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Therapeutic implications of bioactive sphingolipids: A focus on colorectal cancer

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ABSTRACT

Therapy of colorectal cancer (CRC), especially a subset known as locally advanced rectal cancer, is challenged by progression and recurrence. Sphingolipids, a lipid subtype with vital roles in cellular function, play an important role in CRC and impact on therapeutic outcomes. In this review we discuss how dietary sphingolipids or the gut microbiome via alterations in sphingolipids influence CRC carcinogenesis. In addition, we discuss the expression of sphingolipid enzymes in the gastro-intestinal tract, their alterations in CRC, and the implications for therapy responsiveness. Lastly, we highlight some novel therapeutics that target sphingolipid signaling and have potential applications in the treatment of CRC. Understanding how sphingolipid metabolism impacts cell death susceptibility and drug resistance will be critical toward improving therapeutic outcomes.

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Treatment challenges in colorectal cancer

As of 2016, colorectal cancer (CRC) remains the third most frequently diagnosed cancer with an estimated 140,000 new cases per year.¹ Of concern are the findings of a recent study by the American Cancer Society that the incidence of CRC is rapidly increasing among younger adults.² The incidence of rectal cancer, a subset of CRC located in the pelvis within 12 cm from the anus, has doubled between 1989-90 to 2012-13 in adults under 55 year old.² Locally advanced rectal cancer (LARC), either a muscle-invasive primary tumor (AJCC Stage 2) or involved lymph nodes (AJCC Stage 3), is particularly challenging to treat and has high rates of disease recurrence. Neoadjuvant 5-fluorouracil (5FU) chemoradiation (5FU/RT) is the most effective treatment approach for patients with LARC. The landmark randomized German Rectal Cancer Trial (CAO/ARO/AIO-94) showed that LARC patients treated with neoadjuvant 5FU/RT had fewer treatment-associated complications and less local disease recurrence compared with patients receiving surgery followed by 5FU/RT.³ Tumor 5FU/RT pathologic response (AJCC tumor regression grading (TRG)) has since become an established surrogate marker of long-term survival and a useful oncologic benchmark.^{4,5} Approximately 20% of the LARC patients have a pathologic complete response (TRG0) to 5FU/RT, with exceptional long-term outcomes.5,6 Unfortunately in ~80% of cases, therapeutic resistance is evident and contributes to surgical failure, disease recurrence, and ultimately, death of the patient. Further, while additional agents have shown promise in cancer cell lines, therapies combining these agents with 5FU/RT have significantly increased toxicity in patients without improvement in clinical response.⁷⁻¹⁰ To change the therapeutic paradigm, LARC inter-patient heterogeneity must be integrated into clinical algorithms tailoring therapy for individual patients by either identifying more effective strategies or by omitting ineffective treatments to avoid unnecessary toxicity. Improving therapeutic response rates to preoperative therapy should ultimately translate into better outcomes associated with CRC. Given the high rate of resistance, highlighted by the lack of complete response in the majority of rectal cancer patients, exploring novel molecular strategies to enhance conventional therapy for CRC is desperately needed.

Molecular mechanisms of therapeutic resistance in CRC continue to be under intense investigation. Recently, investigations have demonstrated that bioactive sphingolipids play an important role in CRC and impact oncologic therapies such as chemotherapy and radiation. Therefore, dissecting the relationship between bioactive sphingolipids and chemoradiation resistance should provide insight into tumor survival mechanisms and suggest potential novel targets to improve CRC treatment strategies.

Sphingolipid metabolism

Sphingolipids represent a lipid subtype with vital roles in cellular function.¹¹ As structural components, sphingolipids influence the physical properties of membranes, which impacts on cellular signaling.¹² In addition, sphingolipids affect cellular signaling by acting as secondary messengers. An increasing body of literature supports the role of sphingolipids in regulating biologically critical processes such as cell growth and death, autophagy, migration and invasion, and angiogenesis.^{13,14}

