

Molecular mechanisms of polyamine analogs in cancer cells

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The natural polyamines are aliphatic cations with multiple functions and are essential for cell growth. Soon after the critical requirement of polyamines for cell proliferation was recognized, the metabolism of polyamines was pursued as a target for antineoplastic therapy. Initially, much attention was focused on the development of inhibitors of polyamine biosynthesis as a means to inhibit tumor growth. The best-characterized inhibitor is α -difluoromethylornithine (DFMO), an irreversible inhibitor of ornithine decarboxylase. While compensatory mechanisms in polyamine metabolism reduce the effectiveness of DFMO as a single chemotherapeutic agent, it is currently undergoing extensive testing and clinical trials for chemoprevention and other diseases. There has been increasing interest over the last two decades in the cytotoxic response to agents that target the regulation of polyamine metabolism rather than directly inhibiting the metabolic enzymes in tumor cells. This interest resulted in the development of a number of polyamine analogs that exhibit effective cytotoxicity against tumor growth in preclinical models. The analogs enter cells through a selective polyamine transport system and can be either polyamine antimetabolites that deplete the intracellular polyamines or polyamine mimetics that displace the natural polyamines from binding sites, but do not substitute in terms of growth-promoting function. Synthesis of the first generation of symmetrically substituted bis(alkyl)polyamine analogs in the mid-1980s was based on the theory that polyamines may utilize

feedback mechanisms to auto-regulate their synthesis. In the 1990s, unsymmetrically substituted bis(alkyl) polyamine analogs were developed. These compounds display structure-dependent and cell type-specific cellular effects and regulation on polyamine metabolism. More recently, a novel class of analogs has been synthesized, which include conformationally restricted, cyclic and long-chain oligoamine analogs. The development and use of these analogs have provided valuable information for understanding the molecular mechanisms of targeting the polyamine pathway as a means of cancer therapy. *Anti-Cancer Drugs* 16:229–241 © 2005 Lippincott Williams & Wilkins.

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Introduction

Polyamines (putrescine, spermidine and spermine) are naturally occurring, polycationic alkylamines that are absolutely required for cell growth. Most living organisms are able to synthesize intracellular polyamines from the precursor amino acids. The metabolic enzymes controlling polyamine concentrations are highly regulated and respond to changing environmental conditions [1–3]. The significance of polyamines for cell growth and function is reflected by the complex coordination of polyamine transport, uptake, synthesis and catalysis in mammalian cells. It has been realized for more than three decades that there is a strong connection between high levels of intracellular polyamines and increased cell proliferation. Since depletion of natural polyamine pools in cancer cells may result in the disruption of several important cellular functions and cytotoxicity, the polyamine metabolic pathway represents an important target for cancer therapy.

Initially, inhibitors of essentially all of the polyamine biosynthetic enzymes were developed and their anti-neoplastic activities were evaluated. Although these inhibitors may interrupt the polyamine metabolic pathway and lead to the cessation of tumor cell growth in *in vitro* and animal studies, the encouraging preclinical results did not translate well into the clinic [4]. Currently, the most actively pursued approach is the development and use of polyamine analogs as tumor growth inhibitors based upon the rationale that natural polyamines have several feedback mechanisms to regulate their own synthesis by reducing the rate of polyamine transport and increasing the rate of polyamine degradation and export [2,5]. Some polyamine analogs can mimic these regulatory properties of the natural polyamines and induce the catabolic process, inhibit polyamine biosynthesis, deplete the polyamine pools, and ultimately lead to the cessation of tumor cell growth [2,3,6]. In this review, we discuss the complexity of

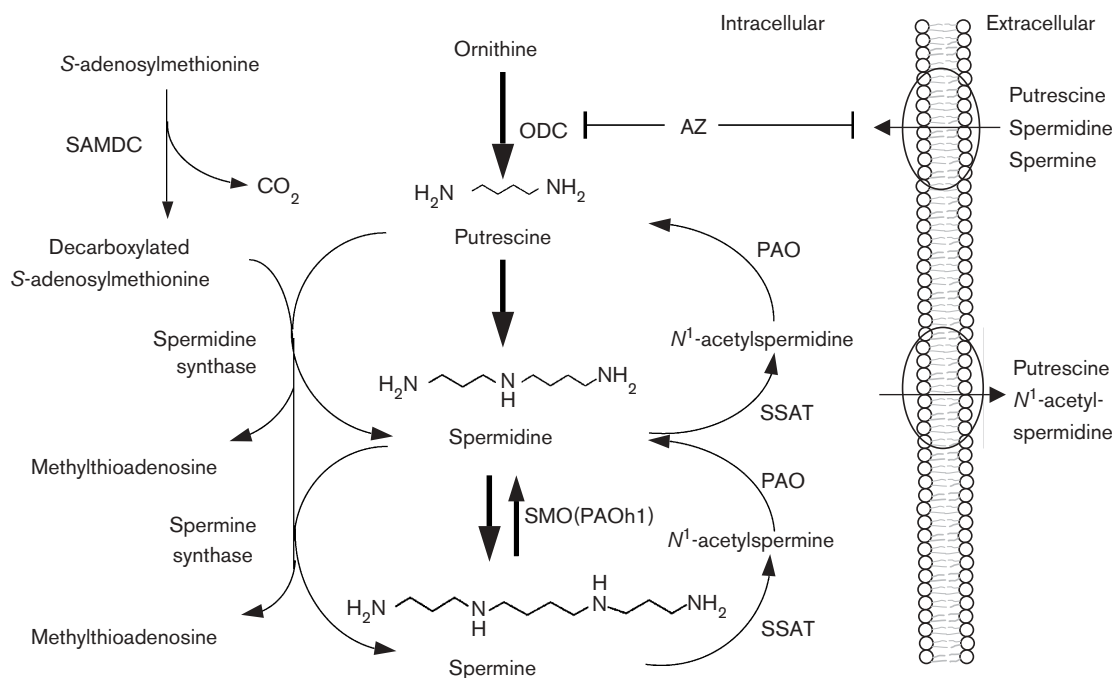
polyamine metabolism, the roles of polyamines in cell growth and tumorigenesis, and the mechanisms of action of different classes of polyamine analogs in cancer cells.

