

NF-κB Signaling Pathways in Carcinogenesis

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Abbreviations

Epidermal growth factor receptor ERB-B2 receptor tyrosine kinase 2
Phosphatidylinositol-4,5-
bisphosphate 3-kinase catalytic sub-
Zinc finger E-box-binding homeobox
Senescence-associated secretory

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Introduction

NF-κB in Tumor Initiation

The potential role of NF-κB in cancer was first investigated in hematologic malignancies, particularly multiple myeloma and leukemia [[1\]](#page-4-0). The focus subsequently shifted to solid tumors after discovery of its aberrant activity in breast cancers by others and us in 1998 [[2\]](#page-4-1). Aberrant NF-κB activity was subsequently reported in prostate, bladder, lung, head and neck, and pancreatic cancers. Mechanistic studies, however, revealed dichotomous role of NF-κB in cancer. NF-κB has been shown to promote senescence and function as a master regulator of senescenceassociated secretory phenotype (SASP) [[3\]](#page-4-2). Soluble factors in SASPs reinforce senescence arrest, alter the microenvironment in the tumor, and trigger immune surveillance. In genetically engineered Kras-induced mouse model of pancreatic cancer, the p65 subunit of NF-κB triggers CXCL1 (part of SASP)-/CXCR2-dependent senescence and inhibits initial steps of carcinogenesis [\[4](#page-4-3)]. DNA damage also triggers NF-κBdependent senescence. Switch to oncogenic function occurs once cells are immortalized. Other genomic aberrations in cancer cells will also determine tumor suppressor function of NF-κB. For example, in tumors where the expression of the anti-apoptotic BCL-2 is dependent on NF-κB, it functions as a survival factor. By

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S. Badve, G. L. Kumar (eds.), *Predictive Biomarkers in Oncology*, https://doi.org/10.1007/978-3-319-95228-4_27

contrast, in tumors with elevated BCL-2 independent of NF-κB, therapy-induced cell death requires NF-κB-mediated induction of SASPs [\[5](#page-4-4)]. In general, senescence is a double-edged sword in cancer because while senescent cells themselves rarely progress into cancer, SASPs from these cells can promote neoplastic progression of nearby preneoplastic cells by providing pro-inflammatory molecules and inducing epithelial-to-mesenchymal transition. Therefore, timing of NF-κB-induced senescence could have profound influence on cancer development. From the practical and clinical angle, this also poses challenge to ascertain whether concurrent presence of activated NF-κB (as determined by nuclear p65 and phosphorylated p65) and SASP is a good or bad prognostic marker.

NF-κB in Tumor Progression/ Metastasis

Overcoming senescence barrier either through inactivation of cell cycle inhibitors such as p16 or overexpression of cell cycle protein cyclin D1 and inactivation of p53 or through telomerase overexpression leads to cellular immortalization. In immortalized cells, NF-κB cooperates with other oncogenes such as RAS or functions downstream of oncogenes such as KRAS, mutant PIK3CA, activated EGFR, or ERBB2 to promote cancer progression. Cancer-promoting functions include upregulation of anti-apoptotic proteins such as BCL-2, BCL-XL, GADD45β, XIAP, cIAP1, and cIAP2; cell cycle proteins such as Cyclin D1; pro-invasion molecules such as MMP9; pro-metastatic molecules such as CXCR4; pro-inflammatory molecules such as IL-1α, IL-1β, IL-6, IL-8, TNFα, and COX2 and metabolic pathway genes such as GLUT3; and genes linked to epithelial-to-mesenchymal transition such as $ZEB1$ and $ZEB2$ [[2,](#page-4-1) [6\]](#page-4-5). Additionally, there is evidence for NF-κB playing a significant role in self-renewal and maintenance of cancer stem cell phenotype, particularly in glioblastoma. NF-κB is essential for lung tumor development upon p53 mutation and KRAS (G12D) expression [[7\]](#page-4-6). Inhibition of NF-κB sensitizes EGFR-

mutant lung tumors to EGFR-targeted therapies. MLL fusion proteins that typically cause leukemia are dependent on NF-κB for transformation. NF-κB is also a central player in epigenetic switch that links chronic inflammation to cell transformation and subsequent metastasis [\[8](#page-4-7)].

Cancer Cell Non-autonomous Functions of NF-κB

Although initial studies on NF-κB were primarily focused on cell autonomous functions, several recent reports highlight its non-cell autonomous roles. NF-κB-inducible cytokines such as IL-6 and IL-8 not only alter tumor microenvironment by attracting different immune cells, but also these cytokines are associated with systemic effects of cancer. IL-6 is a major contributor to cancer cachexia. NF-κB itself contributes to cancer cachexia by blocking myogenic differentiation by affecting skeletal muscle transcription factor-microRNA circuitry and by repressing differentiation factor MyoD [\[9](#page-4-8)].

