

We have developed a series of PARP inhibitors, exemplified by ABT-888, that potently inhibit PARP as chemosensitizing agents. In combination studies, ABT-888 delays DNA repair induced by DNA damaging chemotherapeutics. At the same concentrations, ABT-888 potentiates the cytotoxicity of these agents in human tumor cell lines. The potentiation was demonstrated with a wide range of agents including TMZ, Irinotecan, platinum, etc.. The delay in DNA repair and the potentiation of cytotoxicity by DNA damaging agents are modulated by genetic defects on DNA repair pathways. ABT-888 has excellent PK properties and can be administered orally or parentally. It efficiently crosses blood brain barrier, making it an excellent candidate to combine with TMZ for treating intracranial tumors. ABT-888 is well tolerated in animal models. ABT-888 inhibited PAR formation in tumors from the treated animals, which is used as a biomarker for ABT-888. Treatment of ABT-888 enhances tumor growth inhibition by a variety of chemotherapeutic agents such as TMZ, cisplatin, carboplatin, cyclophosphamide, etc. in multiple xenograft models. ABT-888 was also shown to potentiate the antitumor efficacy of TMZ in a rat 9L glioma orthotopic model. ABT-888 is currently in clinical trials. We will disclose the structure of ABT-888 at the AACR meeting.

**#1458 MG-132 sensitizes TRAIL-resistant prostate cancer cells by activating c-Fos/c-Jun heterodimers and repressing c-FLIP(L).** Wenhua Li, Xiaoping Zhang, Aria F Olumi. *Division of Urologic Surgery, Beth Israel Deaconess Medical Center, Harvard Medical School, Boston, MA.*

Tumor necrosis factor-related apoptosis-inducing ligand (TRAIL) is a promising anti-cancer agent since it induces apoptosis in cancer cells but not in normal cells. Unfortunately, some cancer cells develop resistance to TRAIL-induced apoptosis. Therefore, it is clinically important to determine the molecular mechanisms that differentiate between TRAIL-sensitive and -resistant tumors. Previously, we have shown that the anti-apoptotic molecule, c-FLIP(L) (cellular-FLICE-inhibitory proteins long isoform), is necessary and sufficient to maintain resistance to TRAIL-induced apoptosis. We have found that c-FLIP(L) is transcriptionally regulated by the AP-1 family member protein, c-Fos. Here we report that MG-132, a small molecule inhibitor of the proteasome, sensitizes TRAIL-resistant prostate cancer cells by inducing c-Fos and repressing c-FLIP(L). c-Fos, which is activated by MG-132, negatively regulated c-FLIP(L) by direct binding to the putative promoter region of the c-FLIP(L) gene. In addition to activating c-Fos, MG-132 activated another AP-1 family member, c-Jun. We show that c-Fos heterodimerizes with c-Jun in order to repress transcription of c-FLIP(L). Therefore, MG-132 sensitizes TRAIL-resistant prostate cancer cells by activating the AP-1 family members, c-Fos and c-Jun, which in turn repress the anti-apoptotic molecule, c-FLIP(L).

**#1459 Anticancer activities of proteasome inhibitor MG132 on human malignant pleural mesothelioma cells.** Bao-Zhu Yuan, Joshua A Chapman, David Lowry, Steven H Reynolds. *Nat'l Institute for Occupational Safety & Health, Morgantown, WV.*

Malignant pleural mesothelioma (MPM), caused mainly by occupational or environmental exposure to asbestos, is an aggressive malignancy with a very poor prognosis. The general resistance of MPM to current therapeutic modalities and an ongoing increase in the incidence of MPM demonstrate the need for new treatments for this deadly disease. Proteasome inhibitors have emerged as a category of promising reagents for treating human cancers. MG132, the most commonly used proteasome inhibitor, has demonstrated its ability to induce apoptosis in some types of human cancer. In our initial attempt to determine the therapeutic effect of the proteasome inhibitors on human MPM, we chose MG132 as a prototype of the proteasome inhibitors and characterized its anti-cancer effects on NCI-H2502 and NCI-H2452, two human thoracic MPM cell lines. It was observed that as low as 0.5  $\mu$ M MG132 caused significant cell death in both MPM cell lines. Cell death derived mainly from apoptosis as evidenced by the appearance of a pre-G1 peak in flow cytometry analysis and propidium iodide staining. It was also observed that MG132-induced apoptosis in both MPM cell lines was accompanied by cleavage of the PARP and caspase 3 and 7 proteins. Treatment with either caspase 3 inhibitor or Z-VAD, a broad spectrum caspase inhibitor, resulted in inhibition of MG132-induced apoptosis, suggesting that MG132-induced apoptosis operates through a caspase-dependent mechanism. Further demonstration of caspase 9 activation, which is likely the consequence of MG132-induced Bid protein activation and the subsequent release of the Smac/Diablo protein from the mitochondria, suggests that a mitochondria-caspase 3 apoptosis pathway is responsible for mediating MG132-induced apoptosis in MPM cells. Analysis of MG132's effect on invasion by MPM cells, as tested by an *in vitro* Matrigel invasion assay, demonstrated that 0.1  $\mu$ M MG132, which showed no effect on apoptosis induction, significantly inhibited invasion by both NCI-H2452 and NCI-H2502 cells. MG132-inhibited tumor cell invasion was found to be associated with reduced Rac1 activity. These

