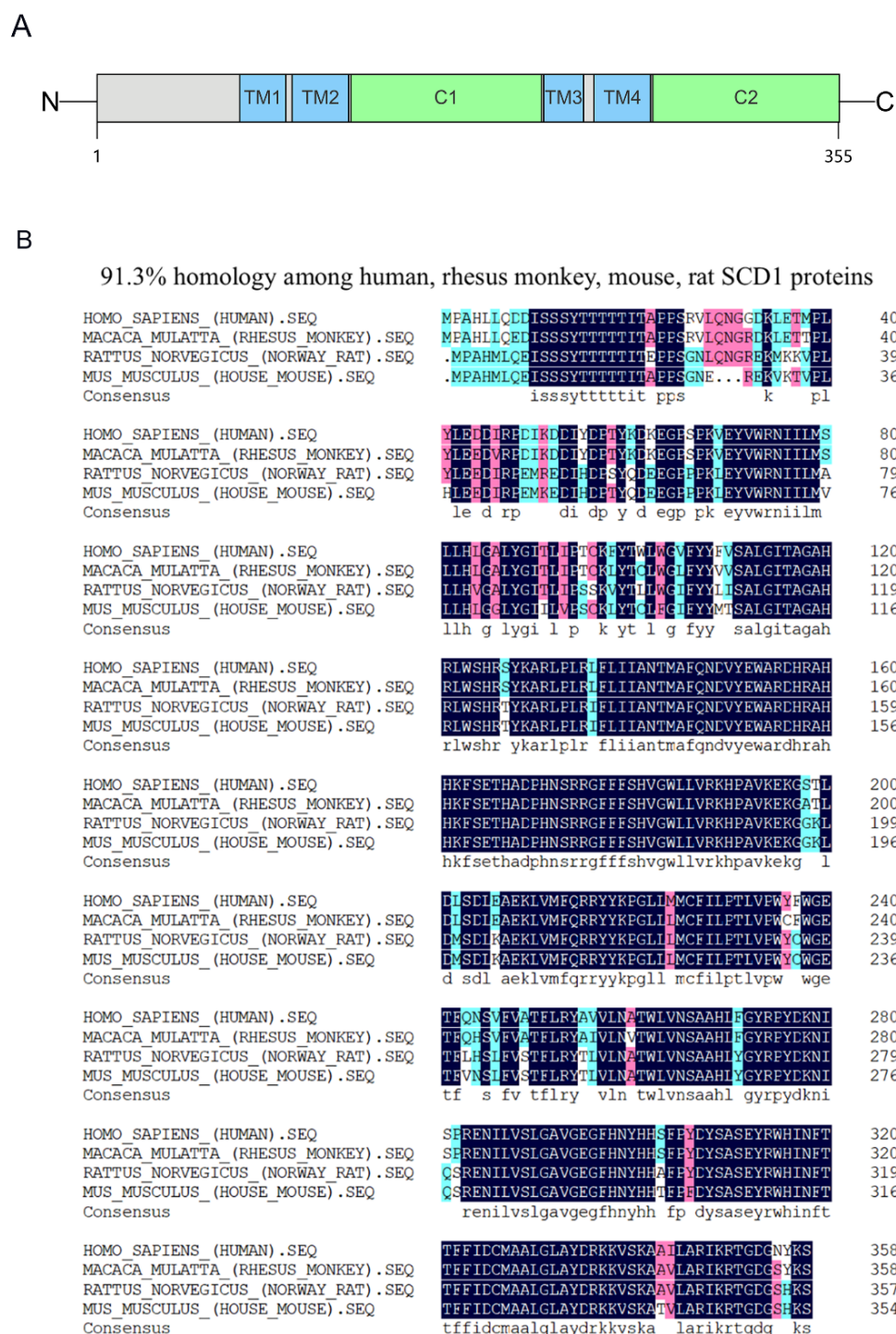


图 1. SCD1 分子结构域及在各物种间蛋白序列比对(参考 NCBI 数据库绘制)

Fig. 1. SCD1 molecular domain map and protein sequence alignment among different species (Drawn by NCBI database). *A*: The molecular structure domain diagram highlights the critical functional regions of the SCD1 protein. N-terminal signal peptide is represented in gray; 4 transmembrane  $\alpha$  helices (TM1-TM4) are represented in blue; and 2 histidine clusters (C1-C2) are represented in green. These domains show a high degree of consistency across species. *B*: This alignment compares SCD1 protein sequences from human (NP\_005054.3), rhesus monkey (XP\_015003410.1), mouse (NP\_033153.2) and rat (NP\_631931.2). Among the 4 sequences, 2 identical residues are represented in blue; 3 identical residues are represented in red; and 4 identical residues are represented in black.



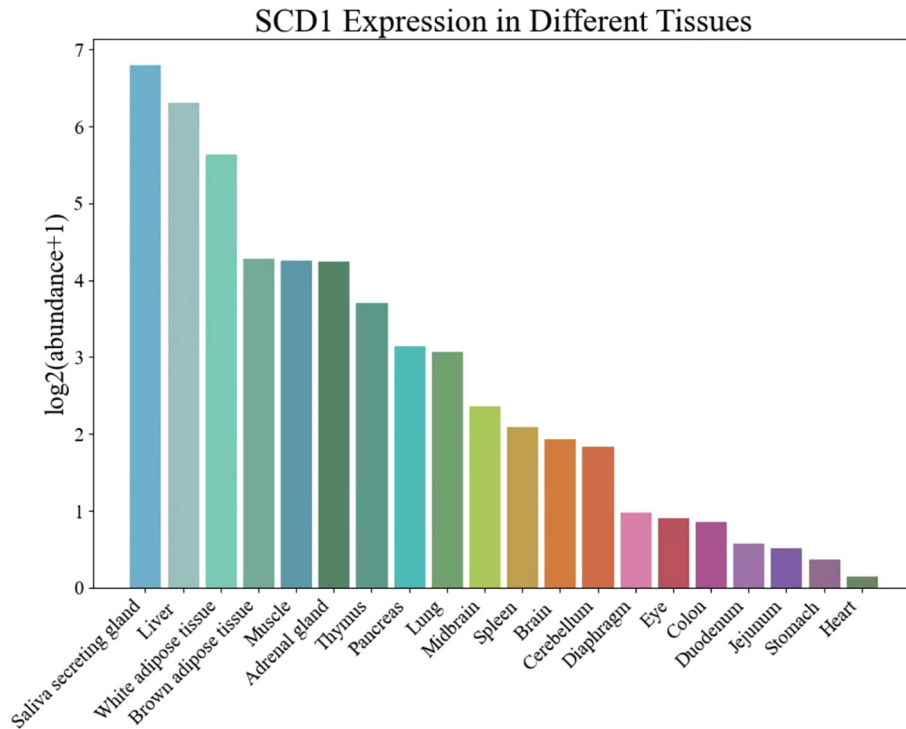


图 2. SCD1 小鼠各组织特异性分布(参考 NCBI 和 PaxDb 数据库绘制)

Fig. 2. Tissue specific distribution of SCD1 in mice (Drawn by NCBI and PaxDb databases).

