

PROSPECTS

Blocking Anti-Apoptosis as a Strategy for Cancer Chemotherapy: NF- κ B as a Target

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Abstract Critical processes underlying cancers must be better understood to develop strategies for treatment and prevention. A chemotherapeutic strategy is proposed that is based upon re-establishment, with a drug, of nullified programmed cell death (apoptosis) in cancer cells, which to survive have mutated to block apoptosis. A chemotherapy that is specific against tumors implanted in mice demonstrated the feasibility of this principle. This therapy is specific because it affects a process unique to cancer cells. It also has the advantage of killing these cells, in contrast to reversibly blocking their proliferation. The anti-apoptotic transcription factor NF- κ B provides a potential therapeutic target in estrogen receptor negative (ER⁻) breast cancers that over-express the epidermal growth factor family of receptors (EGFR). Further investigations of the pathways utilize dominant negative protein inhibitory peptide, and small inhibitory RNAs (siRNAs) to block the production of relevant enzymes. *J. Cell. Biochem.* 92: 646–650, 2004. © 2004 Wiley-Liss, Inc.

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Cell Biology

Both cell proliferation and apoptosis are important for cancer growth [Lin and Karin, 2003]. These processes are intertwined; aberrant proliferation activates tumor surveillance and apoptosis. Growth factors such as excess heregulin, transcription factor E2F1, oncogenic ras, and compounds such as retinoids can increase or decrease both proliferation and apoptosis. Cells die under conditions of stress, based upon pathways linked to the cell cycle [Vermeulen et al., 2003]. In cancers, the cells are metabolically unbalanced, and initially many die; survivors are mutants that developed anti-apoptotic properties [Weinstein, 2000]. An example is the very frequent loss of apoptotic p53; others are cancer-specific increases

in Bcl-2, the inhibitor-of-apoptosis protein survivin [Altieri, 2003], and increases of heat shock proteins [Nylandsted et al., 2000]. Overcoming anti-apoptosis can thus be the basis of therapy [Kim et al., 2002; Orłowski and Baldwin, 2002; Cusack, 2003a]. Compounds such as β -lapachone, geldanamycin, and Go6976 that selectively block anti-apoptosis (or reinstate apoptosis) selectively cause cancer cell death.

An initial demonstration of this principle of a drug which extirpates tumors in vivo is as follows. Mouse mammary epithelial carcinoma cells (CSMLO) subcutaneously implanted in syngeneic mice grew into large tumors which shrank dramatically after i.v. injection with the protein kinase C (PKC- α/β) inhibitor Go6976, a nonglycosidic indolcarbazol analog of staurosporine. The tumors did not reappear during extended treatment, and the action was tumor-specific because the mice exhibited no deleterious effects [Biswas et al., 2001, 2003]. As confirmation, CSMLO cells stably expressing a dominant negative mutant protein (dnIKK β) showed reduced and delayed tumor formation as compared to parental cells or cells plus vector alone.

The anti-apoptotic phenotype of breast cancer cells correlates with interaction of epidermal growth factor (EGF) with its over expressed

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receptor (EGFR+) in estrogen receptor negative (ER-) human and mouse adenocarcinoma cells in culture [Sovak et al., 1997; Biswas et al., 2000]. The DNA binding of anti-apoptotic NF- κ B is activated by EGF [Biswas et al., 2000]. Go6976 decreased NF- κ B activation, and the cells subsequently underwent apoptosis. Gene expression profiling using microarray analysis confirmed that inhibition of NF- κ B activation by Go6976 decreases the expression of anti-apoptotic genes and increases expression of genes associated with the apoptotic phenotype [Biswas et al., 2003]. Go6976 and dominant negative PKC- α block transformation of EGFR expressing cells [Hornia et al., 1999]. These experiments demonstrate the feasibility of rationally targeted therapies with the aim of activating apoptosis via the negation of an anti-apoptotic signaling pathway.

Molecular Mechanisms

Apoptosis and its several signaling pathways are often decreased in cancers [Zhivotovsky and Orrenius, 2003]. Of particular interest here is the increased NF- κ B found in cancer cells, which is reported to block apoptosis. This is especially influential for cancers that over express EGFR (ErbB1) or ErbB2, as compared to those that over express the ER. Serum primarily activates the MAPK pathway in ErbB1 over expressing cells, and the Akt pathway in ErbB2 over expressing cells [Tari and Lopez-Berestein, 2000]. These receptors and their downstream kinases are often over expressed in ER- breast tumors [Romieu-Mourez et al., 2002].