Polyamine function, metabolism and transport

The natural polyamines, putrescine, spermidine and spermine, are found in all eukaryotic cells and are essential for cell growth. Positively charged polyamines can interact electrostatically with negatively charged moieties such as DNA, RNA, proteins and phospholipids to stabilize and alter the structure and function of these molecules [2]. Polyamine-mediated DNA conformational changes play an important role in DNA structural stabilization and may be essential for the normal transcriptional activity of genes [7]. Functional interactions between polyamines and cellular anions may extend to DNA-protein interaction, post-translational modification, protein-protein interaction and enzyme activity [8]. Polyamines have been demonstrated to regulate membrane K^+ channel activity that is important in maintaining normal membrane electrical activity [9]. Polyamines may also function as second messengers in protein phosphorylation and signal transduction by promoting the activity of several important kinases in cell proliferation like MAP kinase and casein kinase II [10,11]. Other important cellular roles of polyamines include anti-oxidant, anti-apoptotic and metabolic regulatory functions [12].

The natural polyamines are synthesized in a highly regulated biochemical pathway [1–3]. As shown in Figure 1, polyamine biosynthesis is rate-limited by the polyamine biosynthetic enzymes, ornithine decarboxylase (ODC) and *S*-adenosylmethionine decarboxylase (SAMDC) (see the end of the article for a full list of abbreviations). The amino acid, ornithine, is converted to putrescine in a decarboxylation reaction catalyzed by ODC. SAMDC forms decarboxylated *S*-adenosylmethionine (dcAdoMet) that serves as an aminopropyl donor in the synthesis of spermidine and spermine by separable aminopropyl transferases. The regulatory protein ODC-antizyme (AZ) can bind to ODC and facilitate the degradation of AZ-ODC complex by the 26S proteasome in a non-ubiquitin-dependent manner [13]. AZ mRNA contains two overlapping open reading frames (1 and 2) and is regulated in a post-transcriptional manner. The presence of increased intracellular polyamine or analog levels can lead to enhanced efficiency of AZ mRNA frameshifting from reading frame 1 which contains an initiation codon to reading frame 2 where most of the functionally active AZ protein is produced [14]. Four forms of AZ (AZ1–4) have been identified, but only AZ1 has strong association with degradation of ODC [14]. The aminopropyltransferase reactions that form natural spermidine and spermine are essentially irreversible, but the catabolism of spermine to putrescine can be accomplished by the activity of two catabolic pathways. The first pathway relies on the acetylation of spermine and

Fig. 1



Polyamine metabolic pathways. See 'Abbreviations' for definitions.

spermidine by the rate-limiting enzyme spermidine/spermine N^1 -acetyltransferase (SSAT) to form N^1 -acetyl-spermine and N^1 -acetylspermidine, respectively. Acetyl derivatives are then cleaved into 3-acetamidopropanal, H_2O_2 , spermidine and putrescine through the action of FAD-dependent polyamine oxidase (PAO). Wang *et al.* and Vujcic *et al.* recently identified a second polyamine catabolic pathway by cloning a variably spliced human spermine oxidase SMO(PAOh1). SMO(PAOh1) uses unacetylated spermine as substrate and is inducible by specific polyamine analogs [15,16]. These findings indicate a complex polyamine catabolic pathway that may contribute to polyamine homeostasis and response to polyamine analogs.

Eukaryotic polyamine uptake is known to be an energy-dependent and carrier-mediated process, and is critical in maintaining cellular polyamine homeostasis. Although much is recognized about the polyamine transport system in prokaryotes, considerably less is understood about the cell membrane polyamine transport system in eukaryotic cells. Soulet *et al.* recently proposed a putative two-step process of eukaryotic polyamine transport where polyamines first enter the cells by a plasma membrane carrier and are then sequestered into pre-existing polyamine-sequestering vesicles via a mechanism that requires an outwardly directed H^+ gradient [17]. It is important to note that the feedback mechanism may regulate the polyamine transport system to ensure intracellular polyamine homeostasis. One example is that AZ induction has recently been found to negatively regulate the eukaryotic polyamine transport system [18].

Polyamines and cancer

Polyamine concentrations and biosynthetic enzyme activities are generally higher in tumor cells as compared to their normal counterparts [19,20]. The key polyamine synthesis enzyme, ODC, is considered to be a proto-oncogene product, and a marker of carcinogenesis and tumor progression. Enhanced levels of ODC activity compared with normal tissues have been detected in several solid tumors [21,22]. Overexpression of ODC in NIH 3T3 cells led to the transformation of these cells [23]. Transgenic mice overexpressing the ODC gene developed skin cancer with typical administration of carcinogen alone, while normal mice required the administration of carcinogen and a tumor promoter [24]. In tissue specimens of prostate cancer patients, ODC expression and activity were substantially higher in cancerous tissues as compared with the normal surrounding tissues [25]. Colorectal mucosa specimens from familial adenomatous polyposis (FAP) patients show higher ODC activity and polyamine levels than those from genotypic normal relatives [26]. In estrogen receptor (ER)-positive breast cancer cells, estradiol up-regulates ODC and increases polyamine levels, which

promotes the breast cell proliferation [27]. Antiestrogen tamoxifen-induced growth inhibition of breast cancer cells is associated with the down-regulation of ODC activity and polyamine biosynthesis [28]. Moreover, treatment with ODC or SAMDC inhibitors decreased intracellular polyamine pools and inhibited the growth of a variety of tumor cells [29,30]. In mice bearing human breast cancer MDA-MB-435 xenografts, DFMO almost completely prevented the development of pulmonary metastasis, which supports a role of ODC and polyamines in promoting the distant metastasis of breast cancer [31].

The above studies demonstrate that polyamines are important for tumor development. It is noteworthy that a high level of polyamines has also been found in diseases other than malignant conditions, such as cystic fibrosis, muscular dystrophy, psoriasis and diabetes [32–35]. Elevated polyamines have also been detected after stress including metabolic, mechanical, chemical and ischemia injury [36,37]. These clinical findings suggest that the application of polyamine measurements in body fluids as a diagnostic tool for cancer patients will be limited. Nevertheless, the clear association between elevated polyamine content and tumorigenesis implies that depletion of polyamines could be a practical strategy for antineoplastic therapy, and measuring polyamine levels could be a useful tool in predicting therapeutic efficacy.