率<sup>[8, 49, 50]</sup>。相反, SCD1 的失活将降低透明细胞肾细胞癌的细胞增殖并促使其凋亡<sup>[51]</sup>。与此同时, SCD1 在葡萄糖代谢、自噬和炎症中亦发挥着广泛影响<sup>[6, 9, 47, 52, 53]</sup>。

## 2.1 SCD1 和糖代谢

小鼠全身 SCD1 敲除可增强骨骼肌、棕色脂肪组织(brown adipose tissue, BAT)和心脏等组织的葡萄糖摄取, 促进糖原积累, 增强胰岛素敏感性<sup>[54, 55]</sup>。其中, 骨骼肌是人体最大的代谢器官, 直接调节全身葡萄糖稳态和胰岛素敏感性<sup>[56]</sup>。在骨骼肌过表达 SCD1 小鼠中, 骨骼肌胰岛素受体底物 1 (insulin receptor substrate 1, IRS1)酪氨酸磷酸化和 Akt1 丝氨酸 473 磷酸化水平下调, 提示胰岛素信号传导途径受损<sup>[54, 55]</sup>。相反, 在骨骼肌敲除 SCD1 小鼠中, 骨骼肌胰岛素受体、IRS1、IRS2 和 Akt 磷酸化水平上调, GLUT4 介导葡萄糖转运增强<sup>[54, 55]</sup>。其潜在机制可能是 SCD1 改变了体内如游离脂肪酸(free fatty acids, FFAs)和神经酰胺等脂质含量, 进一步抑制了 IRS1-Akt 信号轴活性及其相关的胰岛素敏感性关键因子(如 AMPK 和 PTP-1B)的激活, 最终导致葡萄糖代谢稳态失衡<sup>[54, 57, 58]</sup>。

高碳水化合物饮食(high carbohydrates diet, HCD)通过 SCD1 引起肝脏脂质生成, 并且这种效应

受到 SREBP1、碳水化合物反应元件结合蛋白(carbohydrate response element binding protein, ChREBP)和 LXR 等应答元件的调节<sup>[36]</sup>。特异性敲除 HCD 喂养小鼠肝脏 SCD1 可增强机体葡萄糖摄取, 并促进肝脏成纤维细胞生长因子 21 (fibroblast growth factor 21, FGF21)的合成和分泌<sup>[59]</sup>。已知 FGF21 能够增强全身胰岛素敏感性并调节全身脂质代谢<sup>[59]</sup>。进一步实验证明, 原代肝细胞经油酸处理后 PGC-1 $\alpha$  表达下调, 并部分抑制了 SCD1 抑制剂诱导的 FGF21 表达<sup>[59]</sup>。另有研究发现, 在 AML12 细胞中 SCD1 的缺失引发 ER 应激, 并通过过量 SFAs 激活 mTORC1-PGC-1 $\alpha$  轴增加 FGF21 蛋白表达<sup>[60]</sup>。已知胰岛素样生长因子结合蛋白 1 (insulin-like growth factor binding protein 1, IGFBP1)作为另一种改善葡萄糖耐量的肝脏细胞因子, 能够增强  $\beta$  细胞增殖、胰岛素信号传导和葡萄糖摄取<sup>[61]</sup>。在 HCD 化合物喂食期间, SCD1 肝脏敲除小鼠肝脏中 IGFBP1 蛋白水平显著增加<sup>[60]</sup>。这种 IGFBP1 水平的变化可能受油酸调节下 mTORC1-FGF21 信号轴调控<sup>[62]</sup>。

## 2.2 SCD1 和脂代谢

SCD1 减弱脂肪组织棕色化。研究发现, 敲除脂肪干细胞中 *Scd1* 基因可增加琥珀酸积累, 增强线

粒体复合体II活性,促进脂肪干细胞向米色脂肪细胞的分化<sup>[63]</sup>。在高脂饮食(high fat diet, HFD)喂养小鼠的腹股沟脂肪(inguinal white adipose tissue, iWAT)中, *Scd1* 基因突变上调 *Ucp1* 和 *Pparγ* 等产热基因转录,诱导脂肪细胞棕色化,降低 iWAT 质量<sup>[63]</sup>。有趣的是,通过冷暴露或激活骨形态发生蛋白(bone morphogenetic proteins, BMP)-Smad 轴将上调 SCD1 蛋白表达,促进 *Fabp4*-*Cre*-*Bmp4*<sup>LoxP/LoxP</sup> 小鼠皮下脂肪细胞的脂质动员和全身能量代谢,诱导脂肪细胞褐变<sup>[64]</sup>。其中,通过上调 SCD1 表达而增加的油酸合成将触发脂解和脂质自噬<sup>[64, 65]</sup>。不同来源的脂肪组织可能是 SCD1 缺失时出现代谢差异表型的潜在原因。先前研究发现内脏脂肪质量的增加与机体代谢功能障碍相关,而增加皮下脂肪则可以维护代谢生理稳态<sup>[66]</sup>。而冷暴露干预促使 FFAs 的脂解和转化,也可能导致产生不一致的实验结果<sup>[67]</sup>。此外,在特异性敲除肝脏 SCD1 的小鼠中发现  $\beta_3$ -肾上腺素受体( $\beta_3$ -adrenergic receptor,  $\beta_3$ -AR)的表达增加和 PPAR $\alpha$  的激活,伴随着 BAT 产热和肝脏脂解的显著增加<sup>[68]</sup>。尽管 PPAR $\alpha$  诱导脂肪酸 $\beta$ -氧化和产热基因的转录,但小鼠肝脏中 PPAR $\alpha$  的缺失无法消除肝脏特异性 SCD1 敲除对脂肪酸氧化增强的效果,提示肝脏 SCD1 仍然存在 PPAR $\alpha$  以外的下游靶点<sup>[68]</sup>。