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Central to sphingolipid metabolism is ceramide, which can be generated de novo, through hydrolysis of complex sphingolipids, or via the salvage pathway (Fig. 1). De novo synthesis of ceramide occurs in the endoplasmic reticulum, where serine palmitoyl transferase (SPT) mediates the condensation of serine and palmitoyl-CoA to generate 3-ketosphinganine, which is then reduced to dihydrosphingosine. Next, ceramide synthases (CerS) add a fatty acid moiety generating dihydroceramide and lastly dihydroceramide desaturase (DEGS) introduces a double bond to generate ceramide. Ceramide serves as a building block for complex sphingolipids such as sphingomyelin and glycosphingolipids. Hydrolysis of these complex sphingolipids is the second metabolic pathway by which ceramide can be generated. Ceramide can also be hydrolyzed by ceramidases into sphingosine. In the salvage pathway, ceramide synthases utilize sphingosine and fatty acids as substrates to regenerate ceramide. Thus ceramide can be generated through 3 metabolic pathways, an apparent redundancy that may be necessary since lipid molecules do not freely diffuse throughout the cytoplasm. Distribution of sphingolipid enzymes across various subcellular compartments can thus fulfill the need for site-specific generation of ceramide. Ceramide can also be shuttled between compartments by ceramide transporters, such as CERT.

In contrast to ceramide, sphingosine and sphingosine-1phosphate are soluble mediators. Sphingosine is generated by hydrolysis of ceramide by ceramidases and serves as a substrate for sphingosine kinases, which through the addition of a phospho-group generate sphingosine-1-phosphate (S1P). The phospho-group of S1P can be removed by S1P phosphatase or S1P lyase, which results in regeneration of sphingosine or irreversible degradation into ethanolamine-1-P and palmitaldehyde, respectively. The latter step constitutes the exit from sphingolipid metabolism. As shown in Fig 1, the majority of enzymatic reactions in the sphingolipid pathway are reversible, which endows cells with flexibility in responding to cellular stimuli. Consequently, cellular stimuli can induce a flux in sphingolipids that involve rapid conversion between ceramide, complex sphingolipids, sphingosine and S1P.

Additional complexity derives from multiple isoforms of sphingolipid enzymes that can vary in subcellular location and pH requirements. For example, there are at least 3 ceramidases and sphingomyelinases with different pH requirements and 2 sphingosine kinases with different subcellular locations. Furthermore, the 6 ceramide synthases (CerS1-CerS6) have preferential substrate utilization and can generate ceramides that vary in acyl chain length ranging from 14–30 (or more)



Figure 1. Sphingolipid metabolism. Central to sphingolipid metabolism is ceramide, which can be generated *de novo* and is used as a building block for complex sphingolipids. It can also be phosphorylated or further metabolized into sphingosine-1-phosphate. See text for more details. Abbreviations: SPT = serine palmityoltransferase; CerS = ceramide synthase; DEGS = dihydroceramide desaturase; SMS = sphingomyelin synthase; SMase = sphingomyelinase; GCS = glucosylceramide synthase; CDase = ceramidase; CK = ceramide kinase.

carbons.15 CerS1 preferentially generates ceramide with 18-carbon fatty acids (C18-Cer) while CerS5 or CerS6 primarily generate ceramide with 16-carbon fatty acids (C₁₆-Cer). Phenotypes observed in CerS-deficient mice suggest that ceramides with different fatty acid chain lengths have distinct biologic roles. For example, CerS3 or CerS4-deficient mice develop problems with skin barrier function and alopecia, respectively,^{16,17} while CerS1- and CerS2-deficient mice exhibit abnormalities in the central nervous system.¹⁸ In addition, CerS2-deficiency (inability to generate C24-Cer) results in a compensatory generation of C₁₆-Cer, which in the liver leads to apoptosis and subsequent development of hepatocellular cancer.^{19,20} Similarly, shifting ceramide composition in cancer cell lines through targeting specific CerS changes cellular signaling and responses.²¹ The tissue distribution of CerS varies and likely reflects the need of specific ceramide species for proper signaling and sphingolipid homeostasis in any given tissue. Several comprehensive reviews on CerS are available.^{18,21,22}

The two isoforms of sphingosine kinase, SK1 and SK2, both utilize sphingosine and generate S1P, but have significant differences in subcellular localization and function.²³ Upon activation, SK1 migrates to the plasma membrane where it releases S1P extracellularly. Extracellular S1P exert its actions in an autocrine or paracrine fashion by binding to 5 different S1P-receptors that via G-proteins regulate a broad range of cellular functions. In contrast to SK1, SK2 appears to have both proand anti-apoptotic functions depending on cell type, stimuli, and subcellular localization.²³ Targeting SK2 in malignancies reduces tumor growth and suggests an anti-apoptotic role for this enzyme in cancer.²³

