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## p53 and Regulation of Bioactive Sphingolipids

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### Introduction

It is becoming increasingly evident in the literature that the sphingolipid metabolizing enzyme sphingosine kinase 1 (SK1) (E.C. 2.7.1.91) may act as an oncogene (Xia *et al.* 2000), (Pitson *et al.* 2005; Vadas *et al.* 2008; Bergelin *et al.* 2009). Although most studies focus on the activation/upregulation of SK1, we are interested in the novel concept of SK1 downregulation as a mechanism of tumor suppression. Our laboratory has previously shown that SK1 proteolysis is p53-dependent in Molt-4 leukemia cells: inactivation of p53 inhibited SK1 proteolysis after genotoxic stress (Taha *et al.* 2004). This review expands on this initial observation and is directed at exploring the p53-dependent regulation of sphingolipids and their metabolizing enzymes. Here, we provide background about the tumor suppressor p53 and discuss the currently known points of connection between the p53 and sphingolipid pathways, along with the therapeutic concept of tumor cell senescence.

### Tumor Suppressor Protein p53

Originally identified in its mutant form, p53 is now known to be one of the most commonly mutated tumor suppressor proteins in human cancers. In fact, 50% of all cancers appear to harbor a mutation in p53 (Vogelstein B 2000; Soussi 2007; Weisz L 2007). In healthy cells, low p53 concentrations are maintained by a negative-feedback loop in which p53 promotes Mdm2 expression which, in turn, tags p53 for nuclear export and proteasomal degradation (Vogelstein B 2000). When stress signals are recognized by the cell, p53 can accumulate in the nucleus and transcriptionally regulate genes to control the cell's fate. For instance, p53 can induce expression of p21, a cyclin-dependent kinase inhibitor, which leads to cell cycle arrest (Bates S 1998). p53 can also activate both death-receptor and mitochondrial apoptotic pathways by inducing the expression of various pro-apoptotic genes (Vousden KH 2002). Additionally, cytoplasmic p53 has been shown to induce non-transcriptional tumor suppressor functions (Green and Kroemer 2009), such that overexpression of a mutant p53 that lacks a DNA binding domain can still induce apoptosis in human cells (Haupt *et al.* 1995; Kakudo *et al.* 2005).

For apoptotic events, p53 must accumulate in the cell. Several kinases can activate p53 via phosphorylation after DNA damage, and such post-translational modifications can protect p53 from degradation. An alternative path for p53 accumulation is through the induction of p19 ARF, which can inhibit p53's degradation by Mdm2. (Evan G 1998). Such cellular stress causes p53 to accrue within the cell, which signals for apoptosis, growth arrest, and

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cellular senescence to prevent tumorigenesis. As reviewed extensively in various reports, (Kim and Deppert 2007; Strano *et al.* 2007; Weisz *et al.* 2007), p53's involvement in tumorigenesis could take one of three forms: (1) complete loss of wild type (WT) p53, leading to loss of the cell's growth-inhibitory response to physiologic or genotoxic stress; (2) a dominant-negative function of mutant p53 such that it can inactivate the tumor suppressive function of WT p53, i.e., inhibiting the formation of tetrameric complexes in cells; and (3) a mutant p53 gain of function such as altered gene expression regulation with oncogenic properties such as chemoresistance conferred by MDR-1, or inhibitory interactions with p53 family members, p63 and p73 (Weisz *et al.* 2007). For twenty years, the creation of genetically altered mice lacking p53 or expressing mutant p53 has produced animals prone to early carcinogenesis, and has thus illustrated the important tumor suppressive functions of p53 *in vivo* (Donehower *et al.* 1992; Jacks *et al.* 1994; Soussi 2007).