Development of polyamine analogs as effective antineoplastic agents

Polyamine biosynthesis inhibitors

Because of the requirement for polyamines in mammalian cell growth and the demonstration of dysregulated polyamine metabolism in tumor cells, polyamine metabolism became a logical target for cancer therapy. Initially, inhibitors of essentially all of the biosynthetic enzymes were developed and their antineoplastic activities were evaluated [2]. However, most work focused on the rate-limiting polyamine biosynthesis enzymes, ODC and SAMDC, as targets. The best characterized among the inhibitors is α -difluoromethylornithine (DFMO), an enzyme-activated, irreversible inhibitor of ODC [38]. Despite success in *in vitro* and preclinical animal models, clinical trials with DFMO were disappointing and the compound failed to demonstrate lasting antineoplastic effect [4]. The major problems of DFMO as a monotherapeutic strategy include poor transport, complex regulatory mechanisms leading to the compensatory changes in metabolism, rapid turnover of ODC and toxicity (thrombocytopenia, anemia, gastrointestinal reaction and ototoxicity) at high therapeutic doses [29]. Currently, DFMO is the front-line agent in the treatment of African trypanosomiasis and is undergoing clinical trials as a potential chemopreventive drug [29,39]. Other representative biosynthesis inhibitors include the ODC inhibitors α -monofluoromethylornithine (MFMO) and

Table 1 Inhibitors of polyamine biosynthesis enzymes

Ornithine decarboxylase	α -difluoromethylornithine (Efornithine, Ornidyl) (DFMO) α -monofluoromethylornithine (MFMO) (2 <i>R</i> ,5 <i>R</i>)- δ -methylacetylenicputrescine (MAP or MDL 72,175)
S-Adenosylmethionine decarboxylase	methylglyoxal bis(guanyldihydrazone) (MGBG) 4-amidinoinidan-a-one 2'-amidinothiohydrazone (CGP-48664 or SAM486A) S-(5'-deoxy-5'-adenosyl)-methylthioethylhydroxylamine (AMA) 5'-{[(Z)-4-amino-2-butenyl] methylamino}-5'-deoxy-adenosine (AbeAdo) [2,2-bipyridine]-6,6'-dicarboximidamide (CGP-39937)
Spermidine synthase	S-adenosyl-1,12-diamino-3-thio-9-azadodecane (AdoDATAD)
Spermine synthase	S-adenosyl-1,8-diamino-3-thiooctane (AdoDATO)

(2*R*,5*R*)- δ -methylacetylenicputrescine (MAP or MDL 72,175), the SAMDC inhibitors methylglyoxal bis(guanyldihydrazone) (MGBG), 4-amidinoinidan-a-one 2'-amidinothiohydrazone (CGP-48664 or SAM486A), S-(5'-deoxy-5'-adenosyl)-methylthioethylhydroxylamine (AMA), 5'-{[(Z)-4-amino-2-butenyl]methylamino}-5'-deoxyadenosine (AbeAdo) and [2,2-bipyridine]-6,6'-dicarboximidamide (CGP-39937), the spermidine synthase inhibitor S-adenosyl-1,12-diamino-3-thio-9-azadodecane (AdoDATAD) and the spermine synthase inhibitor S-adenosyl-1,8-diamino-3-thiooctane (AdoDATO) (Table 1). Unfortunately, these compounds were also of limited clinical usefulness [2,40].

Symmetrically alkylated polyamine analogs

Because of the limited success of polyamine biosynthetic inhibitors as effective chemotherapeutic agents, interest has been turned to the development of polyamine analogs based upon the theory that natural polyamines have several feedback mechanisms to regulate their own synthesis. Symmetrically substituted bis(alkyl)polyamine analogs represented the first generation of these analogs. Examples includes *N,N'*-bis(ethyl)polyamines, which include *N*¹,*N*¹¹-bis(ethyl)norspermine (BENSpm or BE-3-3-3), *N*¹,*N*¹²-bis(ethyl)spermine (BESpm, or BE-3-4-3), *N*¹,*N*¹⁴-bis(ethyl) homospermine (BEHSpM or BE-4-4-4), *N*¹,*N*¹⁵-bis-[3-(ethylamino)-propyl]-1-17-heptane diamine (BE-3-7-3), *N*¹,*N*¹⁵-bis(ethylamino)-4,8,12-triazapentadecane (BE-3-3-3-3) and *N*¹,*N*¹⁹-bis(ethylamino)-5,10,15-triazanonadecane (BE-4-4-4-4) (Fig. 2) [41]. These analogs utilize the energy-dependent polyamine transport system and are readily accumulated by tumor cells. Their accumulation generally results in a net reduction of the natural polyamine pool, decrease in ODC and SAMDC activities, as well as cytotoxicity in certain types of tumor cells [6,42]. In specific instances in human non-small cell lung cancer (NSCLC), melanoma and breast cancer cells, these analogs result in tremendous induction of SSAT by as much as several thousand fold ('superinduction') [43–45]. The induction of SSAT enzymatic activity is a result of analog-induced

transcription, increased message stability, enhanced translational efficacy and protein stability [46–49]. Studies using melanoma and pancreatic adenocarcinoma cells suggested that cytotoxicity produced by symmetrical bis(ethyl)polyamines analogs is correlated with the induction of SSAT activity in these cells [44,50]. Perhaps the most successful alkylpolyamine to date is BENSpm, an analog that shows exceptional promise as an antitumor agent in both *in vitro* and *in vivo* studies. However, a recent phase II clinical trial of BENSpm in breast cancer patients showed that this analog can be safely administered to women with advanced refractory breast cancer, but was not effective as a single agent [51]. Combination of BENSpm with standard cytotoxic agents is currently being examined [52].