SCD1 调控体内 SFAs 和 MUFAs 水平变化能够作为脂质信号显著影响细胞代谢健康。人类和动物的研究发现,富含 SFAs 的西方饮食将异常提高体内 SFAs 水平,并引发慢性炎症。这进一步干扰代谢稳态,促进各类代谢性疾病的发生和发展<sup>[69-71]</sup>。相反,富含 MUFAs 的地中海饮食存在抗炎效果,可诱导 M2 型巨噬细胞极化和抑制 NOD 样受体热蛋白结构域相关蛋白 3 (NOD-like receptor thermal protein domain associated protein 3, NLRP3)炎症小体激活,以及抑制 SFAs 诱导的胰岛素抵抗和肝脏脂肪变性<sup>[72-74]</sup>。因此,由 SCD1 内源性合成的 MUFAs 在调节组织脂质稳态方面具有关键且独特的功能。

### 2.3 SCD1和炎症

脂肪酸长期以来被认为可调节组织炎症<sup>[75, 76]</sup>。SCD1 通过调节 MUFAs 和 SFAs 间的平衡,对炎症和应激反应存在显著影响<sup>[77]</sup>。SFAs 作为 Toll 样受体(Toll-like receptor, TLRs)家族成员等细胞表面免疫受体的配体,可促进炎症过程<sup>[70]</sup>。SFAs 也可转化为鞘氨醇等中间物质,间接产生促炎效应<sup>[6, 78]</sup>。SFAs 在包括巨噬细胞、内皮细胞、脂肪细胞和  $\beta$  细胞等多种类型细胞中均发挥促炎效应<sup>[6]</sup>。而 MUFAs 通过激

活抗炎机制(如促进 M2 巨噬细胞极化和脂肪细胞 IL-10 分泌)能够逆转 SFAs 对脂肪和肝脏组织的不良影响<sup>[79]</sup>。重要的是,SCD1 活性增强可通过促进巨噬细胞中 MUFAs 积累进而减少由髓样分化因子 88 (myeloid differentiation factor 88, MyD88)介导的炎症反应<sup>[80]</sup>。动物研究表明通过核呼吸因子 2 (nuclear respiratory factor 2, NRF2)-SREBP1-SCD1/2 信号轴控制 MUFAs 代谢能够减轻 MyD88 驱动的 TLRs 诱导的巨噬细胞炎症<sup>[80]</sup>。除此之外,SCD1 也可通过增强 DNA 甲基化来控制炎症基因的表达,从而抑制 *Il4ra*、*Il6st* 和 *Tgfb1* 转录,最终减缓炎症过程<sup>[81]</sup>。相反,当使用化学抑制剂抑制脂肪细胞中 SCD1 活性时,脂肪酸如硬脂酸的水平将上调,这会显著加剧包括 IL-6、单核细胞趋化蛋白-1 (monocyte chemotactic protein-1, MCP-1)和趋化因子配体 5 (regulated on activation, normal T-cell expressed and secreted, RANTES)在内的炎症标志物分泌,最终引发细胞发生炎症应激<sup>[82]</sup>。

总而言之,SCD1 能够减弱上述如 SFAs、MyD88 和 TLRs 等各种炎症因子促炎症作用<sup>[83]</sup>。然而,目前尚不清楚累积的 MUFAs 如何解决 MyD88 驱动的炎症,需待未来进一步研究。

### 2.4 SCD1和自噬

在营养缺乏条件下,自噬在蛋白质、脂滴(lipid droplets, LDs)和糖原的分解代谢中起着关键作用,以维持正常代谢平衡<sup>[84, 85]</sup>。自噬的特征是通过包裹部分细胞质和需要降解的细胞器及蛋白质形成自噬体<sup>[86, 87]</sup>。自噬体与溶酶体融合,分解其内容物,以维持细胞内稳态和器官更新<sup>[86, 87]</sup>。脂质是细胞膜结合的信号分子,通过特异性招募调控膜变形、扩展和囊泡运输的细胞质蛋白效应因子,介导自噬小体组装和形成<sup>[88]</sup>。其中,SCD1 调控哺乳动物细胞和组织中的脂肪酸构成,并显著下调自噬水平<sup>[7]</sup>。在小鼠胚胎成纤维细胞中 SCD1 缺失阻碍了 Akt 磷酸化激活,随后增加叉头盒蛋白 O1 (Forkhead box O1, FOXO1)核转位,最终激活 *Atg4b*、*Atg12* 和 *Becn1* 等自噬相关基因的转录<sup>[89]</sup>。有趣的是,这种在 SCD1 抑制下的自噬诱导并不依赖于经典 mTOR 信号调控<sup>[89]</sup>。此外,去乙酰化酶 3 (Sirtuin 3, SIRT3)和 NADH: 泛醌氧化还原酶 B9 亚基(NADH: ubiquinone oxidoreductase subunit B9, NDUF9)通过调节 SCD1 表达,从而调控各组织自噬、分化和炎症等生物事件<sup>[41, 52, 90, 91]</sup>。

3 SCD1 水平异常诱发代谢性疾病的流行病学证据

日益增加的流行病学研究证据表明, *Scd1* 基因遗传多态性与代谢性疾病风险因素显著相关, 如血脂分布异常、胰岛素抵抗和慢性炎症<sup>[92–96]</sup>。在一项基于 2152 名成年人的队列研究中, *Scd1* 基因中包含 rs1502593 的两种常见连锁遗传型与代谢综合征 (metabolic syndrome, MetS) 患病率增加显著相关<sup>[97]</sup>。SCD1 潜在高概率的错义多态性已被多项研究报道与代谢性疾病发生有关<sup>[93, 94, 97]</sup>。一项关于 *Scd1* 基因多态性与糖尿病潜在机制的体外研究中, 功能获得性 rs2234970 多态性通过增强 mRNA 稳定性, 抑制蛋白降解, 能够提高 SCD1 蛋白表达和酶活性<sup>[98]</sup>。这与随后胰岛素抵抗风险增加显著相关, 表明 rs2234970 多态性的变异将进一步增加糖尿病的患病

风险<sup>[98, 99]</sup>。  
上述流行病学证据表明, 单核苷酸多态性导致 SCD1 的不同突变及其后续功能改变可能对包括肥胖、MetS 和糖尿病在内的多种代谢性疾病的发生存在显著影响<sup>[24, 100]</sup>。