Ceramide has been associated with cell death, growth inhibition and differentiation whereas S1P regulates proliferation, motility, angiogenesis and inflammation. Sphingolipid homeostasis is critical for normal cellular function, which it evident from rare mutations that result in lipid storage diseases as well as phenotypes of various sphingolipid gene knockout mice. Alterations in bioactive sphingolipids have been associated with a variety of human diseases, including cancer.

The role of dietary sphingolipids in CRC carcinogenesis

The per capita consumption in the United States from sources rich in sphingolipids such as dairy products or soy is estimated at 0.3-0.4 g/day.²⁴ Considering this significant intake of dietary sphingolipids, the impact of dietary sphingolipids on intestinal carcinogenesis has been investigated in different animal models. Models of intestinal carcinogenesis include treatment with chemical carcinogens such as 1,2-dimethylhydrazine (DMH) or azoxymethane (AOM) and APC^{min} mice, which serve as genetic model for inactivating mutations of the Adenomatous Polyposis Coli (APC) gene. Inactivation of APC, a negative regulator of the Wnt signaling pathway is the most common gene abnormality in CRC and mutations are found in 81% of the CRC cases in the Cancer Genome Atlas. Mutation of APC results in accumulation of β -catenin, which acts as a transcriptional co-activator of TCF/LEF gene family and promotion of CRC.²⁵ APC^{min} mice, which express a truncation mutation of the APC gene, develop up to 100 polyps in the small intestine as well as colon tumors. Multiple studies have confirmed that

dietary sphingolipids, including sphingodienes, glucosylceramide, and sphingomyelin reduce carcinogenesis in both APC^{min} mice as well as in DMH treated mice.²⁶⁻²⁹ Adding soy glucosylceramide to the diet of DMH treated mice reduced colonic cell proliferation by about 50% and significantly diminished the number of aberrant colonic crypt foci.²⁷ Similarly, sphingomyelin supplementation reduced the number of aberrant crypt foci by 70% and diminished the number of colonic adenocarcinomas.³⁰ These studies suggested a role for intestinal metabolism of dietary sphingolipids as a critical step in protection from colon carcinogenesis and supported the hypothesis that elucidating signaling events may provide a foundation for possible therapeutic approaches in CRC.

Dietary sphingolipids are metabolized throughout the intestinal tract.³¹ The metabolism of dietary sphingomyelin, which is found in dairy products,³⁰ is initiated by the release of cholecystokinin from intestinal endocrine cells, which stimulates the release of alkaline sphingomyelinase from the gallbladder and the secretion of trypsin from the pancreas. Trypsin cleaves alkaline sphingomyelinase releasing it from the intestinal mucosa and enhancing enzymatic activity in the lumen.³² Ceramide generated by the activity of alkaline sphingomyelinase is then further hydrolyzed by neutral ceramidase into sphingosine, which is absorbed into enterocytes, where it can be converted into S1P. In the DMH model, dietary sphingomyelin resulted in increased mRNA, protein expression and activity of alkaline sphingomyelinase,³³ which presumably increases the generation of ceramide, although this was not directly evaluated in the study. In the APC^{min} model, 3 types of sphingolipid-supplemented diets (ceramide, milk sphingolipids similar to proportions found in dairy products, and a mixture of 60% complex sphingolipids and 40% ceramide) were compared with the control diet. While all sphingolipid-supplemented diets significantly reduced the tumor burden per animal, the mixture of complex sphingolipid and ceramide was the most effective and resulted in the redistribution of β -catenin from a diffuse pattern into localization to intercellular junctions between intestinal epithelial cells.²⁹ Moreover, when CRC cell lines that harbor APC mutations (SW480 and T84) were treated with sphingosine or ceramide, cytosolic and nuclear β -catenin levels were reduced, which ultimately lead to cell death.²⁹ In the APC^{min} model, sphingodienes reduce Wnt signaling via a protein phosphatase 2A/Akt/GSK3β-dependent mechanism.²⁶ Glucosylceramide, which reduced tumor burden in the APC^{min} and DMH models, significantly alters the expression of 96 genes, including decreased expression of the TCF/LEF family member TCF4.²⁷ Taken together, these studies suggest that dietary sphingolipids may inhibit intestinal carcinogenesis through negative regulation of Wnt pathway signaling. The protective effects of sphingolipids in the setting of APC mutations provide supportive evidence of targeting sphingolipid metabolism for therapeutic intervention.