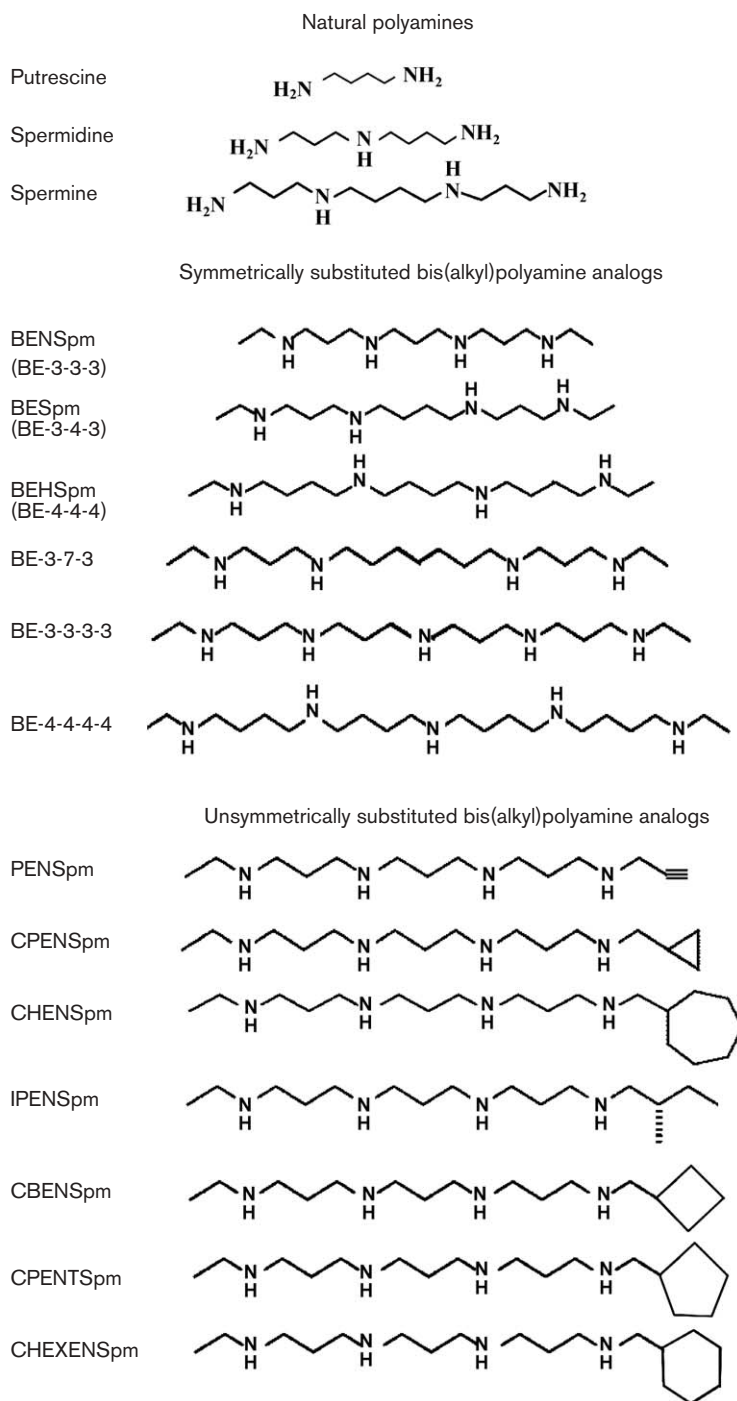
Unsymmetrically substituted polyamine analogs

In the 1990s, a second generation of polyamine analogs termed unsymmetrically substituted bis(alkyl)polyamines was developed by Woster *et al.* at Wayne State University [53]. The first synthesized unsymmetrically substituted analogs were *N*¹-propargyl-*N*¹¹-ethyl-norspermine (PENSpM) and *N*¹-(cyclopropylmethyl)-*N*¹¹-ethyl-4,8-diazaundecane (CPENSpM) (Fig. 2). Both compounds exhibited cytotoxicity and induction of SSAT similar to or greater than symmetrically substituted analogs like BESpm [53]. As with symmetrically substituted analog-treated tumor cells, the cytotoxicity of PENSpM and CPENSpM was correlated with the induction of SSAT mRNA levels and activity [53]. Two other representatives of this class of compounds, *N*¹-(cycloheptylmethyl)-*N*¹¹-ethyl-4,8-diazaundecane (CHENSpM) and (*S*)-*N*¹-(2-methyl-1-butyl)-*N*¹¹-ethyl-4,8-diazaundecane (IPENSpM), exhibited a more potent growth-inhibitory effect against *in vitro* tumor cell growth, but with only modest SSAT induction and polyamine depletion [54]. In addition, CHENSpM and IPENSpM, but not CPENSpM, could alter microtubule polymerization and induce G₂/M cell cycle arrest in tumor cells [55]. These data imply that small modifications in backbone structure of unsymmetrically substituted polyamine analogs have profound consequences on cellular effects and cytotoxicity produced by these compounds.

Conformationally restricted, cyclic and long-chain oligoamine analogs

Recently, a third generation of conformationally restricted analogs has been developed [56–58]. These agents incorporate alterations into the free rotation of the single bonds in otherwise flexible molecules such as spermine or the analogs, thus restricting the molecular conformation that they may assume. Such modifications can alter the ability of analogs to bind DNA, tRNA or other polyamine-binding sites by introducing bends, kinks or loops at their binding domains [56,59]. One class of conformationally restricted analogs comprises the tetramines (homospermine analogs) (Fig. 3). In these compounds, the external

Fig. 2



Chemical structures of natural polyamines, and symmetrically and unsymmetrically substituted bis(alkyl)polyamine analogs.

aminopropyl residues are replaced by aminobutyl residues, making a homospermine backbone. By introduction of alicyclic residues or one or two *cis* double bonds in the homospermine backbone, free rotation at the central part

of the molecule is restricted [58,60]. Another class of conformational restricted analogs is termed pentamine analogs. Their free rotating conformation is also restricted by the introduction of *cis* or *trans* double bonds into the

hydrocarbon skeleton or hydrophilic groups into the pentamine structure [58,60] (Fig. 3). Oligoamine analogs have longer chains than natural polyamines and each NH_2^+ residue is separated by four CH_2 residues [57,58]. Oligoamines consist of synthetic octa-, deca-, dodeca- and tetradecamines. The rationale behind the synthesis of oligoamines is that spermine at a concentration range of 50–100 μM and at near physiological ionic strength, leads to the collapse of DNA [61]. Oligoamines were found to initiate DNA aggregation at much lower concentrations (around 2–4 μM) than spermine and are more cytotoxic against several different types of cultured cancer cells as compared to other analogs tested so far [56,59,62]. In human prostate carcinoma cells, oligoamines are markedly more cytotoxic against tumor cells in culture than many previously described polyamine analogs [57]. A recent study from our group demonstrated that oligoamines effectively inhibit growth of human breast cancer cell lines in cell culture and nude mouse xenografts [62].