4 SCD1 与代谢性疾病

SCD1 的异常表达和活性改变与肥胖、NAFLD 和糖尿病等多种代谢疾病的风险增加相关<sup>[9–11, 13, 14, 101]</sup>。重要的是, SCD1 可能通过调节脂质产生在各组织器官间实现“对话”(crosstalk), 在各种代谢疾病中扮演调控分子信号网络的核心角色。并且, 其下游脂质产物不仅影响能量代谢, 也可作为信号分子来调节生理和病理功能。目前, 已有众多研究报道 SCD1 相关信号轴(见表 1), 其是影响糖脂代谢、炎症反应和自噬异常的共通枢纽<sup>[49, 57, 102, 103]</sup>

表 1. SCD1 影响不同代谢性疾病的机制  
Table 1. The mechanisms by which SCD1 affects different metabolic diseases

| Diseases | Signaling axis                                                                                                                                                                                                                 | Effects of SCD1 on metabolic diseases                                                                                                                                                                                                                                                                                  |
|----------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Obesity  | NDUFB9 <sup>[91]</sup><br>FOXO1 <sup>[104]</sup><br>AMPK-ACC <sup>[26]</sup><br>NRF2-SREBP 1 <sup>[105]</sup>                                                                                                                  | Excess MUFAs production through SCD1 promotes lipogenesis, leading to hypertrophy and proliferation of adipocytes, which promotes fat accumulation and leads to lipid toxicity.                                                                                                                                        |
| NAFLD    | STAT3 <sup>[106]</sup><br>ADGRF1 <sup>[107]</sup><br>FGF1-SREBF1 <sup>[108]</sup><br>AMPK-SREBP1 <sup>[109]</sup><br>NcDase-Wnt <sup>[27]</sup><br>SIRT3 <sup>[90]</sup>                                                       | Overactivity of SCD1 reduces fatty acid oxidation and increases lipid deposition in the liver, contributing to the development of hepatic steatosis, lipid accumulation, and insulin resistance. Furthermore, SCD1 plays a role in promoting inflammation and oxidative stress by altering the fatty acid composition. |
| Diabetes | PPAR-RXR <sup>[110]</sup><br>AKT-mTOR <sup>[62]</sup><br>AMPK-SIRT1 <sup>[102]</sup><br>SREBP1 <sup>[111]</sup>                                                                                                                | High SCD1 activity promotes elevated triglyceride levels and indirectly affects insulin signaling sensitivity. On the other hand, SCD1 is able to maintain the number and size of lipid droplets in $\beta$ cells and the structure and function of islets.                                                            |
| Cancer   | AKT-GSK3 $\beta$ <sup>[112]</sup><br>mTOR-SREBP1 <sup>[113]</sup><br>METTL16-YTHDC2 <sup>[114]</sup><br>Wnt-IGHM <sup>[115]</sup><br>PI3K-AKT-mTOR <sup>[49]</sup><br>FBW7-NRA41 <sup>[13]</sup><br>XBP1-CHOP <sup>[116]</sup> | SCD1 promotes tumor proliferation, growth, migration, invasion and metastasis by maintaining the lipogenesis of cancer cells.                                                                                                                                                                                          |

NAFLD: non-alcoholic fatty liver disease; INSIG1: insulin-induced gene 1; SREBP1: sterol regulatory element-binding protein 1; AMPK: adenosine 5'-monophosphate (AMP)-activated protein kinase; ACC: acetyl-CoA carboxylase; NRF2: nuclear respiratory factors 2; NcDase: neutral ceramidase; SIRT3: Sirtuin 3; PI3K: phosphoinositide 3-kinase; mTOR: mammalian target of rapamycin; FBW7: F-box and WD repeat domain-containing 7; NRA41: nuclear receptor subfamily 4 group a member 1; CHOP: C/EBP-homologous protein; Bax: BCL2-associated X; NDUFB9: NADH: ubiquinone oxidoreductase subunit B9; STAT3: signal transducer and activator of transcription 3; ADGRF1: adhesion G protein-coupled receptor F1; FGF1: fibroblast growth factor 1; RXR: retinoid X receptor; METTL16: methyltransferase-like protein 16; YTHDC2: YTH domain containing 2; MUFAs: monounsaturated fatty acids.



