NLRC3 mediated PI3K-mTOR inhibition takes a toll on colon cancer

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NLRs (nucleotide-binding domain and leucine-rich repeats containing receptors) are pattern recognition receptors, evolutionarily conserved across plants and animals (1). NLRs, initiate immune response against specific damage and pathogen-associated molecular patterns (DAMPs/PAMPs) (2). NLRs also contribute to the function and regulation of multiple innate immune signaling pathways (3) (Figure 1). NLRC3, the newly characterized NLR family member is preferentially expressed in immune cells. NLRC3 performs various innate immune functions such as regulation of several inflammatory signaling pathways (4). NLRC3 attenuates TLR (TOLL-like receptor) signaling via inhibition of MyD88 and TRAF6-dependent activation of NF-κB pathway. In fact, NLRC3 interacts with TRAF6 to promote its degradation (5). According to Gultekin et al., NLRC3 is expressed as a cytosolic protein and promotes negative regulation of inflammasome signaling. Inflammasome represents a multi-protein platform comprising of a NLR family member, apoptosis-associated speck-like protein containing CARD (ASC) and pro-caspase-1, a cysteine protease enzyme (2). NLRC3 specifically inhibits NLRP3-induced ASC speck formation, and subsequent activation of pro-caspase-1 and IL-1β production (6) (Figure 1). Furthermore, NLRC3 interacts with stimulator of interferon genes (STING) to modulate host response towards intracellular DNA, DNA virus, and cyclic di-GMP. NLRC3 impedes STING-TBK1 interaction, reducing STING-dependent innate immune activation and downstream type I interferon production (7). The STING pathway mediates recognition of cytosolic DNA by dendritic cells, generating spontaneous T cell responses against immunogenic tumors (8). However, study by Corrales et al., highlights another major pathway triggered by tumor-derived DNA that activates AIM2 inflammasome. The AIM2 inflammasome activation inhibits STING pathway, by promoting caspase-1 mediated cell death (9). In summary, these findings highlight the complex regulation of inflammasomes, and their association with other innate immune signaling pathways.

Dysregulation of NLR signaling is central to several inflammation-associated diseases including cancer (3). NLR signaling plays a critical role in cancer initiation, development and progression. Moreover, NLRs perform both pro- and anti-tumor regulatory roles in cancer (10). Specifically, colorectal cancer presents disease burden worldwide, and is the third leading cause of cancer death in U.S (11). The dramatic increase in colon cancer incidence and mortality rate, has raised attempts towards potential targets identification, to reduce cancer risk, promote preventive diagnosis and effective treatment strategy. Scientists have identified the novel association of NLR family members, NLRP6 and NLRP12 with tumorigenesis in colorectal cancer (12). Both NLRP6 and NLRP12 are non-inflammasome-forming NLRs and function as negative regulators of inflammation signaling. The Nlrp6−/− and Nlrp12−/− mice showed increased susceptibility towards...
Figure 1 NLRC3 regulates several inflammation-associated pathways. NLRC3 inhibits LPS-induced TL4 signaling and NF-κB activation, via increased TRAF6 degradation. NLRC3 inhibits NLRP3 inflammasome formation, and subsequent IL-1β and IL-18 cytokine production. NLRC3 sensing also inhibits T-cell activation dependent IL-2 release and HSV1, c-di-GMP and dsDNA induced type-I-interferon production.

Inflammation-driven colon tumorigenesis. Importantly, Nlrp6 and Nlrp12 signaling in the hematopoietic cells is critical for protection against colitis and colitis-associated tumorigenesis (13-15). Emerging data provides strong evidence for NLRs-mediated protection in colorectal cancer, and suggest interplay between NLRs and inflammation-associated molecules during colitis and colitis-associated tumorigenesis.

Recently, Karki and Man et al., identified NLRC3 as a negative regulator of PI3K-mTOR pathway in colorectal cancer (16). NLRC3 expression was significantly higher in the colon tumor tissue as compared to the non-tumor/normal. The study utilized azoxymethane (AOM)-dextran sulfate sodium (DSS) model for colorectal tumorigenesis. The reduced body weight and colon length, and increased dysplasia and adenocarcinoma in the AOM-DSS treated Nlr3-/- mice, signifies the protective role of NLRC3 in colitis-associated colorectal cancer. To assess the contribution of any gene-deletion induced microbiota alterations, the mice were co-housed during the course of experiments. However, results remained unchanged in the co-housed and non-co-housed experimental groups confirming the lack of microbiota contribution. Interestingly, the Nlr3-/- deficient mice when treated with AOM-DSS had significantly increased weight loss, tumor size and tumor number per colon as compared to the wild type controls. Histopathological analysis further revealed higher colon inflammation, ulceration, hyperplasia and extent of severity/damage in the Nlr3-/- mice as compared to wild-type mice (16). Based on the preliminary
observations, NLRC3 emerges as a potential innate immune regulator for protection against colorectal tumorigenesis.