The role of the gut microbiome in CRC

Symbiosis between the host and the gut microbiome is critical for intestinal health. Metagenomic studies show that patients with CRC have altered gut microbiomes when compared with healthy controls and some studies have linked specific microbes to increased tumorigenesis.^{34,35} While sphingolipids have been primarily studied in eukaryotes, microbes such as the soil dwelling Sphingomonas as well as human commensals including Bacteroides, Porphyromonas, and Prevetella from the Bacteroidetes family contain sphingolipids.^{36,37} Bacteroides fragilis, a prevalent gram-negative microbe within the intestine, encounters a significant amount of stress within the constantly changing host environment. B. fragilis sphingolipids do not seem to play a role in growth of the microbe but rather are essential for survival under stressful conditions.³⁸ Interestingly, germ-free mice or mice colonized with B. fragilis lacking the bacterial equivalent of SPT were found to have an increase in invariant natural killer T cells (iNKT), which play a key role in innate and adaptive immunity.³⁹ This increase in iNKT cells was unique to the gastro-intestinal niche and facilitated an increased inflammatory response following induction of colitis.³⁹ B. fragilis sphingolipids were found to inhibit the proliferation of iNKT cells during neonatal development, which suggested a potential role of these lipids in the maintenance of gut homeostasis.³⁹ A proof-of principle experiment demonstrated that oral administration of bacterial glycosphingolipids to young mice resulted in reduced iNKT cells and protection from colitis into adulthood.³⁹

A subset of B. fragilis known as enterotoxigenic B. fragilis (ETBF) is distinguished by its ability to secrete a pathogenic enterotoxin. ETBF are increased in patients with inflammatory bowel disease and CRC.^{40,41} Recently it was shown that ETBF induce the release of intestinal mouse derived exosome-like nanoparticles (IDEN) that contain S1P and mediate tumor promotion through induction of pro-inflammatory Th17 cells.⁴² Comparing the effect of secreted particles from nontoxigenic (NTBF) and ETBF, researchers found that NTBF particles induced immunosuppressive Treg cells (CD4⁺FoxP3⁺) and exerted a protective effect in the DSS-induced colitis model, whereas particles from ETBF induced inflammatory Th17 cells.⁴² The effect of ETBF was recapitulated with NTBF engineered to express the enterotoxin. Further studies led to a model in which particles from ETBF induced intestinal epithelial cells to secrete S1P-containing IDENs. These IDENs then had a 2-fold function in promoting the inflammatory response: (i) IDENs recruit inflammatory T cells (CD4⁺IL17A⁺CCR6⁺) from the periphery into the intestinal tissue and (ii) IDENs were taken up by intestinal macrophages resulting in enhanced production of prostaglandin E2, which promotes the proliferation of the recruited pro-inflammatory CD4⁺IL17A⁺CCR6⁺ T cells.⁴² Since the risk of CRC in patients with inflammatory bowel disease is increased by approximately 2-3-fold,⁴³ these studies suggest that toxin-induced S1P release may contribute to colon tumorigenesis.

Certain probiotic bacteria can also influence host sphingolipids and/or their enzymes. *Lactobacillus rhamnosus* GG reduces levels of lysophosphatidylcholines, sphingomyelins, and glycerophosphatidylcholines, whereas *Lactobacillus brevis* and *Streptococcus thermophilus* increase neutral sphingomyelinase levels, which may be involved in ceramide induced immune cell apoptosis.^{44,45} Taken together these studies show that microbes can either directly, through their own sphingolipids, or indirectly, through modulation of host sphingolipids, impact intestinal inflammation, which has been associated with CRC carcinogenesis. These findings suggest novel opportunities to utilize microbes or nano-particle mediated delivery as preventative or possibly as therapeutic strategies in CRC.