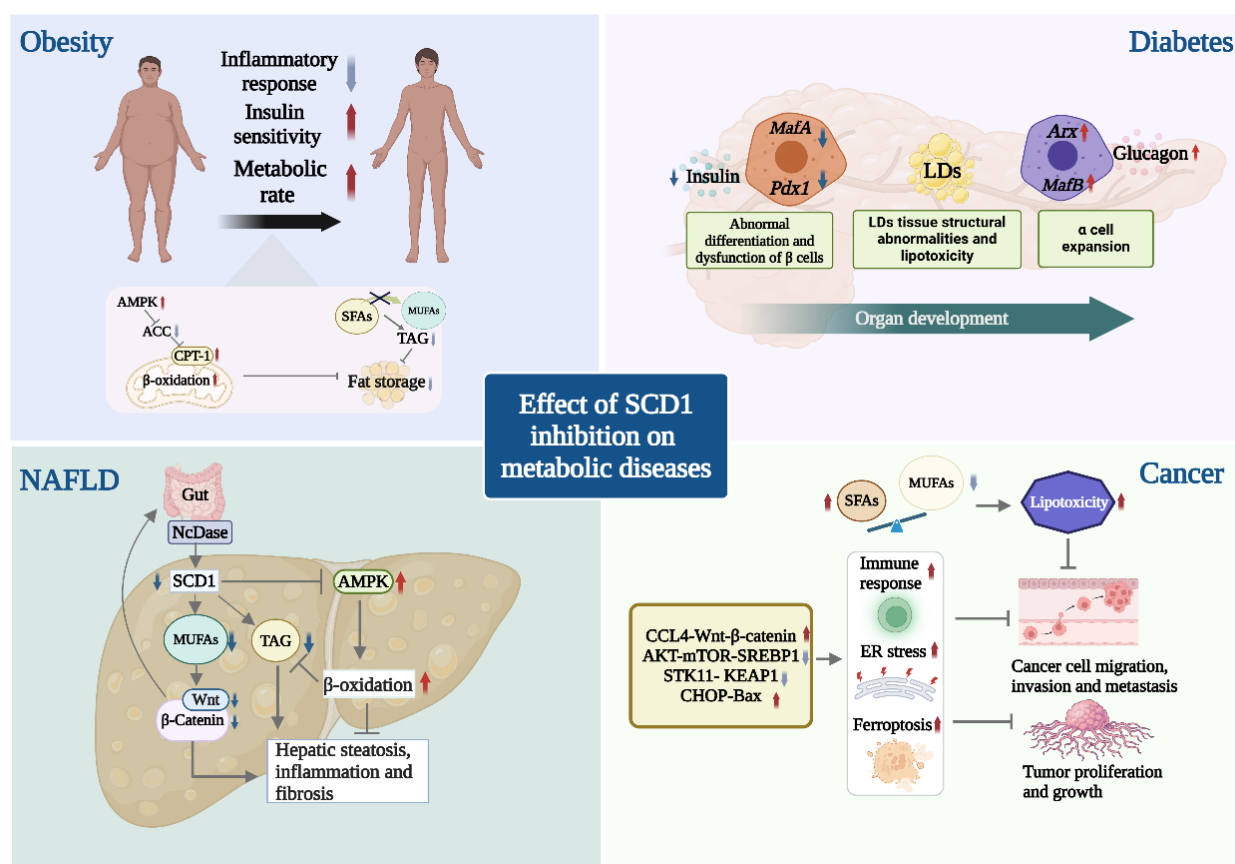


图 3. 抑制SCD1在各类代谢性疾病中的作用

Fig. 3. The impact of SCD1 inhibition on various metabolic diseases. Inhibition of SCD1 improves obesity by activating the AMPK-ACC signaling axis, increasing  $\beta$ -oxidation of fatty acids in mitochondria and reducing lipid synthesis through decreased conversion of MUFAs. This mechanism also enhances hepatic fatty acid oxidation, improves hepatic lipid accumulation, and alleviates hepatic steatosis in NAFLD, while attenuating the fibrosis and inflammation responses driven by the NcDase-Wnt- $\beta$ -Catenin signaling axis in non-alcoholic steatohepatitis (NASH). Furthermore, loss of SCD1 adversely affects islet structure and functionality, reducing the number and size of LDs in  $\beta$ -cells, thus exacerbating susceptibility to lipotoxicity, and accelerating structural and functional changes associated with type 2 diabetes mellitus (T2DM). Eventually, it leads to abnormal differentiation and dysfunction of  $\beta$ -cells. Additionally, inhibiting SCD1 suppresses ER stress, ferroptosis, and immune responses in cancer cells, hindering their proliferation, migration, invasion, and metastasis. TAG, triacylglyceride; LDs, lipid droplets; NcDase, neutral ceramidase; SFAs, saturated fatty acids; ER stress, endoplasmic reticulum stress. The other abbreviaions are the same as those in Table 1.

(图3)。因此, 深入理解SCD1在各种代谢性疾病中的调控机制对于疾病治疗具有深远的临床意义。

#### 4.1 SCD1和肥胖

SCD1是体重调节的关键因子。Cruz-Color等研究者对38名肥胖患者进行脂肪组织活检, 发现SCD1在皮下和内脏脂肪组织高表达, 其中皮下脂肪组织中SCD1的表达随身体质量指数的增加呈显著上升趋势<sup>[117]</sup>。全身SCD1基因敲除小鼠能够抵抗HFD引起的肥胖, 同时胰岛素敏感性和新陈代谢率增强<sup>[118]</sup>。SCD1基因缺失ob/ob小鼠(*Scd1*<sup>-/-</sup> *leptin*<sup>ob/ob</sup>)低代谢表型逆转, 能量消耗显著增强, 肝脏脂质沉

积显著减少<sup>[119]</sup>。然而, 与*leptin*<sup>ob/ob</sup>小鼠相比, *Scd1*<sup>-/-</sup> *leptin*<sup>ob/ob</sup>小鼠表现出胰岛素抵抗、低胰岛素血症和高血糖, 伴随出现异常胰岛功能<sup>[118]</sup>。基于此, SCD1在肥胖进程中可能发挥双重作用: SCD1缺失减轻了肥胖表型, 但伴随胰岛功能障碍和糖代谢紊乱<sup>[118]</sup>。近期研究也发现SCD1缺乏将引起胰岛内 $\beta$ 细胞向 $\alpha$ 细胞分化, 胰岛微结构破坏并出现胰岛功能障碍<sup>[28]</sup>。其中, SCD1参与调节关键 $\beta$ 细胞基因(如*Pdx1*和*MafA*)启动子甲基化状态, 抑制SCD1会导致低甲基化并改变 $\beta$ 细胞染色体内的甲基化水平分布<sup>[28]</sup>。同时, 在INS-1细胞中过表达SCD1将增

加SFAs向MUFAs转化<sup>[104, 120]</sup>。棕榈油酸作为MUFAs之一，能够增强FoxO1转录活性，使INS-1细胞对棕榈酸引起的脂毒性侵害具有抵抗力，从而保护它们免受脂毒性诱发的ER应激和细胞凋亡<sup>[104, 120]</sup>。有趣的是，特异性敲除小鼠肝脏SCD1能够降低肝脏脂肪生成酶活性，减少肝糖异生，从而避免HCD而不是HFD导致的(皮下和内脏)脂肪组织的脂质堆积，改善小鼠肥胖和胰岛素抵抗<sup>[121]</sup>。已知肝脏*Scd1*启动子甲基化水平与全身脂肪含量和血清瘦素水平显著正相关，并受到高胆固醇饮食和HFD的正向调节<sup>[122]</sup>。这提示我们SCD1功能可能具有组织特异性作用，并且在肝-脂肪轴的组织间“对话”中存在关键作用，是肝脏-脂肪轴间交流的介质<sup>[123]</sup>。

过表达小鼠骨骼肌中SCD1显著减弱AMPK磷酸化，AMP/ATP比例降低，小鼠表现出对HFD诱导的脂质积累和胰岛素抵抗的易感性<sup>[26]</sup>。在正常原代人骨骼肌细胞中，过表达SCD1将减少45%油酸氧化，并且细胞内甘油三酯和磷脂水平分别增加22%和27%<sup>[9]</sup>。同时，AMPK和ACC2磷酸化水平受到抑制，导致肌细胞脂肪酸氧化率下降<sup>[9, 124]</sup>。此外，与正常人相比，在重度肥胖患者骨骼肌中，SCD1显著增加与脂肪酸氧化率下降、甘油三酯合成增加以及MUFAs水平上升呈显著正相关，这可能是受表观遗传和/或遗传机制驱动<sup>[9, 31]</sup>。