## p53 and Ceramide

Both p53 and ceramide have been shown to regulate cell cycle arrest, senescence, and apoptosis. Early work in our laboratory showed that gamma irradiation induced p53 and led to a p53-dependent increase in ceramide in human leukemia and mouse fibrosarcoma cell lines. However, ceramide was also observed to accumulate irrespective of p53 upregulation in other growth-suppressive pathways (Dbaibo *et al.* 1998). Later, this p53-dependent ceramide generation was shown to be driven by *de novo* ceramide synthesis through upregulation of ceramide synthases (CerS), particularly CerS5, leading predominantly to C<sub>16</sub>-ceramide accumulation (Panjarian *et al.* 2008). Likewise, several other reports suggest that ceramide accumulation is an important downstream mediator of the p53 response (Kim *et al.* 2002; El-Assaad *et al.* 2003; Villani *et al.* 2006), whereas others have shown that p53 and ceramide are concomitantly upregulated in response to various cell stressors (El-Assaad *et al.* 2003; Nasr *et al.* 2005; Villani *et al.* 2006), and that ceramide can accumulate and signal for apoptosis, irrespective of p53 status (Yang and Duerksen-Hughes 2001; Deng *et al.* 2009). Beyond apoptosis, ceramide has also been shown to mediate G1 arrest in a p53-independent manner through its induction of p21 to inhibit CDK2, leading to Rb dephosphorylation in hepatocarcinoma cells (Kim *et al.* 2000).

## Effects of Exogenous Ceramide Administration on p53

Some investigators claim that ceramide can act upstream of p53, such as in Ras-induced senescence (Castro *et al.* 2008). However, most of these studies involve exogenous ceramide treatment and may have limited applicability for understanding temporal relationships in endogenous cellular responses. Nonetheless, exogenous C<sub>2</sub>-ceramide treatment in primary cortical neuron cultures can increase c-jun, c-fos, and p53 mRNA (Willaime *et al.* 2001) and increase cellular p53 in mouse fibroblasts (Yang and Duerksen-Hughes 2001). Treatment with C<sub>2</sub>-ceramide has also been shown to increase p53 and p27 in a mouse B cell lymphoma cell line; however, when these cells were transfected with the transcription factor Nur77, which is associated with resistance to ceramide-induced apoptosis in these cells, p53 was markedly downregulated after ceramide treatment (Bras *et al.* 2000), suggesting a criticality of p53 functions downstream of exogenous ceramide addition. Ceramide can also activate PP2A which inhibits Bcl-2 phosphorylation, leading to increased p53/Bcl-2 binding and apoptosis (Deng *et al.* 2009), whereas Bcl-2 overexpression can suppress C<sub>2</sub>-ceramide-mediated apoptosis (Kolettas *et al.* 2006). Treatment of colon adenocarcinoma cells with C<sub>16</sub>-ceramide lead to differential expression of 51 proteins, one of which was Btf, which regulates p53 and other apoptosis-related proteins such as Mdm2, BAX, and pBcl-2 in these cells (Renert *et al.* 2009). Furthermore, treatment with 20  $\mu$ M C<sub>8</sub>-ceramide has been shown to induce increased expression of p53 leading to cell cycle arrest of endothelial cells in the

G1 phase (Lopez-Marure *et al.* 2000), and treatment with C<sub>2</sub>-ceramide has been shown to inhibit cell proliferation in WT, but not p53-deficient, tumor cells (Pruschy *et al.* 1999).

In contrast, C<sub>2</sub>-ceramide treatment has not been shown to change p53 levels while inducing PARP-mediated apoptosis in retinoblastoma cells (Vento *et al.* 1998). Other studies indicate that p53 protein was decreased after ceramide treatment in Bel7402 cells (Zhu *et al.* 2000), despite the occurrence of G0/G1 arrest in ceramide-treated B lymphoma Raji cells that lack TNF- $\alpha$  receptors (Kuroki *et al.* 1996). Furthermore, treatment of p53 WT or p53 mutant cells with exogenous ceramide led to apoptosis in both cell lines, whereas treatment with radiation only led to apoptosis in p53 WT cells, indicating that radiation-induced apoptosis is p53 dependent, and ceramide-induced apoptosis is not (Shi *et al.* 2001).

## p53 and Glycosphingolipids

Although most of the work connecting p53 and sphingolipids has focused on ceramide, considerable evidence relates p53 and more complex members of the sphingolipid family. For instance, adenovirus-delivered WT p53 decreases the expression of glycosphingolipids, namely gangliosides GD1 and GM1b, in human glioma cells (He *et al.*; He *et al.* 2007). Interestingly, the ganglioside GM1 increases in density on the surface of neuroblastoma cells during the switch from proliferation to differentiation, and the sugar chain of this glycosphingolipid was found to be a ligand for galectin-7 (p53-induced gene 1) which exerts neuroblastoma growth control. So, in addition to p53-linked intracellular interactions with sphingolipids, galectin-7 appears to connect p53 to glycosphingolipids on the cell surface (Kopitz *et al.* 2003).