Expression of sphingolipid enzymes in the gastro-intestinal tract and alterations in CRC

Ceramide synthases

Ceramide synthases (CerS1-CerS6) vary in mRNA distribution across tissues and preferentially generate ceramides with specific chain lengths. The first mammalian ceramide synthase was discovered in 2002 and subsequent overexpression and knockout experiments have indicating that altering the composition of ceramide species influences cell physiology and pathology.^{21,46} CerS-deficient mice are viable and do not have any obvious defects in the gastrointestinal tract, suggesting that CerS family members have the ability to functionally compensate under normal physiologic conditions.^{21,46}

CerS mRNA expression varies among tissues but is not necessarily reflective of protein expression or activity. The biscistronic mRNA for CerS1, a CerS that preferentially generates C₁₈-Cer, is detected in the small and large intestine.⁴⁷ CerS1 protein however was only detected in CRC cell lines and not in colonic biopsies,⁴⁷ which is consistent with data in the protein atlas http://www.proteinatlas.org/ENSG00000223802-CERS1/ tissue. CerS2, which preferentially generates very long chain ceramides such as C24-ceramide, is ubiquitously expressed and has been suggested to function as a housekeeping gene.^{48,49} CerS3 appears to primarily play a role in the skin and testis and although its mRNA has been detected in the gastrointestinal tract, it is unclear whether the protein is expressed. CerS4 has been implicated in folate stress, yet RNAi targeting of this protein did not protect cells (including HCT116 colon cancer cells) from methotrexate toxicity nor did it noticeably affect proliferation.⁵⁰ Similar to CerS2, CerS5 and CerS6, which both preferentially generate C₁₆-Cer, are ubiquitously expressed.⁵¹ CerS5 and CerS6 appear to be involved in cell stress responses.²¹ CerS6 mRNA is highly expressed in the intestinal tract and the C₁₆-Cer content in CerS6-deficient mice is reduced to 25%.^{22,52} Reduced CerS6 expression has been associated with epithelial to mesenchymal transition in a panel of cancer cells that included several CRC cell lines.53

Sphingomyelinases

In healthy human colon tissue sphingomyelinase activity is detected at acidic, neutral, and alkaline pH, indicating that 3 sphingomyelinases (SMase) known as acid SMase (aSMase), neutral SMase (nSMase) and alkSMase, are expressed. AlkSMase plays an important role in digestion with expression highest in the ascending colon and lowest in the rectum.⁵⁴ A similar gradient was found for nSMase but not for aSMase.⁵⁴ In comparison to surrounding normal tissue, SMase activity in colorectal cancer is reduced by 75%, 50%, and 30% for alkSMase, nSMase and aSMase, respectively.⁵⁴ These results are consistent with immunohistochemical staining for alkSMase described in the protein atlas, although results are considered uncertain as they are based on only one antibody (http://www.proteinatlas.org/ENSG00000182156-ENPP7/tissue#gene_infor mation).

Ceramidases

Five different ceramidase genes encode enzymes that hydrolyze ceramide at different pH optimums. Little is known about alkaline ceramidases but interestingly ACER3-deficiency has recently been associated with colitis-induced colon cancer in mice due to an increase in $C_{18:1}$ -Cer and increased expression of proinflammatory cytokines.⁵⁵ Acid (ASAH1) and neutral ceramidase (ASAH2) have been more extensively studied and both have been identified as potential therapeutic targets in malignancies.

ASAH1 is primarily found in the lysosomal compartment and mutations in its gene result in rare genetic disorders as well as other diseases.^{56,57} Knockout of ASAH1 results in death early during embryonic development, indicating that other ceramidases are unable to compensate for its enzymatic activity.⁵⁸ Although ASAH1 expression is essential for development, expression in the normal colon is relatively low.⁵⁹ Ceramide has been shown to transcriptionally induce ASAH1 expression either in response to radiation or as a consequence of CerS6 (and possibly CerS3–5) overexpression.^{60,61} Thus it is possible that expression of ASAH1 increases in response to cell stresses that elevate intracellular ceramide within the intestinal system.