We demonstrated that DNA binding activity and transactivation of NF- κ B responsive genes can be stimulated by EGF-EGFR interaction in ER- human and mouse mammary epithelial adenocarcinoma cells in culture. Interaction of EGF with its receptor ErbB1 activates cell proliferation and also blocks death signals [Navolanic et al., 2003]. The signaling cascade includes PKC which is upstream of and activates I κ B kinase (IKK), a complex composed of three subunits IKK α , IKK β , and IKK γ /Nemo. NF- κ B is complexed with the inhibitory protein I κ B and maintained in an inactive state in most cell types. IKK phosphorylates I κ B which is then ubiquitinated and hydrolyzed by proteasomes. NF- κ B is then released to enter the nucleus, where it is further regulated by phosphorylation, acetylation, and interactions with coactivators and corepressors to transcribe

both anti-apoptotic and proliferative genes [Nakshatri and Goulet, 2002].

The NF- κ B/Rel transcription factor family regulate proliferation, survival, and transformation. They are heterodimeric complexes, of which the most predominantly detected is the p50/p65 (RelA) complex. NF- κ B is ubiquitously expressed and its activity is tightly controlled in normal cells. Its basal level can be activated by several signaling mechanisms, a variety of extracellular signals such as chemotherapeutic drugs, inflammatory cytokines, mitogens, growth factors, bacterial and viral infections, irradiation, and oxidative stress. It can transactivate over 200 target genes, involved in cellular processes that include anti-apoptosis (Bcl-2 and survivin), proliferation (cyclin D1), immune response, and adhesion/metastasis (uPA, integrin) [Lin and Karin, 2003].

Heterodimerized ErbB2/ErbB3 (Her2/neu), another EGF family receptor, is activated by heregulin [Stoica et al., 2003]. Over expression of ErbB2 is detected in about 30% of ER- human breast cancers. The prognosis of these patients is poor, although current therapies with Herceptin, an anti-Her2/neu antibody, in combination with other chemotherapeutic agents show some if limited success. ErbB2 activates NF- κ B via signaling that includes PI3-kinase, PDK1, Akt, protein kinase 2 (CK2) [Litchfield, 2003] and CKBBP1. Akt generates anti-apoptosis [Hill and Hemmings, 2002]. These pathways interconnect, in that PKC inhibits the Akt survival pathway [Tanaka et al., 2003], as do the ER α and Akt pathways [Stoica et al., 2003].

Tests of the Model

Our original hypothesis is that elevated NF- κ B in tumors provides a difference specific to cancer cells that suggests it as a target for drug therapy. NF- κ B is an attractive target because it often prevents apoptosis. Go6976 might nullify this anti-apoptosis by inhibiting PKC and thereby IKK activity; then I κ B is not phosphorylated nor degraded, NF- κ B activation is blocked and decreases because of its short half life. NF- κ Bs transcription of anti-apoptotic genes is thereby prevented, increasing apoptosis.

Consistent with our results, Go6976 inhibits PKC- α/β and enhances apoptosis, as do GF109203X and Ro318425 which are also selective inhibitors of PKC- α . They also enhanced

heregulin-induced apoptosis of SkBr3 cells that over express Her2. This action is via down regulation of Bcl-2 and activation of caspases-7 and -9, and poly-ADP-ribose polymerase cleavage [Le et al., 2002]. Unlike Go6976, several other PKC inhibitors did not protect neurons from cytotoxicity, suggesting Go6976 might alternatively block other enzymes [Jeohn et al., 2000].

We are currently exploring the effects of specific inhibition of NF- κ B activation in cancer cells by small inhibitory RNA (siRNA) targeted at the IKK complex in ER- human breast cancer cells in culture. This approach is in principle like the stable expression of dnIKK β mutant protein. Both are believed to block IKK β activity and should thereby down-regulate NF- κ B activation and permit apoptosis.

We have targeted two of the three proteins in the IKK complex. IKK β (IKK2) is a kinase which activates NF- κ B by phosphorylating the inhibitory protein I κ B α at serine residues 32 and 36, causing its dissociation from the p65/p50 complex (the primary NF- κ B heterodimer), and subsequent degradation by the 26S proteasome. IKK γ (Nemo) is the regulatory protein that is considered to control the kinase activities of both IKK α and β . In initial studies with the MDA-MB-231 ER- human mammary adenocarcinoma cell line we have successfully knocked down both IKK- β and IKK- γ with siRNAs. They could be reduced by greater than 85 and 98%, respectively. Surprisingly, we observed no significant changes in the level of active nuclear NF- κ B, growth characteristics, or apoptosis.