Recent studies have demonstrated a role for NF-κB in resistance to immune therapy. PD-L1, expressed mainly by cancer cells, plays a significant role in creating antitumor immunity. Therefore, several antibodies targeting PD-L1 have entered clinic. Although correlation between PD-L1 levels and response to therapy is yet to be established, NF-κB has been shown to increase PD-L1 at both transcriptional and posttranscriptional level. Pro-inflammatory cytokines such as TNF α induce the expression of COP9 signalosome 5 (CSN5), which deubiquitinates PD-L1 and stabilizes the protein.

By increasing PD-L1 protein, NF-κB could reduce the effectiveness of PD-L1-targeting antibodies (Fig. [27.1\)](#page-2-0) [[10\]](#page-4-9).

Current NF-κB-Targeted Therapies and Future Strategies for Treatment

Based on multiple functions ascribed to NF-κB in cancer and therapeutically targetable signaling cascades involved in NF-κB activation,

Fig. 27.1 Canonical and noncanonical pathways of NF-κB activation. Steps associated with NF-κB activation in response to extracellular signals are shown. Different members of NF-κB and IκB family are listed. In mammals, the NF-κB family is composed of five related transcription factors: p50, p52, RelA (p65), c-Rel, and RelB. IKK complex is the main signaling hub that integrates extracellular/membrane-activated signals to nuclear events by activating NF-κB. Potential therapeutic agents

there have been several attempts to develop drugs targeting NF-κB [\[11\]](#page-4-10). Commonly used drugs such as aspirin and food ingredient curcumin inhibit NF-κB. Several groups including us have shown anti-NF-κB activity of parthenolide, an active ingredient in the herb *Tanacetum parthenium* or feverfew used for migraine. Clinically, bortezomib used to treat multiple myeloma is a proteasome inhibitor that prevents IκB degradation and restricts NF-κB activation. However, for most of these drugs, NF-κB is one of their targets, and their clinical activity cannot be solely due to NF-κB inhibition. A highly specific NF-κB inhibitor is yet to enter clinic, although such an inhibitor may never be discovered. An ideal NF-κB inhibitor should dampen overactive NF-κB rather than completely eliminate its activity because basal NF-κB is necessary for normal function of the immune system and to prevent infection. Therefore, there is still considerable interest in

that may reduce NF-κB activation or reduce the activities of downstream targets of NF-κB are also indicated. For example, ABX-IL8 may inhibit IL8-dependent signaling, and Plerixafar prevents CXCR4 activation by inhibiting CXCR4/CXCL12 interaction. Infliximab is a TNFα antagonist, which can block NF- κ B activation by TNF α as well as block the effects of TNFα downstream of NF-κB activation

both academia and industry to develop drugs that reduce but not eliminate NF-κB activity. Because of the role of NF-κB in upregulating anti-apoptotic proteins, these inhibitors will likely work as chemosensitizing agents rather than displaying single agent activity. However, chemosensitizing function NF-κB inhibitors needs to be tested with individual chemotherapeutic drugs because in certain instances, therapy-induced NF-κB, particularly therapies that promote replication stress, could augment cell death by activating extrinsic pathway of cell death through expression/ activation of FAS-FASL-dependent cell death machinery [[12\]](#page-4-11).

Several of downstream targets of NF-κB can be targeted therapeutically (Table [27.1](#page-3-0)). In fact, drugs targeting TNF, IL-1, IL-6, IL-8, CXCR4, and PD-L1 are already in clinical use and can be exploited to treat cancers in which NF-κB pathway is essential for cancer cell survival.

Summary

Although NF-κB was discovered more than 30 years ago, its regulation and function still remain at the forefront of research not only from the oncology point of view but also with other diseases such as autoimmune disorders, microbial infections, and neurodegenerative diseases. While constitutive activation of NF-κB has been reported in advanced stages of many cancers, causes of this activation vary widely and may involve both genomic and nongenomic events. Despite early-stage-specific tumor suppressor role described for this transcription factor complex, this observation should not hinder clinical development of NF-κB inhibitors because NF-κB has already transitioned to oncogenic role at the time of clinical manifestation of the disease. Thus, studies focusing on NF-κB, both at regulatory and functional level, will continue to unlock mysteries surrounding cancer progression and potentially to new cancer therapies. Reliable biomarkers of constitutive NF-κB activity in cancer are yet to be identified because of close link between NF-κB activation and inflammatory process. Although drugs that directly target NF-κB are yet to enter clinic, several drugs that inhibit the function of proteins overexpressed as a consequence of increased NF-κB activity in cancer are showing promising results.

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