ASAH2 is the enzyme that hydrolyses ceramide generated by alkSMase during digestion of dietary sphingolipids. ASAH2deficient mice develop normally but tissues have elevated ceramide and reduced levels of sphingosine.⁶² RNAi targeting of ASAH2 in HT29 and HCT116 colon cancer cell lines increases ceramide, leading to 50% loss in viability *in vitro* and delayed xenograft growth *in vivo*.⁶³ In the AOM carcinogenesis model, ASAH2-deficient mice developed significantly fewer tumors with a 93% reduction in adenocarcinomas and an 82% decrease in total colon tumors compared with wild type mice.⁶³

Sphingosine kinases

Both sphingosine kinases play important roles in CRC and are potential therapeutic targets. The SK1/S1P pathway was found to positively regulate cyclooxygenase-2 (COX-2), an enzyme involved in inflammatory pathways that is overexpressed in several epithelial cancers. SK1 expression is elevated in the majority of human CRC specimen (78-89%).^{64,65} A similar increase in SK1 expression was detected in the AOM carcinogenesis model.⁶⁴ Elevated expression of SK1 also increases the susceptibility to colitis-associated cancer.⁶⁶ Conversely, SK1deficient mice had significantly less aberrant crypt foci formation and reduced colon cancer development.⁶⁴ Recently, it was shown that targeting SK1 with RNAi in CRC cell lines SW480 and HCT116 increases E-cadherin expression, decreases vimentin expression and reduces viability and migration.⁶⁵ These observations suggest a role of SK1 in epithelial to mesenchymal transition, proliferation, and migration in the absence of inflammatory signaling.

SK2 also increases in CRC (51/64; 80%), although 21/64 apparently normal tissues also stained positive for SK2.⁶⁷ In serum-free cultures of HCT116 CRC cells, SK2 has been suggested to function as a survival factor.⁶⁸ Recently targeting of SK2 by RNAi was shown to decrease expression of cMyc in the CRC cell line LoVo, which is consistent with cMyc as a down-stream target of SK2.^{67,69} cMyc plays an important role as a

target gene of TCF-4 during aberrant signaling of mutated APC,⁷⁰ which highlights SK2 as a potential novel target in CRC.

Sphingolipids and radiation sensitivity

With the central role of sphingolipids in mediating apoptosis, a growing body of evidence implicates sphingolipids as critical mediators of anti-neoplastic treatment approaches such as chemotherapy and radiation. Given the importance of radiation in the therapeutic armamentarium for LARC, understanding the close association between sphingolipid metabolism and radiation may open opportunities to define novel therapeutic strategies for the future.

Early studies supported a direct link between radiation sensitivity and sphingolipids by demonstrating that radiation activates aSMase to generate ceramide with resultant formation of ceramide-rich domains in the plasma and mitochondrial membranes.⁷¹⁻⁷³ Investigations definitively demonstrated that endothelial derived aSMase was a central mediator of the radiation efficacy highlighting the importance of the tumor stromal compartment in mediating the ceramide response.⁷³ An elegant study using a model of radiation-induced apoptosis in *Caenorhabditis elegans* germ cells confirmed a central role of ceramide in the radiation response.⁷⁴ Investigators demonstrated that loss-of-function mutations in CerS genes, *hyl-1* and *lagr-1*, completely inhibited the apoptotic response that could be rescued with exogenous delivery of C₁₆-Cer.⁷⁴

Recent investigations have demonstrated that radiation efficacy is driven in part by the resulting accumulation of ceramides to enhance apoptosis mediated by multiple pathways including protein phosphatases PP1 and PP2A, cathepsin D and protein kinase $C\zeta$.⁷⁵ Studies in cancer cell lines demonstrated the complex regulation of radiation-induced ceramide generation and apoptosis.⁷⁶ In contrast to CerS2 overexpression that resulted in partial protection from apoptosis, CerS5 and 6 enhanced radiation-induced apoptosis and increased C_{16} -Cer.⁷⁶ Further, many agents and stimuli that lead to cell death require CerS, particularly CerS6, activity for effect, suggesting a critical role in radiation response.⁷⁷⁻⁸⁴

