Targeting constitutive NF-κB specifically in tumor cells

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NF-κB transcription factors are the key regulators of innate and adaptive immune responses, inflammation, and cell survival [1]. Considerable evidences indicate that NF-κB plays important roles in carcinogenesis, cancer progression, metastasis and drug resistance, and therefore NF-κB has been regarded as one of the most important targets for cancer therapy [2]. However, the application of inhibitors that target NF-κB or NF-κB pathways for the treatment of human cancer is impeded by severe side effects due to the indiscriminate inhibition of NF-κB in normal immune cells.

Traditionally, there are two NF-κB activation pathways that play important roles in immune responses. The first pathway, also called canonical or classical NF-κB activation pathway, is triggered in response to microbial and viral infections and exposure to proinflammatory cytokines that induce activation of the tripartite IKK complex and the consequent phosphorylation-related degradation of IκBs, leading to the nuclear translocation of NF-κB complexes, predominantly p50/RelA and p50/c-Rel dimmers [3]. The second pathway, also called non-canonical or alternative NF-κB activation pathway, involves different signaling molecules and leads to the predominant activation of p52:RelB NF-κB dimers by inducing processing NF-κB2/p100 precursor that binds to the RelB NF-κB subunit in the cytoplasm [4].

As more and more reports suggest that NF-κB is activated in most cancer cell lines and tumor cells, especially in those tumor cells that resist to chemo- or radiotherapy, it becomes a consensus for the research community that NF-κB plays a major role in tumor development and cancer therapy-resistance. However, we seemed ignore the differences between NF-κB (acute) activation in immune responses and NF-κB constitutive activation in tumor cells, and also ignore or sometimes confuse the differences between the activation of NF-κB pathways and the activation of NF-κB itself. In fact, as demonstrated in both transgenic/knockout and allograft/xenograft mouse models in our studies [5, 6], the constitutive activation of NF-κB in tumor cells is different from NF-κB (acute) activation in immune cells, and the constitutive activation of NF-κB in tumor cells doesn’t dependent on traditional NF-κB pathways, which are normally related to the activation of IKKβ or IKK complex [5]. This can explain why the IKKβ/IKK inhibitors are not as powerful for the inhibition of tumor cell proliferation and tumor growth as expected or as what theoretically they should do.

Importantly, our studies have shown that it is the constitutive NF-κB activation that drives tumor prostate cancer recurrence and therapy-resistance. And the constitutive NF-κB activation is established and maintained by a feed-forward signaling circuit composed of 1xBα/NF-κB(p65), miR-196b-3p, Meis2, PPP3CC, where the constitutive 1xBα/NF-κB(p65) drives the expression of miR-196b-3p that inhibits the expression of Meis2 and PPP3CC, down-regulated PPP3CC is unable to dephosphorylates p-IκBα, leading to constitutive IkBα phosphorylation and NF-κB(p65) activation (Figure 1). Therefore, targeting this constitutive NF-κB signaling circuit by interrupting its non-IκBα/ NF-κB(p65) components would be a highly efficient way for the treatment of recurrent and therapy-resistant prostate cancer, which will avoid the side effects related to indiscriminate IKK/NF-κB inhibition in normal cells.

The constitutive NF-κB signaling circuit controls the expression of a group of stem cell transcription factors, including Twist2, Sox2, Oct4, and Nanog, that drives the

![Diagram of constitutively activated NF-κB signaling circuit in recurrent or therapy-resistant prostate tumor.](https://example.com/diagram.png)
proliferation and self-renewal of cancer stem cells, leading to tumor recurrence and cancer therapy-resistance [5]. Although these stem cell transcription factors may not be the downstream targets of the individual non-IκBa/NF-κB(p65) components of the circuit, disrupting this circuit by targeting any of the non-IκBa/NF-κB(p65) individual components blocks the expression of these transcription factors and significantly impairs castration-resistant prostate cancer (CRPC) development [5]. Therefore, disrupting this constitutive NF-κB signaling circuit by targeting any of its individual non-IκBa/NF-κB(p65) components would be as powerful as direct inhibition of NF-κB for the treatment of CRPC.

The non-IκBa/NF-κB(p65) components in constitutive NF-κB signaling circuit include miR-196b-3p, Meis2, and PPP3CC (Protein Phosphatase 3, Catalytic Subunit, Gamma Isozyme). Further studies are needed to distinguish the best suitable one among these three non-IκBa/NF-κB(p65) components for targeting constitutive NF-κB signaling in tumor cells. It should be mentioned that the phenomenon and the role of constitutive NF-κB signaling in tumor recurrence and therapy-resistance were found in CRPC cells, further studies are needed to define if the constitutive NF-κB signaling circuit also exists in other types of recurrent and therapy-resistant cancer cells. We predict that the constitutive NF-κB signaling may be a common phenomenon in most recurrent and therapy-resistant cancer cells, however, the components of the constitutive NF-κB signaling may be different in different types of tumors, as evidenced in our previous studies on breast cancer [7].