These observations suggests that other pathways are involved in the activation of NF- κ B. Other kinases can also phosphorylate I κ B [Bennett et al., 1996]. Highly purified PKC preparations did not phosphorylate I κ B α , whereas a crude preparation that contains contaminating CK2 did so [Janosch et al., 1996]. CK2 is constitutively active and it constitutively phosphorylates I κ B α in murine tumor cells. Inhibitors of CK2 induce apoptosis of Jurkat cells [Meggio and Pinna, 2003]. PI3-Kinase also directly interacts with I κ B α and contributes to the phosphorylation of p65, thereby by-passing the IKK complex [Kang et al., 2003]. Decreased NF- κ B does not always cause apoptosis, perhaps because of other survival mechanisms in some tumor cells [Orlowski and Baldwin, 2002]. We are currently investigating these pathways with respect to the constitutive

activity of NF- κ B in MDA-MB-231 and other cells, either alone or in synergy/conjunction with the IKK complex.

DISCUSSION

The primary goal of this study is to define a target-directed drug therapy for ER- human breast cancer patients, by blocking activation of NF- κ B. The rationale for choosing NF- κ B activation as a potential therapeutic target for ER- human breast cancers is based on its over expression in many cancers. Our results identified the importance of rationally targeted therapies with the aim of activating apoptosis via the inactivation of an anti-apoptotic signaling pathway, in particular leading to activation of NF- κ B.

More than 100 compounds including glucocorticoids, proteasome inhibitors, and PPAR γ antagonists decrease NF- κ B activity [Nakshatri and Goulet, 2002]. It should be mentioned that some inhibitors of NF- κ B are toxic due to effects independent of NF- κ B [Pozarowski et al., 2003]. Some of them might prove to be therapeutic or preventive [Lin and Karin, 2003]. For example, genistein, an isoflavone, inhibits the Akt pathway and the activation of NF- κ B; it is apoptotic [Sarkar and Li, 2002]. Many anticancer drugs cause apoptosis, but the increased NF- κ B which they also produce might prevent death. Synergies could exist between apoptotic agents and inactivators of NF- κ B [Weaver et al., 2003].

With this in mind, studies are in progress on cancer therapy with proteasome inhibitors [Kisselev and Goldberg, 2001] such as PS-341 (Velcade, Bortezomid), alone or in combination with other drugs [Voorhees et al., 2003; Cusack, 2003b]. One of the proposed mechanisms of action is based upon inhibition of I κ B proteolysis, with consequent decrease of active NF- κ B and activation of apoptosis. This principle has been demonstrated using a combination of lactacystin, a natural proteasome inhibitor, and doxorubicin which together demonstrated increased levels of apoptosis than when used alone [Tergaonkar et al., 2003]. These studies support the investigation of NF- κ B as a target for chemotherapy.

An important question that has arisen from these investigations suggests caution in applying siRNA in the discovery of new drug targets and it's role in deciphering the mechanisms

underlying the activity of new, supposedly specific, small molecule inhibitors. Go6976 can effectively block the stimulation of NF- κ B by EGF in the ER- setting, and more importantly demonstrated striking anti-tumor activity in vivo [Biswas et al., 2001, 2003]. Contrarily, specific inhibition of IKK β and γ by siRNA, without induction of cell death suggests either that the inhibition of PKC- α and - β by Go6976 has an effect on other pathways that culminate in NF- κ B inhibition, or alternatively also inhibits other kinases that, in conjunction with PKC inhibition, result in the observed anti-tumor effects. The complexity of drug effects and their low specificity [Orlowski and Baldwin, 2002] versus the high specificity of siRNA might be involved. Completeness of the effect depends on siRNA stability, time of exposure, variability of inhibition of IKK and NF- κ B between cell lines, extent of apoptosis signaling mechanisms, etc. Also, depleting a protein with siRNA may produce effects differing from inhibiting its catalytic activity. Apoptosis created from blocking NF- κ B activation may also depend on activating a pro-apoptotic signal. In conclusion, one needs to be cautious with siRNA with regard to predictions regarding the molecular pathways of drug actions.

Drugs are often recognized to modulate several processes. 7-hydroxystaurosporine (UCN-01), an anticancer drug, potently inhibits several kinases including PKC and PDK1 [Komander et al., 2003]. Go6976 is also a potent inhibitor of DNA damage-induced cell cycle checkpoints [Kohn et al., 2003]. Inhibition of NF- κ B transforms deacetylase inhibitor-induced growth arrest to apoptosis [Dai et al., 2003]. CK2 that both phosphorylates I κ B and activates cytoplasmic NF- κ B has more than 300 substrates [Meggio and Pinna, 2003]. Additional drug effects are usually deleterious, e.g., causing host toxicity. But rarely, a drug might modulate more than one desirable target, and could thereby create a synergistic effect. Such results were observed with microarray: Go6976 caused numerous changes of apoptotic genes' expression, which differed from those produced in dnIKK β -mutant cells [Biswas et al., 2003]. Such a "Drug Reinforcement" could be therapeutically beneficial.

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