The generation of ceramide following therapy induces apoptosis but can also transcriptionally activate ASAH1. In prostate cancer cells, ceramide generated following radiation therapy transcriptionally activates ASAH1 expression via an AP-1 mediated mechanism.⁶¹ The increase in ASAH1 resulted in elevation of S1P and therapy resistance and was also associated with relapse following radiation therapy.⁶¹ These results are consistent with several other studies that have associated elevated expression of ASAH1 with proliferation and drug resistance.⁸⁵⁻⁸⁷ In colon cancer cells, overexpression of CerS6, which shifts the cellular ceramide profile toward C₁₆-Cer, also results in transcriptional activation of ASAH1.⁶⁰ However, in contrast to prostate cancer cells, the increase in ASAH1 as a consequence of CerS6 overexpression resulted in increased susceptibility to 5FU chemotherapy, despite transcriptional activation of ASAH1.⁶⁰ Interestingly, ASAH1 expression has been positively correlated with a better prognosis in luminal subtypes of breast cancer and epithelial ovarian cancer.^{88,89} Thus elevated ASAH1 may not be exclusively associated with resistance. The key difference between the prostate and colon cancer



Figure 2. *Cell death susceptibility*. Cells with increased expression of acid ceramidase (ASAH1) may remain susceptible to death stimuli, if ceramide synthases activity prevails over activity of sphingosine kinases.

models appears to be the metabolic flux of the ASAH1 product sphingosine. While S1P levels increased in prostate cancer cells, analysis of metabolic flux in HT29 cells overexpressing CerS6 yielded minimal generation of S1P compared with HT-GFP control cells.^{60,61} This suggests that in cells with high levels of CerS6, the ASAH1 product sphingosine may be preferentially used in the salvage pathway to regenerate ceramide whereas in cells where ASAH1 is associated with resistance, sphingosine maybe preferentially used by SK to generate S1P. The influence sphingolipid metabolic flux may have on cell death susceptibility is illustrated in Fig. 2.

Sphingolipid metabolism as a therapeutic target

As ceramide has emerged as an important mediator of cell death responses, therapies to directly target sphingolipid metabolism have been developed. Strategies include antibodies that bind S1P, inhibitors of enzymes involved in driving sphingolipid metabolism toward S1P, and the use of ceramide or ceramide analogs.⁹⁰⁻⁹³ The S1P antibody Sphingomab slowed tumor growth in a preclinical study of renal cell cancer and appeared to exert this effect by targeting the vasculature, since blood flow to tumors was reduced when measured by MRI.⁹⁴ The humanized S1P antibody ASONEP was well tolerated but overall progression-free survival in patients with advanced renal cell carcinoma was less than 2 months (http://www.prnewswire.com/news-releases/lpath-reports-results-for-aso nep-phase-2a-study-in-renal-cell-carcinoma-300053872.html).

Inhibitors that interfere with generation of S1P include inhibitors of ceramidases and sphingosine kinases. Ceramidase inhibitors have been developed for ASAH1 and ASAH2.^{63,95} The ASAH1 inhibitor LCL-521 has shown promising results in head and neck squamous carcinoma and prostate cancer models^{61,96} but has not been evaluated in CRC. The ASAH2 inhibitor C₆-urea ceramide preferentially reduced viability in a panel of human colon cancer cell lines compared with non-malignant rat intestinal epithelial cells.⁶³ C₆-urea ceramide did not significantly impact on levels of sphingosine or S1P but increased ceramide, which enhanced apoptosis and increased autophagic flux.⁶³ Neither LCL-521 nor C₆-urea ceramide have yet been evaluated in clinical trials.

SK are also a key targets in cancer therapy and both SK1 and SK2 inhibitors have been developed.⁹⁷⁻⁹⁹ Of particular interest to CRC is the SK2 inhibitor ABC294640, which protects from



Figure 3. Sphingolipid therapeutics under clinical investigation. ASONEP is an antibody designed to target S1P. ABC294640 is an inhibitor of sphingosine kinase 2. Ceramide nanoliposomes are pegylated liposomes containing C_6 -ceramide.

colitis-driven CRC in the AOM/DSS model.¹⁰⁰ ABC294640 also enhances the efficacy of 5FU and significantly reduced the growth of HT29 xenografts.¹⁰¹ The Phase I trial for ABC294640 was successfully completed¹⁰² and additional clinical trials in hematological malignancies (NCT02229981, NCT02757326) and hepatocellular carcinoma (NCT02939807) have been initiated. As an inhibitor of SK2, treatment with ABC294640 decreases cMyc, which is a target gene of TCF-4 during aberrant signaling of mutated APC, and may therefore be a promising agent in the treatment of CRC.^{69,70,103-105}

