Different Faces of Fas Signaling in Cancer Cells

Magdalena Szarynska* and Agata Olejniczak Kęder
Department of Histology, Poland

Abstract
Fas signaling pathway is known to be engaged in elimination of unwanted cells by apoptosis, however, new discoveries presented its alternate face, namely FasR and FasL play pro-cancerous functions facilitating cancer progression, invasion and metastasis. Here, we provide a brief summary of knowledge concerning Fas signaling pathway merits and disadvantages in cancer. Fas molecules were found to be engaged in many pathways induced by different therapeutically compounds, such as aspirin and antibiotics. This suggest that Fas signaling pathway may be employed as adjuvant factor for immunotherapy of variable design. Although, the literature presents some first optimistic results, we must intensify efforts to finally introduce Fas-associated procedures into standard therapeutic options.

Introduction
The Fas signaling is known to exert very variable effects in wide spectrum of cells. The FasR/FasL (CD95/CD95L) proteins can be associated with both positive (physiological) and negative (pathological) effects. Nowadays, there is a growing interest in the elucidation of the Fas signaling role in the pathogenesis and progression of various cancers. Additionally, it was proven that the expression of FasR/FasL in colorectal cancer is associated with worse prognosis, metastasis and recurrence [1-6] the aspects of cancer biology which cancer stem cells are responsible for [7]. On the one hand, Fas signaling pathway is considered as a potential target for anticancer therapy and, on the other hand, is hoped to be exploit as therapeutic tool. The approach to this issue is dynamically update since the state of knowledge concerning Fas signaling functions is rapidly developing. The most established pro-apoptotic activity of FasR/L signaling is the elimination of non-CSC cancerous, virus-infected or useless/autoreactive T cells by cytotoxic T lymphocytes [8]. Drug therapy combining the multi-kinase inhibitor Sorafenib and the histone deacetylase inhibitor Vorinostat was shown to activate Fas-mediated apoptosis by promoting receptor tyrosine phosphorylation or contribution to FasR activation via initial facilitation of ROS generation and subsequent FasL expression [9]. Additionally, Fas signaling was proven to be associated with HSPs. Although it was originally demonstrated in rat global brain ischemia model, we assume that similar relationships exist in cancer cells as well. Inhibition of HSP90 proteins reduced FasL expression and induced neuroprotective effect [10]. Moreover, the heat shock proteins accumulation was demonstrated as a specific mechanism increasing protein stability and reducing a turnover during Fas-mediated apoptosis in Jurkat cells [11].

The expression of FasR/FasL by cancer cells implies that they are themselves resistant to Fas-mediated apoptosis [8]. One of the mechanisms used by the cancer cells to acquire insensitivity to pro-apoptotic signals is elimination (or rather prominent reduction) of the FasR from their surface what disrupts the expression of factors engaged in apoptosis [12-14]. This is so called “cancer evasion mechanism” [15]. It claims that tumor cells escape from immune cells attack and attract (or kidnap) immune elements for their own good. The decreased FasR level diminished the apoptosis probability, but the level of FasR in cancer cell membranes necessary for their survival was estimated to be 1000-times lower than the level required for its pro-apoptotic signaling [16]. The current observations regarding cancer cells expression of FasR and their resistance to apoptosis induction suggest that Fas signaling molecules play also non-apoptotic functions within cancer tissue [17,18]. The observed preservation activity of Fas signaling for normal stem cells homeostasis and embryogenesis of the nervous system [16,19-22] suggested that Fas signaling pathway may also have a non-apoptotic function in cancer stem cells (CSCs) [23].

The group of Marcus Peter showed that the elimination of either FasR or FasL causes death of cancer cells (in vitro and in vivo) through a process termed DICE (death induced cell elimination) [24].
by CD95 or CD95L elimination). DICE is a necrotic form of mitotic catastrophe characterized by cell swelling, ROS production causing DNA damage and activation of caspase-2 following mitochondrial outer membrane permeabilization [8, 24]. During DICE many different pathways directing cells to death were proved to be stimulated thus it is hardly possible to modulate that process. These observations proved that DICE is a naturally occurring antitumor defense mechanism, which can eliminate cancer cells devoid of FasR. The pro-survival activity of FasR is believed to be mostly relevant to cancer cells, since cancer cells rarely or even never have mutated or deleted both alleles of FasR. Additionally, it was shown that none of the normal tissues during embryonic development in FASR/Fasl knockout mice showed a growth defects or signs of cell death [8, 24]. Based on the evidence from both FasL and FasR knockout mice that did not show any signs of cell death or growth deficiencies in any tissue outside the immune system, it was predicted that DICE preferentially affects cancer cells (especially CSCs) with little effect on normal cells [8, 24].

In spite of such optimistic results of experiments conducted on genetically modified animals [8, 24], mice injected with agonistic anti-FasR antibodies died after their INK and c-Jun had become phosphorylated [16]. With regard to all these observations, the clinical application of Fas-mediated procedures raise huge controversies because of major side effects, for instance massive apoptosis induced in the liver [25]. In our previous work, we found significant positive correlations between Fas expression and some CSC-like markers, what seemed to indicate the cancer progression promoting role of FasR/FasL signaling [26]. We demonstrated increased expansion of CD133+ cells within colonospheres following FasR stimulation, indicating specific clonal selection of cells sensitive to Fas-mediated supporting treatment. We also found that adherent HCT116 and HT29 CRC lines cells markedly decreased their proliferation rate not associated with increased apoptosis after during the incubation with anti-FasR agonistic antibody. That suggested the engagement of FasR in the senescence induction accompanied by the cell cycle arrest as was earlier presented by Raats et al. [27]. Additionally, these observations not necessarily were associated with particular genetic features of cancer cells (such as KRAS mutations). Fas signaling pathway was found to be engaged in ASA-mediated anti-cancerous effect. Chen et al. [28] described that ASA could induce apoptosis in Cox-independent way through p300-Ach3K9-FasL axis, which specifically targets CSCs. However, many crucial aspects concerning CSCs were left open. Unfortunately, authors did not mention nor evaluated any other possible effects which could be triggered by Fas pathway, including DICE. The literature and our previous data proved that thorough analysis of Fas signaling functions for CRC progression can be a source of very interesting discoveries, crucial for Fas associated procedures application. It seems to be extremely valuable, since CSCs were evidenced to be unusually sensitive to FasR/Fasl-based procedures [18, 29]. However, much needs to be learned before these new therapeutic options can be applied in human cancer.

References