Alteration of polyamine metabolic enzyme activities by polyamine analogs

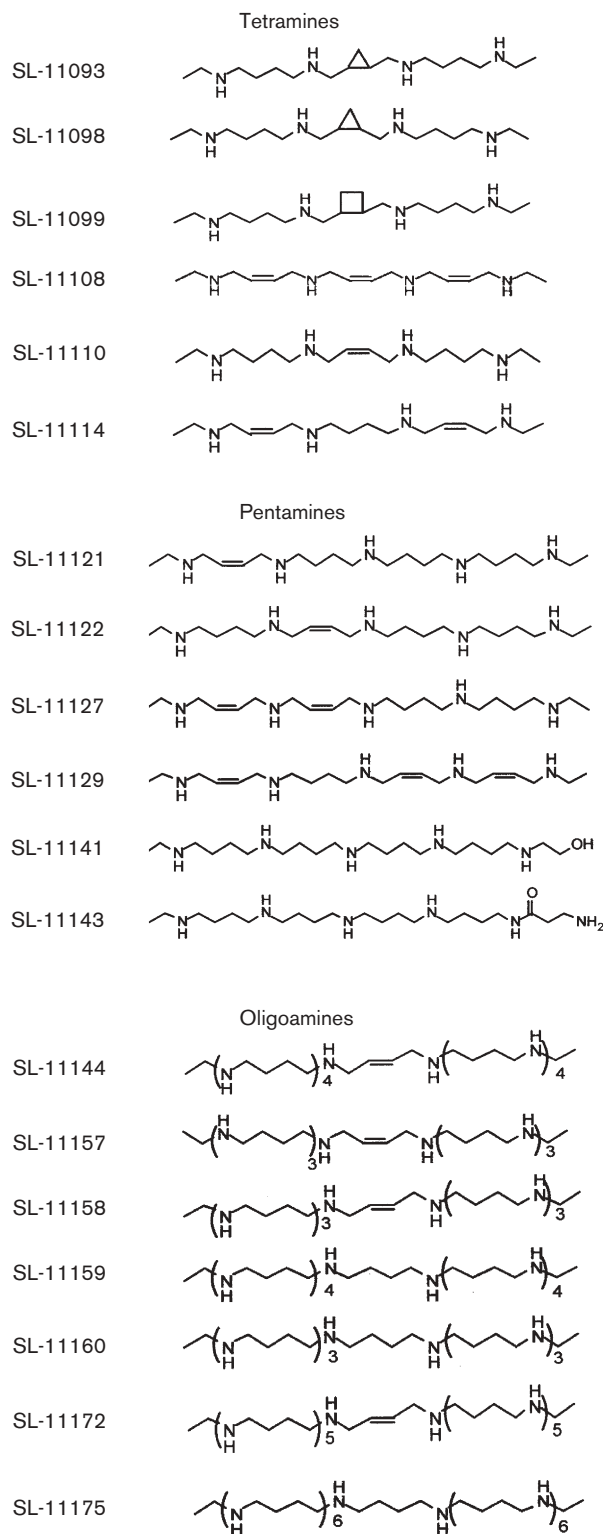
The requirement for polyamine action in cell growth and cancer progression has underscored the rationale of targeting polyamine metabolism as a therapeutic strategy. Although the *in vitro* success of DFMO did not translate well into the clinic, it indeed provides a proof of principle for targeting the polyamine metabolic pathway as a strategy for antineoplastic intervention. New strategies using the polyamine analogs may circumvent some problems encountered with DFMO for the following reasons: (i) unlike DFMO, polyamine analogs utilize the polyamine transporter system and compete with the uptake of the natural polyamines; (ii) polyamine analogs indirectly down-regulate ODC activity through induction of ODC antizyme, which may prevent the rapid turnover of ODC seen in DFMO-treated cells and therefore produce a steady net reduction in natural polyamines; and (iii) some polyamine analogs induce the activity of the rate-limiting catabolic enzymes, SSAT and SMO (PAOh1), that produce cytotoxic H_2O_2 , and therefore are more effective in tumor growth inhibition.

A recent study of 24 polyamine analogs, including tetramines, pentamines and various oligoamines, demonstrated that almost all of these analogs stimulated ODC-AZ synthesis and there was a good correlation between AZ induction and cytotoxicity [60]. Although the AZ induction contributes to down-regulation of ODC activity and depletion of polyamines by many of these analogs, it may also inhibit the activity of polyamine transport system and therefore limit the uptake and effectiveness of polyamine analogs [18]. Better understanding of the mechanisms underlying the inhibition of AZ on polyamine transport system may aid in the development of more effective agents that circumvent this problem.

Recently, there has been increasing interest in the role that polyamine catabolism plays in determining tumor cell response to analogs. This interest is a direct result of the findings that several different polyamine analogs such as BENSp_m and CPENSp_m significantly up-regulate the activities of polyamine catabolic enzymes. As described above, induction of SSAT in response to analogs and, in some cases, 'superinduction' of several thousand fold are the most interesting features of the response of this enzyme to the polyamine analogs [3,41]. The molecular mechanisms of the transcriptional regulation of SSAT are currently not fully understood. By using deletion/mutation and reporter construct strategies, a 9-bp consensus sequence (5'-TATGACTAA-3') that functions as a polyamine response element (PRE) and its associated transcription factors have been recently identified in the SSAT promoter [63]. This element was subsequently used to probe an expression library produced from BENSp_m-induced lung cancer cells, which led to the identification of NF-E2-related transcription factor (Nrf-2), a protein that was constitutively bound to the PRE [63]. Importantly, Nrf-2 expression was only observed in the analog-responsive cell lines with increased SSAT expression, suggesting that Nrf-2 is a specific regulator of analog mediated SSAT induction. A yeast two-hybrid screen using the leucine-zipper domain of Nrf-2 as bait identified a transcription co-factor designated polyamine-modulated factor-1 (PMF-1) [64]. Like SSAT, PMF-1 expression significantly increased with analog treatment in analog-sensitive tumor cell lines. Because of the lack of a DNA-binding domain, PMF-1 appears to modulate SSAT transcription indirectly through a unique leucine-zipper-coiled-coil interaction with Nrf-2 [65].