Cationic ceramides, developed based on structure-function and targeting properties, induce anti-proliferative responses, including autophagic cell death, apoptosis and growth inhibition.^{93,106,107} Among the cationic ceramides, LCL-30 has been investigated in CRC.^{108,109} Due to the cationic charge LCL-30 preferentially localizes to the mitochondria and triggers the apoptotic pathway.¹⁰⁹ Treatment of mice bearing CT26 tumors with LCL-30 mono-therapy resulted in a highly significant reduction in tumor burden.¹⁰⁸ The anti-tumor efficacy of LCL-30 was greater than doxorubicin although combination of LCL-30 and doxorubicin did not result in a benefit over LCL-30 alone.¹⁰⁸

Another approach to utilize ceramides as therapeutics involves the encapsulation within non-toxic, non-aggregating, nanoscale (<100 nm) vehicles. Nanotechnology has the potential to improve the pharmacokinetics, biodistribution, and toxicological profiles of exogenously administered bioactive ceramide. In addition, drug encapsulation within nanotechnology can address inherent issues of poor solubility, precipitation, non-tumor cell delivery (liver cells or macrophages) and immunogenicity of bioactive lipids. Ceramide has been encapsulated within nanoscale liposomes, calcium phosphosilicate nanocolloids, nanoemulsions, polyethyleneoxide-modified polyepsiloncaprolactone nanoparticles (PEO-PCL) and linear-dendritic nanoparticles.^{110,111} One of these preclinical approaches, the C₆-ceramide nanoliposome (CNL) has just entered the clinic for solid tumors under FDA IND 109471 (NCT 02834611). CNL is a homogeneous 85nM, -8mV, nanotechnology that intercalates 30 molar percent C₆-ceramide within a 12 molar percent pegylated liposome.¹¹² Compared to positively charged ceramide analogs or vehicles, these anionic particles intravenously deliver therapeutic doses of ceramide to tumors, without inducing liver or renal toxicities in rodent and canine models. Biological efficacy has been observed in models of CRC, hepatocellular carcinoma, head and neck squamous cell carcinoma,

melanoma, breast cancer, pancreatic cancer as well as LGL, BCLL and AML leukemias.¹¹² Alluding to this efficacy, CNL is engineered to exploit the enhanced permeability and retention (EPR) properties of tumors, due to a calculated biologic halflife of between 18-21 hours compared with just minutes for free ceramide. Recently, CNL were shown to inhibit STAT3 signaling in chronic lymphocytic leukemia (Doshi, et al. under review). The STAT3 pathway is constitutively activated in colon-cancer initiating cells (ALDH⁺CD133⁺ subpopulation) and targeting STAT3 through genetic (shRNA) or pharmacological (LLL12) approaches significantly inhibited tumor growth.¹¹³ Thus incorporating CNL into therapeutic regimen in notoriously difficult to treat LARC could be a powerful approach. Moreover, second generation CNL or other nanoscale approaches are already being developed that can be actively targeted to cancer cells through bio-conjugation of small molecules, antibody fragments or aptamers or that can co-deliver synergistic drugs within the ceramide nanoscale delivery platform.

Conclusions

In over a decade, since the landmark randomized German Rectal Cancer Trial, neoadjuvant chemoradiation has remained the gold standard approach for LARC. As our understanding and appreciation of the complex sphingolipid regulatory pathways grow in the context of cancer biology, we have learned that the imbalance of bioactive sphingolipids can have a major impact the therapeutic efficacy of chemotherapy and radiation. In the future, harnessing these critical metabolic pathways through novel therapies such as ABC294640 or CNL, which are currently under evaluation in clinical trials involving cancer patients, holds significant potential to replace or synergize with traditional therapies like radiation and to improve care for patients with this devastating malignancy.

Disclosure of potential conflicts of interest

Penn State Research Foundation has licensed ceramide nanoliposomes to Keystone Nano, Inc. (University Park, PA). MK is co-founder and CSO of Keystone Nano.

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