### Effects of Exogenous Glycosphingolipid Administration on p53

As with ceramide, exogenous administration of gangliosides has revealed functions upstream of p53. GM3 treatment of colon cancer cells was shown to stimulate PTEN expression to inhibit PI3K/AKT/MDM2 survival signaling which led to the accumulation of p53 protein and increased expression of p21 as well as a p53-independent increase in p27 (Choi *et al.* 2006). The ganglioside GD3 which is overexpressed in some cancers, has been shown to be internalized by activated—but not resting—T cells, and it initiates a series of pro-apoptotic events that include p53 accumulation and the induction of reactive oxygen species, enhancement of Bax accumulation, increase in mitochondrial permeability, cytochrome c release, and the activation of caspase-9 (Sa *et al.* 2009). Interestingly, gangliosides produced by a T cell lymphoma were shown to increase p53 and induce apoptosis in bone marrow cells (Bharti and Singh 2000).

### p53-independent Apoptosis

Similar to ceramide, some studies suggest an independence of p53 and glycosphingolipid-induced apoptosis. Treatment of neuroblastoma cells with fenretinide leads to accumulation of ceramide which is then metabolized to GD3, which can result in p53-independent apoptosis. This effect is enhanced by co-treatment with modulators of ceramide metabolism in various solid tumor cell lines (Maurer *et al.* 1999; Maurer *et al.* 2000; O'Donnell *et al.* 2002; Corazzari *et al.* 2005). Moreover, the chemotherapeutic topoisomerase I inhibitors of the camptothecin family have been proposed to favor ceramide signaling by disturbing sites of synthesis (Golgi) and trafficking of glucosylceramide from the Golgi to lipid droplets, leading to caspase-3 activation and the induction of p53-independent apoptosis in colon HT29 cells (Chauvier *et al.* 2002).

## p53 and Sphingolipid Metabolizing Enzymes

Various enzymes involved in sphingolipid metabolism have been shown to interact with p53 (Figure 1). For instance, p53-dependent expression of acid ceramidase in glioma cells leads to a decrease in ceramide and can protect cells from gamma radiation-induced apoptosis (Hara *et al.* 2004). Ceramide synthase 2-null mice which are unable to produce very long chain (C<sub>22-24</sub>) ceramides accumulate C<sub>16</sub>-ceramide and have increased p53 and p21 messages after the first two weeks of life, leading to defects in liver homeostasis (Pewzner-Jung *et al.*). Interestingly, in the setting of Atm deficiency in which p53 is not notified of unrepaired DNA double-strand breaks, a ceramide synthase-mediated pathway of apoptosis prevails (Rotolo *et al.*).

### p53 and the Sphingomyelinase (SMase) Pathway

Because sphingomyelin is the most prevalent sphingolipid in cells, serving as an important cell membrane component, several studies of p53 and the sphingomyelinase pathway of ceramide formation propose topologically separate signaling mechanisms in which cell stress induces the membrane-based sphingomyelin pathway, whereas p53-dependent apoptosis occurs secondary to DNA damage. For instance, in some tissues such as the lung and heart endothelium and the pleura and pericardium mesothelium, radiation-induced apoptosis appears to be primarily dependent on acid SMase and, for the most part, independent of p53. In contrast, thymic apoptosis appears to be highly dependent on p53 and dependent to a lesser degree on acid SMase (Santana *et al.* 1996). Furthermore, ceramide generated by activation of acid SMase has been shown to be involved in p53-independent Fas-induced apoptosis (Sawada *et al.* 2002). The SMases and p53 are also involved in ceramide formation after TNF- $\alpha$ -induced caspase 8 activation which results in apoptosis. In glioma cells possessing WT p53, TNF- $\alpha$  stimulated ceramide formation via the activation of both neutral and acid SMases, accompanied by superoxide anion production, and induced mitochondrial depolarization and cytochrome c release, whereas p53-deficient cells were partially resistant to TNF- $\alpha$  and lacked superoxide anion generation and neutral SMase activation (Sawada *et al.* 2004). Thus p53 appears to be required for the activation of neutral, but not acid, SMase.