observations demonstrate proapoptotic and anti-invasion activities human MPM cells and lead to the suggestion that MG132 may be a for clinical treatment of human MPM.

**#1460 MS-275 up-regulates FAS (CD95/APO1) expression caspase-8 mediated apoptosis in medulloblastoma cells.** Doll Neeta S. Kang, Johannes Wolff, Vidya Gopalakrishnan. *University of Texas Cancer Center, Houston, TX.*

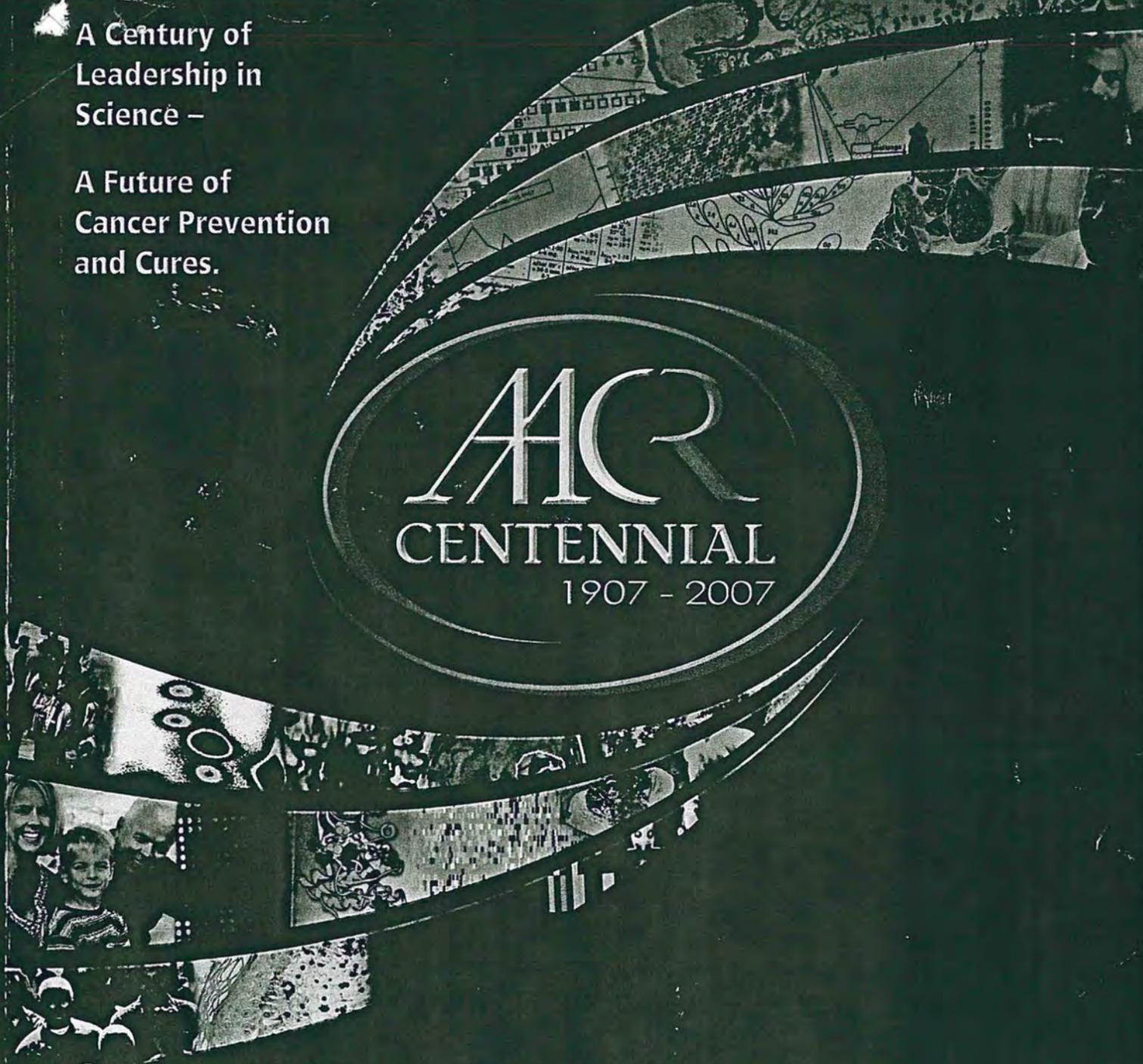
Histone deacetylases (HDACs) are key enzymes that modulate cell fate and regulate gene expression in cells by deacetylation of co-residues in histones. Since aberrant expression of genes involved in differentiation and apoptosis is an important mechanism of tumorigenesis, we examined the effect of the histone deacetylase inhibitor MS-275, on medulloblastoma cell viability and gene expression *in vitro*. We demonstrated that medulloblastoma cell lines, Daoy and D283, are sensitive to treatment and exhibit a rapid accumulation of cells with a sub-G1 population. This is accompanied by an increase in the expression and activity of caspase-3 as well as an induction of poly ADP-ribose (PARP) cleavage was also observed as early as 24-48 hours post-treatment with MS-275. To further confirm the involvement of the extrinsic pathway of apoptosis in response to MS-275, we measured the expression of the FAS/Apo-1 by real-time PCR, western blotting and immunofluorescence. These experiments revealed that MS-275 not only increased the expression of FAS receptor, but also promoted a clustering of receptors as evidenced by immunoprecipitation of FAS receptor on the surface of MS-275 treated cells. FAS ligand expression remained unchanged following drug treatment. It appears to be necessary for MS-275 induced apoptosis, providing further evidence for oligomerization of FAS receptors as a possible mechanism by which it promotes cell death. Finally, the introduction of a dominant negative FAS receptor into Daoy cells interfered with the ability of MS-275 to promote apoptosis. This first demonstration that up regulation of FAS receptor and caspase-8 by MS-275 plays an important role in mediating apoptosis in medulloblastoma cells.