During the search for the human PAO gene, Wang *et al.* identified and characterized a protein, named human polyamine oxidase (PAOh1) [15], with a molecular weight similar to that of the PAO previously found by Holttä [66]. Vujcic *et al.* used a similar technique to find the identical clone termed spermine oxidase (SMO) [16]. The function of SMO (PAOh1) as a spermine oxidase has been confirmed [15,67,68]. To date, at least four splice variants of human and mouse SMO (PAOh1) genes have been identified [69,70]. These variants have unique biochemical properties that are capable of using multiple substrates including spermine, spermidine and acetyl-spermine [69]. However, the *in situ* relevance is not known. Rapid increase of SMO (PAOh1) mRNA levels in response to polyamine analog results in the enhanced oxidase activity, which occurs in a tumor and agent specific manner [71]. These findings suggest the potential of the newly identified SMO (PAOh1) family as a target for antineoplastic intervention. The contribution of analog-induced SMO (PAOh1) activity to analog cytotoxicity is currently being assessed and the preliminary results suggest that it does play a role in tumor cell response to specific agents. Understanding of the

Fig. 3



Chemical structures of tetramines, pentamines and oligoamines.

regulation of both SSAT and SMO(PAOh1) will provide valuable information in more selectively targeting cancer cells with newly designed analogs.

Disruption of polyamine–DNA interaction by polyamine analogs

Although the rapid cell death produced by several polyamine analogs has been well characterized, the exact mechanisms involved in this cytotoxicity have not been fully elucidated. A number of studies have suggested that DNA is a major target for the function of polyamines [72]. Polyamine–DNA interactions play a pivotal role in DNA conformational transitions, condensation/decondensation and stabilization [73,74]. Functional interaction between DNA and polyamines extends to DNA–protein binding, particularly those involving gene regulatory proteins [75]. As a result of the structural similarity, polyamine analogs might play a similar function leading to interference with the normal interactions of polyamine and DNA. Tumor cells treated with cytotoxic analogs are more sensitive than untreated or non-toxic analog-treated cells to nuclease digestion, suggesting a correlation between the ability of an analog to alter the chromatin structure and the analog cytotoxicity [76]. On the basis of computer modeling and physical/chemical studies of polyamine–DNA interactions, several polyamine analogs were synthesized by modifying the charge distribution along the surface of the aliphatic polyamine backbone. These compounds exhibited enhanced cell killing activity in human brain tumor cell lines [77]. A recent study has demonstrated a general correlation between the effects of newly developed long-chain oligoamine analogs on *in vitro* DNA aggregation and their cytotoxic effects in prostate cancer DuPro cells. By comparing the concentrations of oligoamines required to induce DNA aggregation with the concentrations necessary to kill tumor cells, it is apparent that oligoamines with a greater ability to aggregate DNA are more cytotoxic [57]. These studies provide evidence to suggest that modification of polyamine analog structures may affect their DNA binding abilities and, in turn, their cytotoxicity.

Effect of polyamine analogs on cell cycle regulation

It has been known for many years that natural polyamines are needed for normal cell cycle progression. Cellular polyamine levels and ODC activity peak at specific points during the cell cycle. ODC is activated in a biphasic manner with a first burst at late G₁ phase and a second one during the S/G₂ transition [78]. Early studies indicated that polyamines were essential for cells to enter S phase and polyamine depletion led to the inhibition of G₁/S phase transition [79]. Recent studies using an ODC inhibitor demonstrated that inhibition of ODC activity can lead to G₁ arrest in a variety of cell lines including tumor cells [80,81]. In human melanoma cells

expressing wild-type p53, treatment with BENSp_m activated the p53/p21/pRb cell cycle regulatory pathway leading to cell cycle block at G₁ phase, but in human melanoma cells lacking p53, no G₁ arrest was observed after BENSp_m treatment [82]. Other studies indicated that treatment with BENSp_m caused a delay of S phase progression and prolongation of the other cell cycle phases occurred at later time points [83]. Our recent studies indicated that the oligoamine analogs down-regulated cyclin D1 protein, inhibited CDK activity, activated the p53/p21/pRb pathway and induced cell cycle arrest at G₁ phase in breast cancer MCF-7 cells (Huang *et al.*, unpublished data). These results clearly suggest that the polyamines and regulatory enzymes play important roles in the progression of each phase of cell cycle and disruption of polyamine metabolism by polyamine biosynthesis inhibitors or analogs may lead to the cell cycle arrest in cancer cells.

Webb *et al.* found that the unsymmetrically substituted polyamine analogs, CHENSp_m and IPENSp_m, produced a significant G₂/M arrest in human NSCLC H157 cells [55]. As compared with spermine, these analogs exhibited different effects on tubulin polymerization, both in the presence and absence of microtubule-associated proteins (MAPs). Using immunohistochemical staining for tubulin, CHENSp_m was found to alter the microtubule density in the putative centrosome adjacent to the nucleus, but did not affect the cytoplasmic microtubules. The mechanism underlying the effect of these analogs on tubulin polymerization is not clear. It is possible that the loss of mitochondrial membrane potential is correlated with the observed alteration of tubulin polymerization in analog-treated tumor cells [84]. These findings indicate a completely novel antitumor mechanism for a subclass of structurally similar unsymmetrically substituted analogs. Their ability to change cell cycle transition and tubulin polymerization is evidently structure dependent.

Polyamine analog-induced programmed cell death

Interaction with chromatin, displacement of natural polyamines from their cellular binding sites, induction of polyamine catabolism and depletion of mitochondrial DNA have been proposed as possible mechanisms underlying the antitumor action of polyamine analogs [41,85]. Recent studies indicated that programmed cell death might be a common characteristic for polyamine analog-induced cytotoxicity in cancer cells. Polyamine analog-induced apoptotic cell death was first observed by our group following CPENSp_m treatment of the breast cancer cell lines, MCF-7 and MDA-MB-468, as well as the H157 NSCLC human lung tumor cell line [45,86]. To date, most of the bis(ethyl)polyamine analogs have been found to induce apoptosis in a variety of tumor cell lines [87–90]. However, the mechanisms involved in

polyamine analog-induced apoptosis have not been adequately defined. Catalysis of polyamines through analog-induced SSAT and PAO activities produces H_2O_2 as a byproduct, suggesting that analog-induced apoptosis may be, in part, due to oxidative stress resulting from H_2O_2 production. This hypothesis is supported by the finding that co-addition of catalase reduced high-molecular-weight DNA fragmentation, and inhibition of PAO activity by the PAO inhibitor, MDL 72,527, significantly reduced DNA damage and delayed apoptosis in CHENSpm-treated H157 cells [86]. In a recent study, Chen *et al.* demonstrated that SSAT siRNA, which specifically suppressed polyamine analog-induced SSAT activity, prevented the depletion of polyamine pools by analog and inhibited analog-induced apoptosis in human melanoma SKMEL-28 cells [91]. This mechanism of analog-induced cytotoxicity has important implications for the development of new analogs because SSAT induction occurs in some important solid tumors, but not generally in normal tissues.

