RB goes mitochondrial

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The retinoblastoma tumor suppressor RB is well known for its capacity to restrict cell cycle progression at the G1/S transition of the cell cycle by controlling the transcription of cell cycle genes. In this issue of Genes & Development, Hilgendorf and colleagues (pp. 1003–1015) have identified a novel tumor suppressor function for RB independent of its role as a transcriptional regulator, in which RB directly activates the apoptosis regulator Bax at the mitochondria to promote cell death.

Early paradoxical observations on RB and cell death

The cellular mechanisms of tumor suppression include cell cycle arrest, senescence, differentiation, and cell death. When the RB tumor suppressor was first extensively studied in the 1990s, its role as an inhibitor of cell cycle progression was rapidly illuminated. Subsequent studies then indicated that active RB molecules could also contribute to the establishment and maintenance of senescence as well as to promoting differentiation. At the molecular level, these studies identified RB as a transcriptional regulator with the capacity to bind transcription factors such as E2F and recruit chromatin modifiers to the promoters of target genes involved in cell proliferation and differentiation (for review, see Harbour and Dean 2000; Burkhart and Sage 2008; Chinnam and Goodrich 2011).

The connection between RB and cell death was initially a bit surprising. Indeed, induction of cell death was primarily observed as a consequence of the absence of RB or its inactivation in cells. For instance, apoptotic cell death was found to be prominent in mouse embryos deficient for the Rb gene; genetic crosses further demonstrated that E2F and p53 are crucial for the induction of cell death downstream from RB loss (Morgenbesser et al. 1994; Symonds et al. 1994; Macleod et al. 1996; Zacksenhaus et al. 1996; Pan et al. 1998; Tsai et al. 1998; Ziebold et al. 2001). Key downstream targets of E2F in this process are the p53 family member p73 as well as genes encoding cell death machinery components such as Caspases, Apaf-1, and Bcl-2 homology 3 (BH3)-only

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proapoptotic members of the Bcl-2 protein family (Irwin et al. 2000; Moroni et al. 2001; Nahle et al. 2002; Hershko and Ginsberg 2004). RB may also directly bind to and inhibit proapoptotic molecules in some contexts (Adegbola and Pasternack 2005).

These observations led to a model in which cancer initiation following loss of RB—as in retinoblastoma—must happen in a cell type that is inherently resistant to the death resulting from loss of RB function (Chen et al. 2004). These studies also provided a basis for the frequent coalteration of the RB and p53 pathways in cancer cells, as RB pathway mutant tumor cells likely need to abrogate p53 proapoptotic function in order to expand during tumor progression (Sherr and McCormick 2002).

Proapoptotic role for RB

Concomitantly with these studies demonstrating that loss of RB leads to increased cell death, an increasing number of reports have shown a proapoptotic role for RB. For example, studies from the Wang laboratory (Chau et al. 2002) showed that RB is cleaved by Caspases in response to TNF α and that preventing this cleavage reduces the amount of death downstream from the TNF α receptor (but not in response to DNA damage). These data could suggest that either RB needs to be inactivated for death to be induced efficiently, in keeping with the antiapoptotic role of RB described above, or alternatively, cleaved RB is proapoptotic.

Since these initial studies, evidence for a direct role for RB in promoting cell death has accumulated. For example, in Caenorhabditis elegans, genetic crosses have shown that the RB homolog Lin-35 collaborates with E2F to promote apoptosis during development of the anterior pharynx (Reddien et al. 2007). Lin-35 also promotes germ cell apoptosis in part by repressing the expression of ced-9, an ortholog of the anti-apoptotic factor BCL2 (Schertel and Conradt 2007). In addition, in mammalian cells, expression of a form of RB that cannot be phosphorylated on its major CDK phosphorylation sites can stimulate the apoptotic response to $TNF\alpha$ (Masselli and Wang 2006). RB also plays a role in the cell death induced by histone deacetylase inhibitors in colon cancer cell lines (Wagner and Roemer 2005). Furthermore, efficient induction of death by TGFB in some cancer cells requires a complex between RB, E2F-1, and the acetyltransferase P/CAF (Korah et al. 2012). Finally,

a previous report from the Lees laboratory (Ianari et al. 2009) further showed that the phosphorylated forms of RB found in proliferative cells and normally less able to bind to E2F-1 can be induced to interact with E2F-1 in response to genotoxic stress (DNA damage) or oncogenic stress (expression of E1A). In the same study, stressinduced RB/E2F-1 complexes containing P/CAF or p300 were shown to bind to and activate the transcription of cell death genes such as p73 and Casp7 (Ianari et al. 2009).