## 4.2 SCD1、NAFLD和NASH

NAFLD包括其更严重程度NASH全球患病率高达20%~25%，是全球重大公共卫生问题<sup>[125]</sup>。NAFLD主要特征是肝脏脂肪变性、胰岛素抵抗、线粒体功能障碍、炎症和氧化应激<sup>[126, 127]</sup>。SCD1作为临床上肝脏脂代谢稳态的生物标志物，在NAFLD发生和发展中起着重要作用<sup>[123]</sup>。随着脂肪酸在脂肪组织或肝细胞中的过度累积，肥胖人群患NAFLD的风险显著增加<sup>[128]</sup>。能量过剩会启动肝脏AMPK-SREBP1信号轴中SREBP1c的裂解和核转位，随后其与SCD1启动子内SRE基序相结合，引起肝细胞甘油三酯的积累<sup>[109]</sup>。而特异性敲除肝脏SCD1能够减少肝脏脂质合成，保护小鼠免受过量碳水化合物诱发的肝脏脂肪变性<sup>[121]</sup>。

伴随NASH发展，肝脏将发生一系列异常病理变化，尤以肝细胞肿胀、胰岛素抵抗、炎症反应和肝脏纤维化为主要特征<sup>[129, 130]</sup>。此外，与组织再生相关信号通路将异常激活，这会促进贮脂细胞诱导肝脏纤维化<sup>[131]</sup>。动物和临床研究表明，肝脏纤维化

导致肝硬化和肝功能衰竭，并且促进肝癌的发展<sup>[132–134]</sup>。已知中性神经酰胺酶(neutral ceramidase, NcDase)-SCD1- $\beta$ -链蛋白( $\beta$ -Catenin)形成的反馈回路通过调节细胞内鞘氨醇和FFAs水平，促进NASH中的炎症和纤维化<sup>[27]</sup>。在HFD喂养下，NcDase将减少调节肠道微生物群组成的小肠免疫球蛋白A(immunoglobulin A, IgA)的产生，从而降低NASH小鼠肠道中瘤胃菌科的定植<sup>[27]</sup>。瘤胃球菌已知能够抑制SCD1的表达，从而减少小鼠肝脏和回肠中MUFAs生成<sup>[27]</sup>。总体来说，NcDase缺失通过改变肠道微生物群落抑制SCD1的表达，进而产生增强Wnt活性和 $\beta$ -Catenin核转运的SFAs。过量SFAs引起Wnt脂酰化和 $\beta$ -Catenin激活，诱发HFD下肝脏炎症反应<sup>[27, 135]</sup>。并且，这将激活HFD喂养的小鼠肝脏中纤维化基因(如*Il-6h*和*ACTA2*)表达<sup>[27]</sup>。重要的是，Wnt蛋白正向促进NcDase活性，这将最终形成NcDase-SCD1-Wnt之间的正反馈循环，加剧NAFLD病程<sup>[27]</sup>。在另一项关于肠道微生物刺激肝脏再生的研究中，由肠道微生物产生的短链脂肪酸(short-chain fatty acids, SCFAs)通过门静脉运输到肝脏，并在那里转化为SFAs。SFAs随后通过促进肝细胞中SCD1表达来激活肝细胞增殖<sup>[136]</sup>。值得注意的是，SCD1的表达被发现与肝切除术后人肝脏的增殖和再生呈正相关<sup>[136]</sup>。类似地，通过siRNA抑制SCD1显著降低了HepG2细胞的增殖以及肝细胞类器官中的细胞周期标志物<sup>[136]</sup>。基于此，通过改变特定肠道微生物特征和肝脏SCD1表达，诱导NASH中肝部分再生并改善肝功能似乎是一个值得探索的治疗途径<sup>[136]</sup>。然而，应当注意的是，SCD1表达和酶活性的增加可能会导致肝细胞脂质氧化减弱和炎症反应上调，这会间接加剧NASH的进程<sup>[19, 109]</sup>。因此，未来需要进一步探索肠道微生物与其宿主之间这种复杂相互作用背后的分子机制。此外，SCD1的激活在多大程度上能够促进肝细胞的增殖和再生，而不加剧NASH或肝细胞癌发展，值得进一步研究。

## 4.3 SCD1和糖尿病

研究发现，SCD1的活性增强与T2DM的发生和发展呈正相关<sup>[137, 138]</sup>。一项近3000人的大型社区前瞻性队列研究发现，SCD1活性与糖尿病的发病率呈显著正相关<sup>[139]</sup>。然而，最近的证据发现，SCD1表达增加也可能帮助人类抵抗T2DM。已知脂毒性影响胰岛 $\beta$ 细胞的DNA甲基化，导致 $\beta$ 细胞



功能障碍以及加剧 T2DM<sup>[102]</sup>。因此, 肥胖糖尿病患者中 SCD1 蛋白表达或活性的提高可能是机体减轻脂毒性的一种代偿方式<sup>[140]</sup>。SCD1 的激活可以加速酰基辅酶 A 去饱和, 恢复细胞内脂肪酸水平正常化, 这能够缓解肥胖个体因过量 SFAs 引起的脂毒性<sup>[141]</sup>。并且, 长链 SFAs 对人类 EndoC- $\beta$ H1  $\beta$  细胞具有显著脂毒性, 而 SCD1 在这些细胞中的过表达可以逆转 SFAs 对  $\beta$  细胞结构和功能的负面影响, 如 ER 应激和 LDs 形成减少<sup>[142, 143]</sup>。然而, 当下调 SCD1 时, AMPK 被激活, 进而上调 NAD-依赖性去乙酰化酶 Sirtuin-1 (SIRT1) 的表达<sup>[102]</sup>。SIRT1 导致 DNA 甲基转移酶 1 (DNA methyltransferase 1, DNMT1) 去乙酰化, 引起  $\beta$  细胞 DNA 全局甲基化水平下降, 最终影响  $\beta$  细胞 LDs 形态结构以及对脂毒性的抵抗力<sup>[102]</sup>。SCD1 活性也影响胰岛  $\beta$  细胞的自噬小体-溶酶体融合阶段<sup>[144]</sup>。抑制 SCD1 活性将改变 INS-1E 细胞膜磷脂组成, 引发异常自噬小体-溶酶体融合和自噬清除缺陷<sup>[144]</sup>。这些效应加剧了 INS-1E 细胞对棕榈酸诱导损伤的易感性<sup>[144]</sup>。同时, 这种变化干扰了 ER 的正常功能, 将引发线粒体介导的细胞凋亡, 抑制  $\beta$  细胞增殖及其功能, 最终导致胰岛  $\beta$  细胞死亡<sup>[144]</sup>。