Although the production of H_2O_2 by SSAT and PAO activity evidently plays an important role in apoptosis induced by some analogs, it is obviously not the only mechanism by which polyamine analogs kill tumor cells. For example, CHENSpm does not superinduce SSAT, but still induces significant apoptotic cell death in tumor cells [84]. Also, a novel oligoamine analog, SL-11144, induced oligonucleosome DNA fragmentation and activated apoptotic pathways in several human breast cancer cell lines without superinduction of SSAT activity. Multiple apoptotic mechanisms were associated with SL-11144 cytotoxic effects in specific breast cancer cell lines [62]. These results show that polyamine analog-induced apoptosis may occur through multiple structure-related and cell type-specific mechanisms. Typical features of apoptosis including cytochrome *c* release, activation of caspase-3 and cleavage of poly(ADP-ribose) polymerase (PARP) have been observed in oligoamine SL-11144-treated breast cancer MDA-MB-435 cells [62], BENSpm-treated melanoma SK-MEL-28 cells [87], and CPENSpm- and CHENSpm-treated lung cancer NCI H157 cells [86]. In H157 cells stably transfected with a Bcl-2 cDNA expression vector, CHENSpm-induced cytochrome *c* release, caspase-3 activation and PARP cleavage were significantly inhibited [84], suggesting that Bcl-2 activity may modulate the effect of polyamine analog on mitochondrial function and the activities of downstream apoptotic effectors. However, Bcl-2 did not abrogate CHENSpm-induced DNA fragmentation, indicating that a caspase-independent pathway also existed and could be activated by analog treatment. Similar results were observed in caspase-3-deficient MCF-7 breast cancer cells treated with oligoamine SL-11144, indicating that caspase-3 activation is not necessary for DNA fragmentation in this cell line [62]. These results demonstrate that both classical and non-classical apopto-

tic response pathways are operative in polyamine analog-induced programmed cell death. Treatment with the c-Jun NH₂-terminal kinase (JNK) inhibitor, SP600125, or overexpression of dominant-negative mutant c-Jun (TAM67) in human breast cancer MDA-MB-435 cells significantly increased their susceptibility to oligoamine-induced apoptosis, suggesting that an oligoamine-inducible JNK/AP-1 signaling pathway plays an anti-apoptotic role in this line in response to polyamine analog cytotoxicity [92]. In another study, the combination of the MAPK/ERK specific inhibitor, PD98059, with BENSpm nearly doubled the proportion of apoptotic cells in three melanoma lines as compared to BENSpm treatment alone [93]. These results indicate that the combination of polyamine analogs and other agents that target analog resistance mechanisms may significantly enhance the efficacy of tumor cell responsiveness to polyamine analogs.

Effects of polyamine analogs on cancer gene expression and ligand-receptor interaction

Polyamine analogs are known to alter the expression of a number of genes that are associated with the regulation of tumor cell proliferation, differentiation and apoptosis. BESpm treatment of the colon cancer cell line, CaCO₂, decreased c-*myc* oncoprotein in association with ODC inhibition, SSAT induction and polyamine depletion [94]. Gundogus-Ozcanli *et al.* reported that polyamines stimulated casein kinase II (CKII)-mediated phosphorylation of *myc* by 2- to 20-fold, and the polyamine analogs, BESpm and BE4x4, inhibited cancer cell proliferation partially through interference with CKII activity, suggesting that inhibition of cell proliferation by these polyamine analogs may involve the down-regulation of CKII activity [95]. Polyamines can also enhance binding of the transcription factor NF- κ B to its response element (NRE) in the promoter of a variety of target genes, a possible mechanism for the ability of polyamines to promote proliferation of certain cancer cell lines [96]. Treatment with the polyamine analogs, BENSpm and BESpm, significantly inhibited the transactivation of NF- κ B with a loss of the anti-apoptotic protein Bcl-2 in human breast cancer cells, suggesting that polyamine analog-induced down-regulation of NF- κ B signaling pathway may contribute to growth inhibition of breast cancer cells [97].

The effect of polyamine analogs on ligand-receptor interactions and function has also been examined. ERs are critical transcription factors in regulating the expression of a number of genes involved in cell proliferation and differentiation [98]. Polyamine biosynthesis inhibitors disrupted several ER-co-activator interactions and exogenous spermidine reversed this inhibition, indicating a specific role of polyamines in the interaction of ER with its co-activators to regulate ER mediated gene expression

[99]. Further studies showed that increased polyamine concentrations may alter DNA structure harboring estrogen response elements (ERE) to promote conformational changes and enhance the binding of ER to ERE [100]. Treatment of MCF-7 cells with the analogs, BENSp_m and BESp_m, resulted in transcriptional inhibition of an ERE-driven reporter plasmid through the disruption of the association between ER α and co-activator CBP/p300 [97]. The impact of polyamine analogs on other ligand–receptor systems has also been evaluated. Spermine, a natural tetramine, has been shown to exhibit concentration-dependent biphasic effect of binding of MK-801 to *N*-methyl-D-aspartate (NMDA) receptor [101]. Bergeron *et al.* used a group of spermine homologs and analogs to demonstrate that the effect of these analogs on MK-801 binding in the NMDA channel was related to their length and charge state [102]. This finding provides a structure–activity relationship to guide the design of more effective MK-801 agonists and antagonists for cancer therapy.