To elucidate cell-specific regulation of NLRC3 during colorectal tumorigenesis, Karki and Man et al. performed bone-marrow chimera experiments (16). Nlrc3−/− mice showed significant increase in tumor size and number as compared to wild-type mice, after wild-type bone marrow transplant. Similarly, wild-type mice when transplanted with Nlrc3−/− bone marrow had significantly increased tumor size with respect to wild-type mice that received wild-type bone marrow. Notably, Nlrc3−/− mice, with Nlrc3−/− bone marrow showed highest tumor load compared to the other transplants (16). Previous studies demonstrated cell-specific role of inflammasome-forming NLRs, NLRP3 and NLRC4 in colon cancer (17,18). Both Nlrp3−/− and Nlrp4−/− mice were highly susceptible to DSS-induced colitis and suffered loss of epithelial integrity during colitis-associated tumorigenesis. The authors further confirmed these results by generating mice lacking NLRC3 in either hematopoietic or myeloid or intestinal epithelial cells. The Nlrc3−/− mice showed higher tumor development, followed by the mice lacking Nlrp3 in hematopoietic cells. Hence, the inhibitory effect of NLRC3 on tumor growth is significant in epithelial cells as compared to the hematopoietic cells.

In their work, Karki and Man et al. investigated the effect of NLRC3 on cell growth and proliferation in the AOM-DSS treated mice. Interestingly, Nlrc3−/− mice showed increased cellular proliferation in intestinal epithelial cells, which was further confirmed by increased organoids formation by the colonic epithelial stem cells and primary fibroblasts (16). The mammalian target of rapamycin (mTOR), acts downstream of phosphoinositide 3-kinase (PI3K)/Akt pathway, and its activation has been implicated in cell metabolism, proliferation, growth, migration including colon tumorigenesis (19,20). Therefore, Karki and Man et al. compared the phosphorylation levels of mTOR and mTOR-associated signaling proteins in the colon tissue of AOM-DSS treated mice. As expected, the colon of Nlrc3−/− mice at day-4 had increased phosphorylation levels of S6 kinase, 4E-BP1 and AKT at Ser473, the downstream targets of mTOR, as compared to the wild-type mice. Immunoblotting and immunofluorescence analysis showed high levels of phosphorylated AKT, total AKT, phosphorylated 4EBP1 and GAPDH in the colon tissue of Wild type, Nlrc3−/− and Nlrc3−/− mice (day-14) and in the IGF-1-treated primary fibroblasts (16). The data shows early dysregulation of mTOR in the Nlrc3−/− mice, that occurs just 8 days after injection of azoxymethane. These preliminary findings implicate NLRC3 as potent regulator of the mTOR signaling pathway.

The overactivation of Wnt-signaling promotes colorectal cancer invasion, metastasis and metabolism (21). However, Karki and Man et al. didn’t find any difference in expression of genes involved in Wnt-signaling pathways. No significant difference in the production of inflammatory mediators and immune cell infiltration was observed in the Nlrc3−/− mice at day-8. The data clearly indicates absence of NF-κB activation at early stage of tumorigenesis. Surprisingly, the pro-inflammatory mediators, IL-1β, IL-6, TNF-α, GCSF, KC, MCP-1 and MIP-1α in the colon tissue, had significantly increased relative gene and protein expression levels in treated Nlrc3−/− mice at day-14 (16). Therefore, the dysregulation of mTOR signaling was followed by NF-κB signaling activation at a later stage in tumorigenesis. Additionally, the expression of IL-17 and IL-22 was elevated in the colon tissue of Nlrc3−/− mice whereas the expression of IL-23, IFNβ and IFNγ remained unchanged at day-14. As expected, the phosphorylation levels of Iκ-Bα and STAT3 were found to be increased in the Nlrc3−/− mice as compared to the wild-type mice at day-14. Consistent with the observations that Nlrc3−/− mice had increased expression of pro-inflammatory mediators, the authors observed significantly higher number of macrophages, CD11b+ CD11c+ cells, neutrophils, and NK cells, in the colon tissue of Nlrc3−/− mice at day-14 (16). The results demonstrate NLRC3-controlled reduction in intestinal inflammation and immune cell infiltration during colorectal tumorigenesis. Several upstream signaling molecules regulate the phosphorylation and activation of mTOR in the PI3K–Akt–mTOR pathway (22). The phosphorylation of Akt at the Thr308 site by the kinase PDK1 activates the mTOR regulated pathways (23). The authors found increased phosphorylation of AKT at Thr308 site in the Nlrc3−/−, followed by the heterozygous Nlrc3−/+ with respect to wild-type. Therefore, NLRC3 suppresses colon tumorigenesis in gene-dose dependent manner, via PI3K-mTOR-AKT pathway response.