When fibrosarcoma cells with WT p53 were treated with neutral SMase, cell proliferation was reduced. However, neutral SMase treatment did not affect the proliferation of p53-deficient tumor cells, suggesting that p53 can be important for stress-induced apoptotic signal transduction cascades generated at the plasma membrane (Pruschy *et al.* 1999). One report suggested that p53 may modulate ceramide generation by activation of neutral SMase through the formation of superoxide anion in response to etoposide treatment (Sawada *et al.* 2001). However, in another study, Akt degradation and ceramide accumulation resulting from oxidative stress-induced neutral SMase activity may be p53 independent (Martin *et al.* 2002). Even so, a connection between the “membrane/sphingomyelin” and “DNA damage/p53” stress responses is evidenced by the coordinate regulation of neutral SMase and SK, because our laboratory has shown that various DNA damaging agents that induce p53 result in p53-dependent down regulation of SK1 (Taha *et al.* 2004). Importantly, SK isoforms and neutral SMase collaborate in their regulation of chemosensitivity by controlling ceramide formation and the downstream Akt pathway in human colon cancer cells (Nemoto *et al.* 2009).

### Connections between p53 and SK

As mentioned above, the SK1/S1P pathway has been shown to play a role in cancer progression and in tumor chemoresistance. Whereas abundant data have established that p53 is a tumor suppressor whose loss leads to carcinogenesis—a novel finding—our laboratory

reported that treating Molt-4 human leukemia cells with the DNA-damaging agent actinomycin D (act D) decreases SK1 (Taha *et al.* 2004). Because many chemotherapeutic agents can up-regulate p53 in cells via DNA damaging capacities, the requirement for p53 in act D-mediated SK1 down-regulation was investigated. Specifically, Taha and colleagues used a Molt-4 cell system expressing either the retroviral empty vector LXS<sub>N</sub> or the vector into which was inserted the E6 gene of human papilloma virus (Taha *et al.* 2004). The E6 protein has been shown to target p53 to ubiquitination and subsequent proteasomal degradation (Kessis 1993; Dbaibo *et al.* 1998). Treatment of cells for 24 hr with 10 ng/ml act D profoundly decreased SK1 protein in vector control cells; in contrast, cells overexpressing E6 failed to show loss of SK1 in response to act D (Taha *et al.* 2004). Of note, p53 was significantly induced in LXS<sub>N</sub> cells, and this was not observed in E6-expressing cells. Similar to SK1 protein, SK activity was also regulated by p53, in which act D markedly decreased (~50%) enzymatic activity in control cells but not in E6-transfected cells (Taha *et al.* 2004). These data strongly suggest that act D-mediated SK1 loss in Molt-4 cells depends on the accumulation of p53. Based on this initial observation by Taha *et al.*, further work in our laboratory established a conserved connection between p53 and SK1 in response to genotoxic stress in various cell lines (unpublished data).

Whereas p53 has been shown to regulate SK1 (Taha *et al.* 2004), SK2 appears to share an important function of p53, namely induction of the cell cycle inhibitor p21 (Sankala *et al.* 2007; Hait *et al.* 2009). The finding that p53 regulates one SK isoform and the other isoform regulates one of its main effectors, suggests that SK is an important connection between the p53 and sphingolipid pathways. Interestingly, SK1 is stimulated to stabilize HIF-1 $\alpha$  in response to reactive oxygen species (Ader *et al.* 2008), which has been shown to antagonize the p53 homologue in *C. elegans* and DNA damage-induced apoptosis (Sendoel *et al.*). This SK1 regulated protein has also been shown to be upregulated in p53 KO tumors (Samper *et al.* 2009). Another possible connection between p53 and SK1 is their role in autophagy. For example, cytoplasmic p53 has been shown to inhibit the AMP-dependent kinase, a positive regulator of autophagy, and activates the mammalian target of rapamycin (mTOR), a negative regulator of autophagy (Mathew *et al.* 2007), SK1 has been shown to inhibit mTOR and induce autophagy (Lavieu *et al.* 2006). Thus, p53-dependent downregulation of SK1 could contribute to signaling for autophagy.

### Connection between p53 and S1P Lyase

Downstream of SK is the terminal enzyme of sphingolipid catabolism, S1P lyase (E.C. 4.1.2.27). When S1P lyase is overexpressed in HEK293 cells, it potentiates apoptosis in response to DNA damage. This effect appeared to be independent of ceramide generation, but required p53, as well as caspase 2, p53-inducible death domain protein, and p38 MAP kinase (Oskouian *et al.* 2006). We found that cathepsin B or caspase-dependent degradation of SK1 can lead to decreased S1P. Thus the potentiated apoptosis observed in S1P lyase-overexpressing cells may depend on both the decreased production and increased degradation of cytoprotective S1P.