A way to reconcile these different studies is that RB has both anti-apoptotic and proapoptotic roles, depending on the cellular context: In normally slowly proliferating or quiescent cell populations, acute loss of RB function (by deletion but also by rapid hyperphosphorylation) can induce cell death in part by deregulated activity of E2F and activation of p53. This proapoptotic consequence of RB loss may, for example, help prevent cancer initiation from quiescent populations of stem cells. In cells that are already proliferating, RB molecules (including phosphorylated molecules usually thought to be ''inactive'') may be able to sense genotoxic or oncogenic stresses and complex specifically with E2F-1 to activate the expression of cell death genes and promote apoptosis (Fig. 1). While the detailed mechanisms underlying the formation and activation of RB/E2F-1 proapoptotic transcriptional complexes are not fully established, the specific interaction between RB and E2F-1 relies on the C-terminal domain of RB and post-translational modifications of both proteins (Dick and Dyson 2003; Rubin et al. 2005; Julian et al. 2008; Carnevale et al. 2012). This proapoptotic stress sensor function of RB may provide a safety mechanism to eliminate cells that have committed to

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divide but may carry genomic alterations due to replication stress.

Direct activation of the apoptotic machinery by RB at the mitochondria

These previous studies building a case for a proapoptotic role for RB have often focused only on mechanisms linked to the transcriptional regulation by RB of genes involved in cell death. However, the fastest way to trigger cell death in cells is independent of the transcriptional machinery and involves changes in the membrane potential of mitochondria in cells (Wang 2001). In this issue of Genes & Development, Hilgendorf et al. (2013) provide compelling evidence that a fraction of RB is at the mitochondria, where it can activate Bax and promote cell death in response to $TNF\alpha$ treatment or other stresses. First, the investigators showed that RB can promote $TNF\alpha$ -induced apoptotic cell death in a translation-independent manner in RAT16 cells using treatment with cycloheximide to inhibit translation. The investigators then showed that this effect is dependent on the Bax gene and not Bak using immortalized mouse embryonic fibroblasts deficient for each of these two proapoptotic genes. Intrigued by these observations, Hilgendorf et al. (2013) sought to determine whether RB could localize to the mitochondria, where Bax normally resides. Indeed, they found that a fraction of the RB pool is constitutively mitochondrial in IMR90 human cells and mouse liver cells, including phosphorylated forms of RB. RB can directly interact with Bax, conformationally activate it, and trigger mitochondrial outer membrane permeabilization (MOMP)

> Figure 1. Anti- and proapoptotic functions of the RB transcriptional regulator. (Top) During normal cell cycle progression, RB phosphorylation by Cyclin/CDK complexes results in the dissociation of RB and E2F transcription factors and activation of E2F target genes. RB's ability to restrict cell cycle progression is thought to be one of its major tumorsuppressive roles in cells. (Bottom left) When RB is inappropriately inactivated in quiescent cells or cells exiting the cell cycle—for example, by gene mutation, protein degradation, or hyperphosphorylation—activation of E2F in the wrong context may result in the activation of cell death genes (in addition to cell cycle genes), which may help suppress cancer initiation. (Bottom right) In response to certain stresses (e.g., DNA damage or oncogene activation), RB can specifically bind to E2F-1 even in cycling cells and promote the expression of cell death genes, thereby thwarting the expansion of abnormal cells and suppressing cancer development.

and release of cytochrome c , an essential step in the initiation of apoptosis. Furthermore, a truncated form of RB containing only its ''small pocket'' (a domain known to be essential for RB's tumor suppressor function) is sufficient to promote cell death induced by $TNF\alpha$ treatment in the presence of cycloheximide. Strikingly, an RB mutant lacking the nuclear localization signal but carrying a mitochondrial import signal was found to be sufficient to drive apoptosis. The remarkable ability of this mutant to suppress the growth of $Rb^{-/-}$; $p53^{-/-}$ osteosarcoma tumor cells to an extent comparable with nuclear RB suggests that the proapoptotic action of RB at the mitochondria may be as critical as its transcriptional tumor suppressor role, at least in certain contexts, and further demonstrates that this proapoptotic role of RB is p53-independent.