The authors further evaluated NLRC3-mediated protection mechanism against colorectal cancer, using the Apcmin/+ mice, a spontaneous model of colon cancer. They found increased cellular proliferation, dysplasia and tumor burden in the colon of Apcmin/+ Nlrc3−/− mice. Moreover, the colon of Apcmin/+ Nlrc3−/− mice, showed decreased proliferation, inflammation damage and tumor load in the presence of mTOR/PI3K/mTOR-PI3K inhibitors (16). All above investigations present NLRC3 as a negative regulator
of mTOR to prevent colorectal cancer progression. NLRC3 interactions with the PDK1 and p85 subunit of PI3K were confirmed by co-immunoprecipitation assays, in the IGF-1 activated 

\[ \text{Nlrc3}^{-/-} \] primary fibroblasts/bone-marrow-derived macrophages (BMDMs). Increased phosphorylation and activation of p85 subunit was observed in the colon tissue of AOM-DSS treated 

\[ \text{Nlrc3}^{-/-} \] mice as compared to the wild-type. Collectively, the findings demonstrate NLRC3-mediated disruption of p85-p110\( \alpha \) PI3K subunits interactions, by direct interaction with the p85 subunit. Researchers also found increased activation of mTOR signaling, in response to LPS-induced TLR4 activation, in the 

\[ \text{Nlrc3}^{-/-} \] primary BMDMs and 

\[ \text{Nlrc3ld/ld} \] mice \((\text{ld}-\text{large deletion})\) (16). Here, the 

\[ \text{Nlrc3ld/ld} \] mice represent the independently generated NLRC3-deficient mice in the laboratory.

Recent findings by Karki and Man et al., highlights the protective role of NLRC3 against colorectal cancer, via inhibition of the PI3K-mTOR-AKT signaling pathway. The results explain the molecular mechanism of NLRC3 mediated inhibition of cellular proliferation, inflammation and immune cell infiltration in colon cancer (16). The emerging evidences suggest cross-talk between NLRC3 and major intracellular signaling pathways, involved in cell growth, proliferation and inflammation. NLRC3 is a negative regulator of inflammation and it would be interesting to find the association of NLRC3 with inflammasome-forming NLRs, during colorectal tumorigenesis. Several new questions arise from these findings: (I) How is NLRC3 expression in epithelial cells but not hematopoietic cells critical to tumorigenesis? (II) How does NLRC3-mediated signaling vary across cell lineages? (III) Is NLRC3 expression variable or is the difference in signaling arising from NLRC3-interacting signaling proteins? From the preferential expression of NLRC3 in innate immune cells, it becomes important to further understand the structural and activation mechanism of NLRC3 and its downstream target molecules. These questions and many more arising from this seminal work will fuel future research in comprehending the cellular and molecular pathways underlying NLRC3-mediated protection against colorectal cancer.

Inflammation and innate immune cell infiltration are major components of the tumor microenvironment. The significance of NLRs in generation of host innate immune responses has led to the development of NLR-targeted drugs against inflammation-associated diseases and cancer (24,25). The newly identified role of NLRC3 in protection against colorectal cancer has generated a lot of curiosity regarding the expression and function of NLRC3 in other cancers. NLRC3, similar to its other family members, presents a promising option for development of NLR-targeted strategies, to foster host immune responses against the tumor-microenvironment.

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**Footnote**

*Conflicts of Interest:* The authors have no conflicts of interest to declare.

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