### p53 and Sphingolipids in Animal Models

Hu and co-workers reported that, in a mouse model of azoxymethane-induced colon cancer—whose chow was supplemented with sphingomyelin, WT (p=0.15) and p53 HZ (p=0.12) mice had a reduced but nonsignificant tumor incidence. Thus, increasing intestinal SMase activity and suppressing proliferation, the additional dietary sphingomyelin did not promote apoptosis and did not significantly protect mice (Hu *et al.* 2008). Long-term treatment with the marine sponge glycolipid,  $\alpha$ -galactosylceramide, can significantly delay sarcoma development in p53 KO mice; however, it did not affect lymphoma development in these mice. Effects of  $\alpha$ -galactosylceramide on sarcomagenesis are thought to rely on its

activation of natural killer cells and CD8<sup>+</sup> cytotoxic T-cells to secrete anti-angiogenic interferon gamma (Hayakawa *et al.* 2003). Furthermore, in a dextran sodium sulfate (DSS)-induced colitis model which has been shown to lead to enhanced neoplasia in p53 KO mice (Chang *et al.* 2007), a COX-2 inhibitor reduced the incidence of colon cancer (Mukawa *et al.* 2008); likewise SK1 KO mice with no colonic COX-2 response were also protected from the development of DSS-induced colitis (Snider *et al.* 2009). While not directly linked, these studies suggest a connection between the p53 and sphingolipid pathways in various animal models of cancer. Moreover, they predict a complex and interwoven role of these pathways in controlling proliferation, angiogenesis, and inflammation, all of which are critically involved in carcinogenesis.

## Tumor Cell Senescence, p53, and Sphingolipids

Data from our laboratory and others have suggested a link between the p53 tumor suppressor protein and changes in sphingolipids (Summarized in Figure 2). The pro-apoptotic sphingolipid, ceramide, has been shown to increase in a p53-dependent manner in response to genotoxic stress (Vousden KH 2002). In fact, several inducers of apoptosis have been shown to increase ceramide in cells concomitant with p53 activation (Haupt *et al.* 1995; Kakudo *et al.* 2005; Green and Kroemer 2009). Although ceramides have been reported to induce apoptosis in cells irrespective of p53 status (Chipuk *et al.* 2003; Chipuk *et al.* 2004), p53 activation may enhance the effects of ceramide (Evan G 1998; Mathew *et al.* 2007). Work in our laboratory suggests that treatment of Molt-4 leukemia cells with the act D results in proteolysis of SK1 in a p53-dependent manner (Taha *et al.* 2004). Furthermore, Oskouian *et al.* showed that overexpression of S1P lyase can potentiate an apoptotic response to DNA damage in a p53-dependent manner (Strano *et al.* 2007). These studies suggest that some enzymes controlling cellular levels of ceramide and S1P appear to act as effectors of the p53 tumor suppressor pathway.

Similar to the aforementioned studies connecting the sphingolipid and p53 pathways, the majority of studies to examine effects of chemotherapeutic agents on cancer cell responses have focused on apoptotic mechanisms. Apoptosis may be regarded as the most desired outcome and a more straightforward aspect to study (with several well-defined events such as caspase activation, cytochrome c release, and DNA fragmentation). However, recent studies indicate that, in addition to apoptosis, tumor cell senescence may be a crucial determinant for the *in vivo* response to chemotherapy (Berns 2002; Roninson 2002; Ricci and Zong 2006), thus raising the possibility that mechanisms of tumor cell senescence could be attractive therapeutic targets.

## Tumor Cell Senescence as a Chemotherapeutic Mechanism

The connections between sphingolipids and p53 discussed above suggest that p53-dependent regulation of sphingolipid could also be an important component of the senescence response. Recent studies that illustrate the concept of tumor cell senescence are contributions of Lowe's group who suggest that Bcl-2 overexpression was more effective at inhibiting tumor shrinkage in mice than was the loss of p53 (Schmitt *et al.* 2002). However, these studies also showed that in the long-term, p53 null-tumors were more resistant to chemotherapy than Bcl-2-overexpressing tumors. These studies suggest that inhibition of apoptosis was not the sole determinant of tumor response outcomes. In fact, when these tumors were examined closely, it became evident that tumors overexpressing Bcl-2, but not those that null for p53, could engage a senescence pathway and thus overcome resistance to chemotherapeutic agents. In more recent studies, Lowe's group developed an elegant model in which they could conditionally regulate endogenous p53 in a p53-deficient mouse hepatocarcinoma model. They reported that with this model, reactivation of p53 led to tumor regression