Summary and future directions

A number of polyamine analogs exhibit encouraging effects against tumor growth in both cell culture and animal studies. A wide range of molecular targets for polyamine analogs has been unveiled and the regulation of analogs on the activity of these targets is summarized in Table 2. AZ induction has been found in response to the treatment with many analogs, suggesting that this is

perhaps a common mechanism shared by different classes of analogs. The association of ‘superinduction’ of SSAT activity with a cytotoxic response to certain polyamine analogs has been noted. The complexity of the polyamine catabolic pathway is suggested by the identification of the SMO(PAOh1) genes. Cell cycle arrest and programmed cell death appear to be the common features of analog cytotoxicity in cancer cells. Alteration of DNA structure, production of toxic chemical and oxidative stress by catabolic enzyme induction, altered cancer-related gene expression, and activation of classic or non-classic apoptotic pathways have been proposed as the likely mechanisms by which polyamine analogs induce cell cycle arrest and apoptosis in tumor cells. These effects are clearly both cell type specific and structure dependent.

Despite the disappointing results obtained from early clinical trials of DFMO and BENSp_m, polyamine metabolism remains a rational target for cancer therapy. Increasing knowledge about the molecular mechanisms of polyamine analogs in cancer cells over the last two decades should guide optimum development and use of polyamine analogs in cancer treatment. Development of analogs that target a unique function of polyamine metabolism in tumor cells may possibly provide selectivity in cancer therapy. However, several barriers remain. For example, the molecular characterization of the eukaryotic polyamine transport system is still elusive.

Table 2 Molecular targets of polyamine analog action

Target	Active analog	Cellular effect	References
Polyamine metabolism			
AZ	symmetrically substituted bis(alkyl)polyamines; tetramines; pentamines; oligoamines	AZ induction; ODC inhibition; transport inhibition; polyamine depletion	[14,60]
ODC/SAMDC	symmetrically substituted bis(alkyl)polyamines; oligoamines	ODC inhibition, polyamine depletion; growth inhibition; cell cycle arrest	[3,6,42,62]
SSAT/PAO/SMO(PAOh1)	symmetrically and unsymmetrically bis(alkyl)polyamines	elevated enzyme activity; H ₂ O ₂ production; growth inhibition; apoptosis	[3,15,16,41]
DNA	symmetrically substituted bis(alkyl)polyamines; oligoamines; 5me-4-4-4; Q-Amm-4-4-4; BE-Q-Amm-4-4-4	altered chromatin structure; DNA collapse; altered transcription; apoptosis	[57,76,77]
Cell cycle	symmetrically and unsymmetrically bis(alkyl)polyamines	cell cycle arrest; altered tubulin function; growth inhibition	[55,82–84]
Apoptosis			
SSAT/PAO	symmetrically and unsymmetrically bis(alkyl)polyamines	production of H ₂ O ₂ , 3-acetamidopropanal and other toxic chemicals	[86,91]
caspases/PARP	oligoamines; symmetrically and unsymmetrically bis(alkyl)polyamines	induction of apoptosis	[62,87,86]
Bcl-2	CHENSp _m ; SL-11144	down-regulation of Bcl-2; inhibition of cytochrome c release; caspase-3 activation	[63,84]
AP-1	oligoamines	cell survival; apoptosis resistance	[62,92]
ERK	BENSp _m	apoptosis resistance	[93]
Cancer genes			
<i>c-myc</i>	BENSp _m	ODC inhibition; SSAT induction; polyamine depletion	[94]
casein kinase II	BESp _m ; BE4x4	growth inhibition	[95]
NF- κ B	BENSp _m	disruption of NRE activity; loss of Bcl-2 function	[96,97]
Ligand–receptor			
ER α	BENSp _m ; BESp _m	disruption of the association of ER α with co-activator; down-regulation of ERE activity	[97,100]
NMDA	BENSp _m ; BESp _m ; BEHSp _m ; tetraazaoctadecanes (5,4,5)	growth inhibition	[102]

Since elevated polyamine transport levels usually occur in rapidly dividing tumor cells, a better understanding of transport pathways may help in design of potent drugs to inhibit the transport system and disrupt the polyamine metabolism in tumor cells with minimal effects on normal tissues. It is possible that new techniques such as cDNA microarray and proteomic analysis will facilitate the identification and assessment of other specific genes involved in polyamine analog-mediated tumor growth inhibition. In addition, the combination of a polyamine analog with classic anticancer agents may enhance the effectiveness of these agents without increased toxicity [103]. Information from these studies will be important to lay the framework for further clinical trials.

Abbreviations

AbeAdo, 5'-[(Z)-4-amino-2-butenyl] methylamino-5'-deoxyadenosine; AdoDATAD, δ -adenosyl-1,12-diamino-3-thio-9-azadodecane; AdoDATO, δ -adenosyl-1,8-diamino-3-thiooctane; AMA, δ -(5'-deoxy-5'-adenosyl)-methylthioethylhydroxylamine; AZ, antizyme; BEHSpm, N^1,N^{14} -bis(ethyl)homospermine; BENSpm, N^1,N^{11} -bis(ethyl)-norspermine; BESpm, N^1,N^{12} -bis(ethyl)spermine; CGP-48664, 4-amidinoindan-a-one 2'-amidinohydrazone; CHEMSpm, N^1 -(cycloheptylmethyl)- N^{11} -ethyl-4,8-diazaundecane; CPENSpm, N^1 -(cyclopropylmethyl)- N^{11} -ethyl-4,8-diazaundecane; DFMO, α -difluoromethylornithine; ER, estrogen receptor; ERE, estrogen response elements; IPENSpm, (δ)- N^1 -(2-methyl-1-butyl)- N^{11} -ethyl-4,8-diazaundecane; MAP, (2R,5R)- δ -methylacetylenicputrescine; MAPs, microtubule-associated proteins; MFMO, α -monofluoromethylornithine; MGBG, methylglyoxal bis(guanyldihydrazide); NMDA, N -methyl-D-aspartate; Nrf-2, NF-E2-related transcription factor; ODC, ornithine decarboxylase; PAO, polyamine oxidase; PENSpm, N^1 -propargyl- N^{11} -ethyl-norspermine; PRE, polyamine response elements; PMF-1, polyamine-modulated factor-1; SAMDC, δ -adenosylmethionine decarboxylase; SMO(PAOh1), spermine oxidase; SSAT, spermidine/spermine N^1 -acetyltransferase.

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