此外, SCD1 还作为胰腺-肝脏轴中介“对话”因子, 调节肝脏代谢灵活性。例如, 研究发现胰腺分泌的胰岛素通过肝脏 SCD1 产生棕榈油酸, 能够激活 Wnt- $\beta$ -Catenin 通路活性以调节肝脏代谢稳态以及胰岛素敏感性<sup>[145]</sup>。

综上所述, SCD1 在肥胖和糖尿病中发挥不同作用的潜在解释可能是: 当体内出现代谢应激或胰岛素敏感性受损时, SCD1 会被诱导表达。一旦应激恢复正常, 作为保护者的 SCD1 水平会相应地下调。另一方面, 全身性 SCD1 抑制对肥胖的积极影响可能以  $\beta$  细胞的脂毒性为代价<sup>[28, 118]</sup>。全身 *Scd1* 基因敲除导致小鼠胰岛中 LDs 的大小和数量减少, 中性脂质积累降低, 这会减弱脂质储存能力, 并降低小鼠胰岛  $\beta$  细胞对脂毒性的抵抗力<sup>[146]</sup>。总之, SCD1 在肥胖和糖尿病中的作用是错综复杂的。实验模型的差异(人类或啮齿动物样本)、组织类型(脂肪组织或胰岛)以及遗传特征和代谢功能障碍(肥胖或 T2DM)都可能导致上述不一致的研究结果。

#### 4.4 SCD1 和癌症

SCD1 通过调节肿瘤细胞脂肪代谢, 在肿瘤发生、发展、侵袭和转移中发挥着关键作用<sup>[8]</sup>。临床

上已发现在卵巢癌、胃癌、结直肠癌和胰腺癌中 SCD1 表达水平均显著升高<sup>[10, 147, 148]</sup>。在肺癌和乳腺癌中, SCD1 高水平表达与不良预后显著相关<sup>[149]</sup>。其中, SCD1 的上调通过增加 MUFAs 合成, 保护癌细胞免受 SFAs 脂毒性的侵害, 促进癌细胞迁移和肿瘤再生长<sup>[49]</sup>。研究证实, 经油酸处理后癌细胞可逆转由 SCD1 活性抑制引起的癌细胞凋亡和生长抑制<sup>[112, 150]</sup>。除此之外, SCD1 还参与调控肿瘤细胞的压力和生存过程, 如铁死亡。铁死亡是一种铁依赖性、由细胞膜上脂质过氧化物过量累积引发的程序性细胞死亡形式<sup>[151]</sup>。值得注意的是, 脂代谢是影响铁死亡的关键调节通路之一<sup>[152]</sup>。作为调控脂质代谢的关键介质, SCD1 在癌细胞抵御铁死亡中起关键作用<sup>[8]</sup>。已知 SCD1 通过 PI3K-Akt-mTOR 途径赋予癌细胞抵抗活性氧(reactive oxygen species, ROS)诱导的铁死亡, 从而促进癌症复发<sup>[49]</sup>。最近的研究发现, 人类结肠癌中 LINC01606 (一种致癌基因)显著上调, 通过与 miR-423-5p 相互作用来增强 SCD1 的表达, 激活经典 Wnt/ $\beta$ -Catenin 信号通路, 促进癌细胞生长和干细胞化, 并抑制铁死亡<sup>[115]</sup>。与此同时, Wnt/ $\beta$ -Catenin 信号转导又反过来通过转录因子结合 IGHM 增强子 3 (transcription factor binding to IGHM enhancer 3, TFE3) 增加 LINC01606 的转录, 形成正反馈调节回路, 进一步抑制铁死亡并增强癌细胞干性(癌症干细胞的自我更新和分化的能力)<sup>[115]</sup>。

未折叠蛋白反应(unfolded protein response, UPR)是一种生存机制, 由 ER 中未折叠或错误折叠蛋白质的积累而触发。然而, 异常水平的 UPR 与癌症的发生和发展显著相关<sup>[153]</sup>。SCD1 可能也参与调节应激反应途径, 这些途径促进肿瘤生长和转移<sup>[154]</sup>。先前的研究表明, 抑制肿瘤 SCD1 导致糖原合酶激酶  $\beta$  (glycogen synthase kinase 3 $\beta$ , GSK3 $\beta$ ) 去磷酸化活化, GSK3 $\beta$  是 Akt 通路的下游靶点, 通过诱导  $\beta$ -Catenin 和细胞周期蛋白 D1 降解来阻止癌细胞增殖<sup>[112]</sup>。类似地, 在人类癌细胞中, 抑制 SCD1 表达会触发 UPR 介导的 Xbp1 mRNA 剪接、eIF2 $\alpha$  磷酸化以及 C/EBP 同源蛋白(C/EBP homologous protein, CHOP)表达增加等事件, 最终诱导 CHOP 依赖的癌细胞凋亡<sup>[116]</sup>。最近, Ben-David 等研究者通过高通量筛选鉴定出小分子 PluriSIns, 其能够抑制人类多能干细胞(human pluripotent stem cells, hPSCs)中 SCD1 活性。PluriSIns 处理后导致棕榈酸积累和油酸转化减少, 引起 hPSCs 的 ER 应激, 促

进癌细胞凋亡<sup>[155]</sup>。总体而言, SCD1 的异常激活能够促进肿瘤生长、转移、免疫逃逸以及癌症发展中对治疗的抵抗<sup>[156]</sup>。