**#1461 OSW-1 induces apoptosis by activation of calpains and endoplasmic reticulum cell survival mechanism.** Celia Garcia-Perez, Yan Zhou, Weiqin Lu, Zhou Chen, Yumin Hu, Zhi Peng Huang. *UT M.D. Anderson Cancer Center, Houston, TX and Iowa, Iowa City, IA.*

Previous studies showed that OSW-1, a naturally occurring compound isolated from the bulbs of *Ornithogalum saundersiae*, exhibited potent cytotoxicity and selectivity *in vitro*. This compound was shown to effectively kill cancer cells including those resistant to conventional anticancer agents. A unique mechanism of action of OSW-1 not shared by the conventional cytotoxic agents was suggested. Biochemical analysis showed that OSW-1 also caused intracellular calcium release leading to massive cell death. The current study was undertaken to study the mechanism of action of OSW-1 and to investigate how OSW-1 disturbances may influence the fate of the cells. Using calcium fluorescence and flow cytometry, we showed that OSW-1 caused calcium elevation in a concentration- and time-dependent manner. To evaluate whether extracellular calcium entry might play a role in OSW-1-induced calcium elevations, bioassays using calcium-free medium revealed a reduction in calcium elevation. Calcium increase to 50% calcium increase in calcium-free medium treatment, suggesting that OSW-1 may cause a disturbance in the cell membrane that allows extracellular calcium entry. Electron microscopy showed that OSW-1 treatment caused severe swelling of the endoplasmic reticulum and mitochondria, which likely contributed to a further disturbance in cellular homeostasis. To examine the role of calcium increase in the cytotoxicity of OSW-1, we used a calcium chelator to deprive cells of calcium and showed that manipulation significantly reduced OSW-1-induced cell death. Since calpains were observed after OSW-1 treatment, the role of calcium-dependent calpains in mediating OSW-1-induced cell death was examined. The calpain inhibitor ALLN. Co-incubation of OSW-1 and ALLN significantly reduced cell death compared to OSW-1 alone. Importantly, OSW-1 treatment of cells in the presence of ALLN did not affect the degradation of GRP78 (an ER protein that promotes cell survival during ER stress) or the chelation of calcium by BAPTA-AM or inhibition of calpains by ALLN. Together, these results suggest that OSW-1 exerts its cytotoxicity in a calcium-dependent manner not only by activating calcium dependent-apoptosis but also by disturbing the endoplasmic reticulum through calpain-mediated cleavage of the endoplasmic reticulum.

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