Similarities and differences between RB and p53

This study from the Lees laboratory (Hilgendorf et al. 2013) recalls that of RB's fellow tumor suppressor, the p53 transcription factor (Fig. 2; Brady and Attardi 2010), which, in startling findings in 1994, was discovered to induce apoptosis in the presence of transcription and translation inhibitors (Caelles et al. 1994). The story took

Figure 2. Similarities between RB and p53 in the induction of apoptotic cell death. Both p53 (directly) and RB (with E2F1 or via E2F activity) can increase the transcription of proapoptotic genes in the nucleus. Only some of the known targets are shown, with a focus on genes encoding key apoptotic regulators and common targets between p53 and E2F. Components or regulators of the apoptotic machinery at the mitochondria are highlighted in a box. Note that Puma can disrupt the interaction between p53 and Bcl-XL, thereby allowing p53 to activate Bax and Bak. The new study from Hilgendorf et al. (2013) indicates that RB can also directly control Bax at the mitochondria, similarly to p53, which can activate Bax and Bak at the mitochondrial membrane to promote apoptotic cell death.

an even more surprising twist in 2000, when p53 was reported to localize to mitochondria during apoptosis, where it acted directly to promote apoptosis (Marchenko et al. 2000). While greeted with skepticism initially, mounting evidence from different laboratories buttressed the notion of p53 action at the mitochondria, albeit through different mechanisms according to the specific model system examined. The common theme is that p53 interacts with Bcl-2 family members in each case, but the specific players vary. In the original study, $p53$ was found to interact with Bcl2 or Bcl- X_L at the mitochondria to neutralize their activity and thus activate proapoptotic Bax and Bak proteins to drive MOMP (Mihara et al. 2003). Interestingly, tumor-derived p53 mutants fail to interact with Bcl-2 or Bcl- X_L , suggesting the importance of these interactions for p53-mediated tumor suppression. Moreover, as with the RB story here, mitochondrially targeted p53 was found to be sufficient to trigger apoptosis and inhibit lymphoma growth in vivo (Marchenko et al. 2000; Talos et al. 2005). In a subsequent study, p53 was discovered to interact with Bak, freeing it from the influence of the anti-apoptotic Mcl-1 and triggering Bak oligomerization to induce apoptosis (Leu et al. 2004). Finally, p53 can bind $Bcl-X_L$ in the cytoplasm but becomes displaced upon induction of the proapoptotic Bcl-2 family member Puma, liberating p53 to transit to the mitochondrial membrane, where it promotes Bax oligomerization and apoptosis (Chipuk et al. 2004, 2005). Thus, like RB, p53 can promote Bax or Bak oligomerization/ activation and membrane permeabilization Recently, the role of p53 in mitochondrial compromise was found to extend even beyond apoptosis; p53 was found to enter the mitochondrial matrix and drive necrosis, thereby contributing to stroke phenotypes (Vaseva et al. 2012). These observations highlight the similarities between p53 and RB tumor suppressor functions and further provide some possible directions for the future studies of RB's proapoptotic role at the mitochondria and the contribution of this pool of RB molecules to embryonic development, homeostasis, and cancer initiation and progression.

Conclusions

Hilgendorf et al. (2013) have identified a novel function for RB in cells. This proapoptotic role for RB at the mitochondria expands the tumor suppressor activity of an already versatile protein. Future experiments should investigate the physiological contexts in which this novel activity is most significant, including mouse models of cancer. One report in glioblastoma cells suggests that the RB family member p130 can potentiate the cell death induced by DNA damage in glioblastoma cells (Pucci et al. 2002). It would be interesting to know whether p130 and the other RB family member, p107, both of which share functional and structural similarities with RB, also share its capacity to interact with Bax and regulate apoptotic cell death directly.

The ability of two potent tumor suppressors to induce apoptosis via mitochondrial pathways raises the intriguing question of why RB and p53 evolved to participate in

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this alternate cell death pathway. Perhaps the ability to induce apoptosis at the mitochondria provides a means to trigger very rapid cell death in response to stresses without the need for the intermediary steps of transcription and translation. Alternatively, this mitochondrial effect may be one way to fine-tune the cellular response to stress. The double whammy imparted by apoptosis driven both at the mitochondrial membrane and in the nucleus provides a fail-safe mechanism to ensure the certain demise of the cell in the context of tumor suppressor action.

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