## 5 靶向SCD1的疾病治疗新策略

随着对 SCD1 研究的不断深入, 靶向 SCD1 治疗代谢性疾病已受到广泛关注。不同研究相继揭示了 SCD1 活性抑制在 NAFLD、T2DM 及癌症等代谢性疾病治疗中的显著疗效<sup>[8, 16, 123]</sup>。通过合成具有极低半抑制浓度( $IC_{50}$ , 即物质抑制特定生物或生化功能的效力的量度, 数值越低, 药物性能越好)的合成类化合物抑制 SCD1 能够有效改善代谢性疾病。例如, 在 NASH 患者二期临床试验中发现口服 Aramchol 能够改善患者肝脏纤维化、增强肝功能以及血糖控制<sup>[18]</sup>。重要的是, Aramchol 表现出了良好的耐受性和安全性<sup>[18, 19]</sup>。此外, 患者在口服 Aramchol 后糖化血红蛋白(HbA1c)水平得到显著改善, 这表明了 Aramchol 在治疗 T2DM 中的潜在重要临床价值<sup>[18, 157]</sup>。与此同时, 天然化合物如小檗碱和姜黄素等传统中草药也被证明能够有效调节 SCD1 活性, 展现出独特的临床治疗潜力<sup>[109, 158, 159]</sup>。但值得注意的是, 鉴于 SCD1 广泛表达于多种组织和器官中, 全身抑制 SCD1 可能引起系列不良效应(如扰乱细胞内的脂质稳态), 从而损害正常机体生理功能。因此, 需更加注重 SCD1 抑制剂长期使用的安全性、有效性和剂量优化性, 为代谢性疾病治疗提供有力且稳定的治疗。未来或可进一步开发针对特定组织 SCD1 活性抑制策略的设计, 如研发针对肝脏、脂肪组织和骨骼肌的纳米颗粒和脂质纳米颗粒等组织特异性纳米级药物递送方式(一种基于生物纳米结构的给药方式, 主要存在基于脂质或聚合物或白蛋白的三类纳米药物, 已用于癌症和基因治疗)(见图 4), 从而精确靶向特定组织, 最大限度提高药物疗效并减少对其他组织产生潜在副作用<sup>[160–162]</sup>。

## 6 总结与展望

SCD1 作为一个代谢枢纽因子, 其表达受到转录和表观遗传的调控, 通过控制 MUFA 转化来协调糖脂代谢、炎症和自噬等生物过程。在生理病理条件下, 抑制 SCD1 能够促进脂肪组织棕色化, 减少脂质积累以及增强脂解, 在代谢和能量稳态中发挥关键作用。此前已有多种 SCD1 抑制剂在临床前模型测试中显示出良好的抗肿瘤效果。例如,

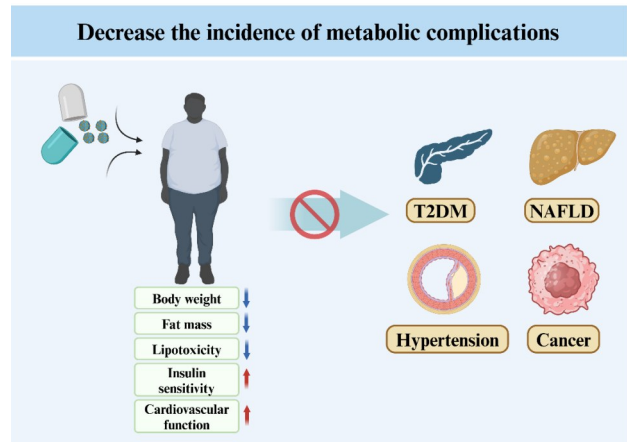


图 4. 特异性精准抑制 SCD1 改善多种代谢性疾病病理状态  
Fig. 4. Precise amelioration of various metabolic disease pathologies through specific SCD1 inhibition. As a key factor in metabolic diseases and their complications, SCD1 is expected to become an important target for the treatment of obesity, diabetes and cardiovascular diseases by developing tissue-specific SCD1 delivery drugs to safely and effectively improve the excessive accumulation of lipids, insulin resistance, inflammation and cardiovascular risk in patients with metabolic diseases. NAFLD: non-alcoholic fatty liver disease; T2DM: type 2 diabetes mellitus.

A939572、MF-438 和 CAY10566 等 SCD1 抑制剂在多种癌症类型(如胰腺癌、食管癌、乳腺癌和肝癌等)中显著降低了肿瘤细胞存活率, 抑制了肿瘤生长<sup>[163–165]</sup>。然而, 值得注意的是, 当前 SCD1 的临床转化十分有限, SCD1 抑制剂的临床应用仍然存在诸多挑战: 1) 现有 SCD1 抑制剂存在一定的脱靶效应, 存在药物相关毒性问题。已知全身 SCD1 抑制或缺乏会对各种器官, 特别是皮肤和眼睛产生不良影响(皮肤功能下降和视力衰退)<sup>[166]</sup>; 2) 对组织器官内特定靶标信号与 SCD1 的相互作用尚不清楚。研究发现皮下脂肪组织中 SCD1 特异性缺失引起葡萄糖转运蛋白 1 (glucose transporter 1, GLUT1) 上调, 但不清楚 SCD1 如何调控 GLUT1 蛋白表达<sup>[167]</sup>; 3) 由于不同癌症类型和亚型对 SCD1 的依赖程度不同, 目前缺乏合适的个体化治疗策略来治疗代谢性疾病患者群体。STK11/KEAP1 共突变的肺腺癌对 SCD1 抑制剂更为敏感, 这表明在这类癌症患者中, SCD1 抑制剂可能更为有效<sup>[168]</sup>。

因此, 为了解决这些问题, 研究者们首先需开发特异性更高、靶点更准确的 SCD1 抑制剂, 通过跨学科合作揭示更多结构信息和潜在的抑制机制,

为合理设计下一代 SCD1 抑制剂提供理论依据。其次,临床上也可根据生物学背景(如组织类型和疾病特征)联合信息与计算科学(如计算机模拟和基于 AI 的医疗数据建模)技术,尝试 SCD1 抑制剂和其他药剂(如自噬抑制剂)的联合治疗,设计有效的个性化治疗策略。已有部分研究显示,SCD1 与铁死亡抑制剂联合使用较单一药剂治疗在肠癌中表现出更好的疗效<sup>[169]</sup>。此外,通过脂质组学鉴定 SCD1 衍生的新型下游脂质介体将有助于阐明 SCD1 复杂的代谢和信号网络。类似于胰高血糖素样肽-1 受体,这些脂质介体的功能操纵能够为开发针对 SCD1 的小分子药物提供新的洞见,它们能够对抗代谢性疾病并改善多个组织和器官的系统代谢。

总之,SCD1 作为代谢枢纽基因,其抑制剂的开发用于治疗各种代谢性疾病具有可观前景,特别是对于肥胖、糖尿病、NAFLD 和 NASH 等代谢性疾病。

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