primarily by cellular senescence, and by triggering an innate immune response that targeted tumor cells and resulted in their clearance *in vivo* (Lozano; Xue *et al.* 2007). Mice with mutations in p53 that abrogate its apoptotic signaling but maintain its induction of p21 and senescence, protect chromosome stability and delay tumor onset compared to full knockout of p53, highlighting the importance of cell cycle arrest in p53-induced tumor suppression (Liu *et al.* 2004). To confirm this connection, compound mutant mice homozygous for the p53R172P mutation that lacks p21, develop tumors with almost the same survival curve as p53-null mice (Barboza *et al.* 2006). Furthermore, restoration of p53 functions in sarcomas from p53-null mice results in tumor regression via a senescence phenotype whereas restoration of p53 function in T-cell lymphomas lacking p53 results in apoptosis and tumor regression (Ventura *et al.* 2007).

In addition, ample evidence in the literature demonstrates that chemotherapeutic agents induce cancer cell senescence (Joyner *et al.* 2006; Probin *et al.* 2007). Interestingly, this program of senescence is somewhat distinct from normal cellular senescence in that it occurs rapidly (within 7 days of treatment) and does not appear to be mediated by telomere shortening but rather by activation of p53 pathways (Roninson 2002). Of note is that ceramide has been shown by our laboratory (Venable *et al.* 1995) and subsequently, by others (Mouton and Venable 2000; Gao *et al.* 2001), to induce a rapid program of cellular senescence in cells that have no telomerase activity, indicating that it can engage a non-telomere-dependent program of cellular senescence. Moreover, we and others have shown that ceramide is significantly increased in cellular senescence (Venable *et al.* 1995; Venable *et al.* 2006). Given the importance of the senescence response for the tumor-suppressive function of p53, the involvement of ceramide in cellular senescence provides another putative intersection point between the p53 and sphingolipid pathways.

In fact, data from our laboratory further suggests that downregulation of SK1 by p53 in response to DNA damage could result in ceramide-induced senescence. As previously mentioned, act D was shown to signal for p53-dependent degradation of SK1 (Taha *et al.* 2004). Other stress-inducing agents such as etoposide and TNF were also shown to induce proteolytic degradation of SK1, leading to alterations in sphingolipid levels (Taha *et al.* 2005; Taha TA 2004). Importantly, knockdown of SK1 results in a 2-fold increase in the upstream signaling molecule ceramide and induces G1 arrest in MCF-7 breast cancer cells (Taha *et al.* 2006). Collectively, these results suggest that basal SK1 may potentially regulate growth-inhibiting lipid ceramide, and that p53-dependent degradation of SK1 may shift the sphingolipid balance toward growth inhibition and cell senescence.

## Summary

Both the sphingolipid and p53 pathways are important regulators—and apparent collaborators—of cell-fate decisions. Whereas some investigations have suggested that ceramide and more complex sphingolipids function upstream of p53 or in a p53-independent manner, other studies propose that p53-dependent alterations in these sphingolipids can also contribute to apoptosis. Further studies focusing on sphingolipid metabolizing enzymes have revealed that they function similarly both upstream and downstream of p53 activation. However, whereas various components of the sphingolipid and p53 pathways may simultaneously function to elicit apoptosis and/or growth inhibition, SMase and SK1 may undergo explicit regulation by p53 that could contribute to ceramide-induced senescence in cells. Thus, we propose that regulation of bioactive sphingolipid signaling molecules could be of therapeutic benefit in the treatment of p53-dependent cancers.

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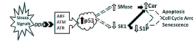
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**Figure 2.**

Proposed points of p53-dependent regulation of the sphingolipid pathway after cell stress.

DDR: DNA Damage Response, ARF: Alternate Reading Frame product, ATM: Ataxia

Telangiectasia Mutated protein, ATR: Ataxia Telangiectasia and Rad3-related protein, p53:

Tumor Suppressor Protein p53, SMase: Sphingomyelinase, SK1: Sphingosine Kinase 1, Cer:

Ceramide, and S1P: Sphingosine-1-Phosphate.