News and Commentary

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New light on Smac mimetics and breast cancer

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The evasion of cell death is an important contribution to cancer development and may also confer resistance to chemo and radiotherapy.¹ Therefore, reinstating a functional cell death machinery is one potential strategy for cancer therapy. One way to promote cell death induction is to use Smac mimetics (SMs),² which are small molecules mimicking the proapoptotic protein Smac. Smac inhibits members of the inhibitor of apoptosis protein (IAP) family of which cellular IAP 1 (cIAP1) and 2 (cIAP2) are important for apoptotic signaling involving the death receptor TNFa receptor 1 (TNFR1). SMs, thereby, re-direct the TNFR1 from NF-kB signaling towards apoptosis. This is achieved by downregulating, and thus inhibiting, cIAP1 and cIAP2 and their function as ubiquitinators of proteins recruited to the TNFR1. Without cIAP-mediated ubiquitination of proteins such as receptor interacting protein 1 (RIP1), the NF-kB signaling cannot be conveyed. Instead, a complex containing RIP1 recruits caspase 8, activating the caspase cascade and apoptosis. In addition to redirecting the TNFR1, SM-mediated downregulation of cIAPs also leads to accumulation of NF-kB inducing kinase (NIK) and the initiation of the noncanonical NF-KB signaling pathway, which can result in induction of TNFa synthesis and possible stimulation of the TNFR1 in an autocrine manner. Whether SM treatment leads to upregulation of TNFa or not has been suggested to determine sensitivity to SMs.3,4,5

Breast cancer is the most common cancer for women worldwide. Screening protocols and advances in therapy have resulted in a 30% increase in 5-year survival between 1975 and 2005 to 90%.⁶ However, for basal-like breast cancers, which typically are estrogen- and progesterone receptornegative as well as lack ERBB2 amplification, there is still no targeted treatment and they are more difficult to treat.7,8 Therefore, strategies to treat this patient group are warranted. Some basal-like breast cancer cell lines, such as MDA-MB-231 are indeed sensitive to SMs suggesting that this pathway may be a potential target for therapy. However, most cell lines are resistant to SM as single treatment. Now, Cornmark et al.9 present a report demonstrating that basal-like breast cancer cell lines can be sensitized to SMs by activating protein kinase C (PKC) (Figure 1). They report that combining the PKC activator TPA with SM leads to cell death in three different basal-like breast cancer cell lines. The cell death can be blocked by using a $TNF\alpha$ -blocking antibody indicating the importance of TNFa in eliciting cell death. When investigating

TNF*a* more closely, they found that PKC activity resulted in increased levels of TNF*a*, both at the mRNA and protein levels. The PKC-induced *de novo* production of TNF*a* was shown to be mediated by the canonical NF- κ B pathway⁹ (Figure 1).

It has previously been described that cancer cell lines sensitive to SM as a single agent are dependent on autocrine $TNF\alpha$ production. However, a majority of cancer cell lines are insensitive to SMs that seems to be due to an inability to induce $TNF\alpha$ expression.³ To investigate mechanisms explaining differences in Smac sensitivity, the response to SM in sensitive MDA-MB-231 and insensitive MDA-MB-468 breast cancer cell lines was compared. One striking difference was that the sensitive cell line had many more upregulated genes after SM treatment compared with the insensitive cell line where few genes were differentially expressed. However, both cell lines responded to SM with increased expression of the BIRC3 gene that encodes cIAP2. Furthermore, both cell lines responded to SM in the same way by initiating noncanonical signaling as detected by p100 processing and p52 nuclear translocations.

The noncanonical NF-kB signaling pathway has been suggested as the mediator of SM-induced TNFa production.3-5,10 Nevertheless, in MDA-MB-468 cells, SM induces the noncanonical pathway and gene expression (BIRC3 induction), but still no increase in $TNF\alpha$. This raises the possibility that other pathways are critical for SM-mediated TNFa synthesis. Therefore, Cornmark et al.9 used a siRNA approach to target either NIK and the noncanonical NF-kB pathway and/or IKK β and the canonical NF- κ B pathway. They found that only downregulation of both pathways resulted in a decrease in SM-mediated TNFa production that also resulted in a suppression of SM-induced cell death. However, although the absolute TNFa levels were suppressed, the relative induction still persisted as TNFa levels were markedly suppressed by the siRNAs also in unstimulated cells. This suggests that pathways other than NF-kB may be necessary to obtain sufficient TNF α levels to induce cell death.

Thus, Cornmark *et al.*⁹ have shown that the SM-sensitive breast cancer cell line MDA-MB-231 needs either the canonical or the noncanonical pathway to mediate TNFa induction and cell death. Even if both pathways are suppressed, SM can still increase TNFa synthesis to some extent, suggesting that other mechanisms are also involved.

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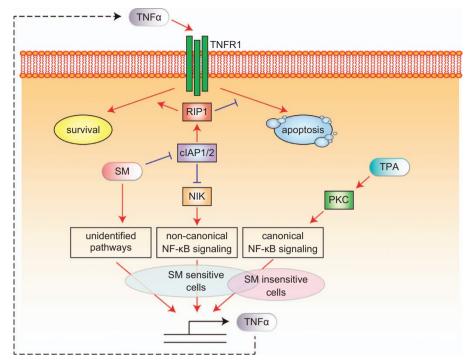


Figure 1 PKC activation induces $TNF\alpha$ production in cells sensitizing them to SMs. As reported by others, SM inhibits clAP1 and clAP2 leading to a redirection via RIP1 of the TNFR1 signaling towards apoptosis. In addition, SM-mediated clAP1/2 inhibition leads to accumulation of NIK, propagation of the noncanonical signaling pathway and $TNF\alpha$ production in SM-sensitive cell lines. However, as the authors report this may not be the sole pathway by which SM mediates $TNF\alpha$ production in SM-sensitive cell lines, other yet unidentified pathways may also contribute to $TNF\alpha$ production. In addition, 12-O-tetradecanoylphorbol-13-acetate (TPA) that activates PKC initiates canonical NF-xB signaling resulting in increased autocrine $TNF\alpha$ production and sensitizes insensitive cell lines to SM

Taken together, these results further support previous studies² that a simultaneous induction of TNFa may be one strategy to increase SM sensitivity. In basal-like breast cancer cells, this can be achieved by PKC activation that may open up for novel ways to target this group of breast cancers.

Conflict of Interest

The authors declare no conflict of interest.

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